Biogenesis pathways of piRNAs loaded onto AGO3 in the *Drosophila* testis

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Supplementary information

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Materials and Methods

Western blotting

Western blotting was performed as described previously (Miyoshi *et al*. 2005). GST-Aub and -AGO3 recombinant proteins were produced in and purified from *E. coli* (Miyoshi *et al*. 2005, Gunawardane *et al*. 2007). Equal amounts of GST-Aub and GST-AGO3 were loaded on a protein gel (×1, ×1/4 dilution, ×1/16 dilution, ×1/64 dilution) for a standard curve. Lysates from testis and ovary (40μg or 80μg ×2 each) were loaded on gels. Anti-GST-Tag antibody was obtained from Medical & Biological Laboratories and used at 1:1000 dilution. After stripping of the antibody, anti-Aub (Nishida *et al*, 2007) and anti-AGO3 (Gunawardane *et al*, 2007) monoclonal antibodies were used subsequently. Each band was quantified using image analysis software, Quant mode, using Image Gauge Ver. 4.21 (FUJIFILM). The background levels of the same lane was subtracted from the quantified values. The expression levels of Aub and AGO3 proteins were compared with those of GST-Aub or GST-AGO3 (×1).

Quantitative RT-PCR

Quantitative RT-PCR was performed as reported previously (Saito *et al*. 2009). Total RNA from each sample (1μg) was used to reverse transcribe target sequences using a Transcriptor First strand cDNA Synthesis Kit (Roche) according to the manufacturer's instructions. The resulting cDNAs were amplified with a LightCycler 480 SYBR Green I Master (Roche). The primers used are shown in Supplementary Table 5.

Supplementary References:

Gunawardane LS, Saito K, Nishida KM, Miyoshi K, Kawamura Y, Nagami T, Siomi H, Siomi MC. 2007. A slicer-mediated mechanism for repeat-associated siRNA 5' end formation in Drosophila. *Science* **315**: 1587-1590.

Miyoshi K, Tsukumo H, Nagami T, Siomi H, Siomi MC. 2005. Slicer function of Drosophila Argonautes and its involvement in RISC formation. *Genes & Dev.* **19**: 2837-2848.

Nishida KM, Saito K, Mori T, Kawamura Y, Nagami-Okada T, Inagaki S, Siomi H, Siomi MC. 2007. Gene silencing mechanisms mediated by Aubergine piRNA complexes in Drosophila male gonad. *RNA* **13**: 1911-1922.

Saito K, Nishida KM, Mori T, Kawamura Y, Miyoshi K, Nagami T, Siomi H, Siomi MC. 2006. Specific association of Piwi with rasiRNAs derived from retrotransposon and heterochromatic regions in the Drosophila genome. *Genes & Dev.* **20**: 2214-2222.

Supplementary Figure 1. Ping-pong signatures of Aub- and Ago3- associated piRNAs in testis are largely restricted to those corresponding to transposons. Shown are the numbers of piRNAs that exhibit ping-pong signatures (black bars) with complementary 5' 10-mers found in pairwise library comparisons (see Supplementary Table 3). Among transposon-derived piRNAs in Aub, ~17% (4539 out of 26781 reads) have ping-pong partner piRNAs, while ~27% (1761 out of 6535 reads) among

transposon-derived piRNAs in AGO3 have ping-pong partner piRNAs. Gray bars indicate the numbers of piRNAs that do not have pairs with the ping-pong signature.

Supplementary Figure 2. The expression levels of transposons in *aub*, *ago3* and *piwi* mutant testes (upper panel) and ovaries (lower panel). The steady-state levels of transcripts from transposons were measured by quantitative RT-PCR in testes or ovaries of *aub*, *ago3* and *piwi* mutants. Note that the expression of transposons was not significantly up-regulated in *ago3* and *aub* mutant testes with the exception of F-elements. In *piwi* mutant testes, the expression of Idefix was highly up-regulated. The expression levels were standardized in each heterozygous fly (*aub*/+, *ago3*/+, *piwi*/+: red line) testes or ovaries by qRT-PCR. The error bars show the mean ± SD for at least three independent experiments.

Supplementary Figure 3. Nearly all *Su(Ste)* piRNAs that associate with Aub or AGO3 correspond to a particular region, *Su(Ste)-4*, of the antisense strand of *Su(Ste)*. Most Aub-associated and AGO3-associated *Su(Ste)* piRNAs are *Su(Ste)-4* and its variants with a 5' deletion and/or a 3' extension (see Supplementary Table 2). A: Whole shot of the piRNA distribution on Su(Ste) (accession Z11734) is shown where the blue bars correspond to the number of unique sequences (dark) or read number (light) of piRNAs from sense strand while the orange bars render the same scheme for antisense strand. Close-up of major hot spots is shown on panel B through F. B: Close-up view of Su(Ste) region 1,172 to 1,214. Sequences showed in bubbles form ping-pong pairs between sense and antisense piRNAs. Light green band represents bases inside of ping-pong base pairs. C: Close-up view of Su(Ste) region 1,241 to 1,286. Two adjacent ping-pong pair patterns are observed from this region. D: Close-up view of Su(Ste) region 1,299 to 1,351. A ping-pong pair pattern is observed among less actively transcribed piRNAs near the hottest spot. E: Close-up view of Su(Ste) region 1,487 to 1,531. One of the major hot spot is shown without observing ping-pong pair. The sequence of *Su(Ste)-6* piRNA (see Supplementary Figure 5) is indicated with underline. F: Close-up view of Su(Ste) region 1,820 to 1,856. A ping-pong pair pattern is observed.

Supplementary Figure 4. Virtually all *AT-chX* piRNAs that associate with Aub or

AGO3 correspond to a particular region, *AT-chX-1*, of the antisense strand of *AT-chX*. Aub-associated and AGO3-associated *AT-chX* piRNAs are *AT-chX-1* and its variants with a 5' deletion and/or a 3' extension (see Supplementary Table 4). (A) Whole shot of the piRNA distribution on AT-chX-A (chrX:21765977-21766248) is shown where the same color bar scheme employed in Supplementary Figure 3. (B) Whole shot of the piRNA distribution on AT-chX-B (chrX:21773610-21773780) is shown. (C) Close-up view of a AT-chX-B region where a ping-pong pair pattern is observed. The sequence of *AT-chX-71* piRNA (see Supplementary Figure 5) is indicated with underline.

Supplementary Figure 5. Genetic requirements for production of *Su(Ste)* piRNAs. (A) The abundance of *Su(Ste)-4* and *AT-chX-1* piRNAs in testes was confirmed by Northern blot analysis. Total RNAs purified from testes were used for northern blotting using DNA oligos recognizing indicated piRNAs. piRNAs [indicated as Su(Ste)-4 piRNA sense] that pair with *Su(Ste)-4* are hardly detected by northern blot analysis. We have tried to find probes that detect piRNAs from the *Su(Ste)* locus other than *Su(Ste)-4* and found two; *Su(Ste)-6* piRNA (as indicated in Supplementary Figure 3E) and *Su(Ste) pair4* piRNA (as indicated in Supplementary Table 3) were detected by northern blots, though weakly. We have tried to find probes that detect piRNAs from the *AT-chX* locus other than *AT-chX-1* and found only one; *AT-chX-71* piRNA (as indicated in Supplementary 4C) was detected by northern blots. (B) Accumulation of *Su(Ste)-6*, *Su(Ste) pair4* and *AT-chX-71* piRNAs in *ago3* and *aub* mutant testes was analyzed by northern blotting. The membrane used in Figure 5A was reprobed with DNA oligos used in (A). Both *Su(Ste)* piRNAs were not detected in testes from males lacking the *Su(Ste)* loci (*6Ste, Ycry*). In both *ago3* and aub mutant testes, *Su(Ste)-6*, *Su(Ste) pair4* and *AT-chX-71* piRNAs were hardly detected. (C) Northern analysis of the *Su(Ste)-6*, *Su(Ste) pair4* and *AT-chX-71* piRNAs in testes mutant for piRNA pathway genes. The membranes used in Figure 5C were reprobed with DNA oligos used in (A).
Sequences of oligos used for northern blotting are shown in Supplementary Table 5.

Supplementary Figure 6. The relative expression levels of Aub and AGO3 in wild-type (yw) testes were roughly determined. GST-tagged Aub and GST-tagged AGO3 recombinant proteins and testis lysates (A) or ovary lysates (B) were probed with

anti-Aub, anti-AGO3 and anti-GST antibodies. Prior to loading on a protein gel, equal amounts of GST-Aub and GST-AGO3 were diluted 1:4, 1:16 and 1:64. Then the signals from the GST fusion proteins were compared with the signals from Aub and AGO3 endogenously expressed in testes and ovaries. It should be noted that the ~1:16 dilution of GST-AGO3 was equal to the amounts of AGO3 in the testis whereas the ~1:4 dilution of GST-AGO3 was nearly equal to half of the amount of AGO3 in the ovary. In contrast, the starting amount of GST-Aub was nearly equal to half of the amount of Aub in the testis whereas the starting amount of GST-AGO3 was nearly equal to the amount of Aub in ovary. Quantification of the corresponding bands (Materials & Methods) revealed that the expression level of AGO3 in the testis was about one tenth of that of Aub, whereas the expression level of AGO3 in the ovary was about 40% of that of Aub.

Supplementary Table 1. Testis AGO3- and Aub-associated piRNAs that correspond to transposons. Transposon-derived piRNAs were identified by using Repeat Masker track and FlyBase natural transposon track. Sense/antisense was determined by comparing the strand of the transposons in the tracks and its associated piRNA mapping strand.

Supplementary Table 2. Su(Ste)-4 piRNA and its variants. Listed are the numbers of Aub- and AGO3-associated piRNAs that correspond to the *Su(Ste)* locus. *Su(Ste)-4* and its variants comprised 87.0% of *Su(Ste)* piRNAs that associate with Aub and 66.8% of *Su(Ste)* piRNAs that associate with AGO3.

Supplementary Table 3. Ping-pong pairs among AGO3- and Aub-associated piRNAs in testis. Some of the piRNAs associated with AGO3 and/or Aub in the testis show complementarity to the first 10 nt of piRNAs associated with AGO3 and/or Aub. Retrotransposons and other genomic regions from where the piRNAs arise are indicated on the right. The sequence of *Su(Ste)-pair4* piRNA (see Supplementary Figure 5) is highlighted.

Supplementary Table 4. *AT-chX-1* piRNA and its variants. Listed are the numbers of Aub- and AGO3-associated piRNAs that correspond to the *AT-chX* locus. *AT-chX-1*

and its variants comprised 90.6% of *AT-chX* piRNAs that associate with Aub and 92.6% of *AT-chX* piRNAs that associate with AGO3.

Supplementary Table 5. Sequences of Primers used for qRT-PCR to examine expression of transposons and oligos for northern blot analysis. All sequences are shown in the 5' to 3' direction.

Supplementary Figure 1
Nagao *et al.*

Supplementary Figure 2
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Supplementary Figure 3
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A

Supplementary Figure 3
Nagao *et al.*

Supplementary Figure 3
Nagao *et al.*

Supplementary Figure 3
Nagao *et al.*

Supplementary Figure 3
Nagao *et al.*

Supplementary Figure 3
Nagao *et al.*

Supplementary Figure 4
Nagao *et al*.

A

B

Supplementary Figure 4
Nagao *et al*.

Supplementary Figure 5
Nagao *et al.*

A

B

C

Supplementary Figure 6
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A **testis**

B **ovary**

Supplementary Table 1
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AGO3-associated small RNAs

No.	Sequence	Read	Genome mismatch	Annotation (category)	Annotation (detail1)	Annotation (detail2)
1	UGUUUCAUCGUUAGACGGCUCGGGC	2642	0	AT-chX		AT-chX-B
2	AAAACCUGAAGAUGAUCCUGAGUCGC	2229	0	transposon	LINE	Helena
3	UGCUCUCGAAUGUAUGCCCGAUAU	2069	0	transposon	LINE	R1-element
4	UCUCAUCGUCGUAGAACAAGCCCGA	1670	0	Su(Ste)		Su(Ste)
5	ACCCUGAACGCGCCCGAUCUCGUC	1655	no match			
6	ACCCUGAACGCGCCCGAUCUCGU	1288	no match			
7	ACCACCCUGAACGCGCCCGAUCUCGUC	1242	no match			
8	GCACACAGAGGAGAGCAGAUGAGAGCA	1079	no match			
9	ACUCGCCGAAUCCCGGGGCCGAGGGAGC	898	no match			
10	UGUUUCAUCGUUAGACGGCUCGGG	875	0	AT-chX		AT-chX-B
11	AUGGCUCUCUCAGUCGCUUCCCGGGA	857	0	ncRNA		rRNA
12	UGCUCUCGAAUGUAUGCCCGAUA	729	0	transposon	LINE	R1-element
13	UGUUUCAUCGUUAGACGGCUCGGGCG	693	0	AT-chX		AT-chX-B
14	UGCUCUCGAAUGUAUGCCCGAUAUA	693	0	transposon	LINE	R1-element
15	ACUCGCCGAAUCCCGGGGCCGAGGGA	649	no match			
16	ACCACCCUGAACGCGCCCGAUCUCGU	624	no match			
17	GGAUCCUCAACUGUGAGGGGAUCCUCAA	621	no match			
18	UGAGACUACAAACGGAUUGCGGACC	606	0	no annotation		
19	ACUCGCCGAAUCCCGGGGCCGAGGGAG	558	no match			
20	AGCUUCAUCCAAGCGAACUGUGCCCGA	479	0	transposon	LINE	R1-element
21	ACGAGUAAUAUGAGCGCCCAGAGU	441	0	transposon	LTR	opus
22	UCAACUGUGAGGGGAUCCUCAA	438	no match			
23	UGCUCUCGAAUGUAUGCCCGAUAUAC	374	0	transposon	LINE	R1-element
24	AAAACCUGAAGAUGAUCCUGAGUC	366	0	transposon	LINE	Helena
25	AAAACCUGAAGAUGAUCCUGAGUCG	357	0	transposon	LINE	Helena
26	UGUUUCAUCGUUAGACGGCUCGG	308	0	AT-chX		AT-chX-B
27	AAACCUGAAGAUGAUCCUGAGUCGC	283	0	transposon	LINE	Helena
28	CCCGAAAGCAGACGAGGUAACGUACC	274	0	transposon	Low_complexity	F-element
29	AACUGAAUAAACGAAAUGGAUGACA	255	0	transposon	LTR	invader4
30	ACGAGUAAUAUGAGCGCCCAGAGUGA	241	0	transposon	LTR	opus
31	AAAAACCUGAAGAUGAUCCUGAGUCGC	241	1	transposon	LINE	Helena
32	UUCAAGGUUACCCCAGGAUGCUGU	238	0	repeat		trf
33	UACUUGUUCCCCGGAUAGUUUAGUUAC	234	0	ncRNA		rRNA
34	GUAACUUCGGGAUAAGGAUUGGCUCU	216	0	ncRNA		rRNA
35	ACUCGCCGAAUCCCGGGGCCGAGGG	211	no match			
36	ACGAGCGCACGUGUUAGGACCCGAAA	201	no match			
37	ACAUUCAUAACCGAGCGACUGUUCAA	188	0	transposon	LTR	GATE
38	AAGAGGGCGUGAAACCGUUAAGAGGU	182	no match			
39	ACCGAAUUAAUCGCGAGGCUGCAAGA	181	0	transposon	LTR	mdg3
40	UGUUUCAUCGUUAGACGGCUCGGGCGG	175	0	AT-chX		AT-chX-B
41	CACCCUGAACGCGCCCGAUCUCGUC	173	no match			
42	CAACUGUGAGGGGAUCCUCAA	173	no match			
43	AACUGAAUAAACGAAAUGGAUGAC	165	0	transposon	LTR	invader4
44	GCCGAACGUAGUCUCUCGAGAGCGA	164	0	transposon	LTR	invader1
45	AAGAGGGACGGCCGGGGGCAUUCGU	162	no match			
46	CAAACCGAUAUGCUAUCUUGGCUGA	158	0	transposon	LINE	Doc
47	UGGGAACACCGCGUGUUGUUGGCCU	152	0	ncRNA		5SrRNA
48	UCUCAUCGUCGUAGAACAAGCCCG	151	0	Su(Ste)		Su(Ste)
49	ACCGAAUUAAUCGCGAGGCUGCAA	147	0	transposon	LTR	mdg3
50	AAGGAGUCUAACACGUGCGCGAGU	147	no match			
51	CAAGCUAGACUAUGUAUCCCCGGGUAA	147	0	transposon	LINE	Doc
52	AAAACCUGAAGAUGAUCCUGAGUCGCA	143	0	transposon	LINE	Helena
53	CCGAACUUGAGACCGCCCCUGGACGACC	139	0	Su(Ste)		Su(Ste)
54	CACCCUGAACGCGCCCGAUCUCGU	137	no match			
55	AACUUCGGGAUAAGGAUUGGCUCU	131	0	ncRNA		rRNA
56	UACGACGAGAACGAGAUCAGCGCGA	124	0	transposon	LTR	frogger
57	CAAACCAAUCCCAACCAAACCCGUCAA	123	0	transposon	LINE	G2
58	UUCAAGGUUACCCCAGGAUGCUGUGC	121	0	repeat		trf
59	AUCCUCAACUGUGAGGGGAUCCUCAA	115	no match			
60	UUUGCAAAAACGAGAGCGAUAGAGC	113	0	transposon	LTR	micropia
61	GCCAAUUACAUUAGACGAGACGGCCC	111	0	transposon	LTR	opus
62	GUGGCUUUGAGAGGCGCCGUAGUUA	109	0	no annotation		
63	AGCUUCAUCCAAGCGAACUGUGCCCG	104	0	transposon	LINE	R1-element
64	ACCCUUGAUAAACGGUUGACUUUCG	103	0	transposon	LINE	Rt1b
65	UGAGAUAUGCUAGCAGACUGGAGA	102	0	transposon	LINE	G2
66	UCGGCCCGAACCCUUCUGGUGUGU	102	0	transposon	LINE	baggins
67	AACUGAAUAAACGAAAUGGAUGACAC	101	0	transposon	LTR	invader4
68	AAGAGGGACGGCCGGGGGCAUUCG	99	no match			
69	ACCCUUGAUAAACGGUUGACUUUC	98	0	transposon	LINE	Rt1b
70	UCGGCCCGAACCCUUCUGGUGUGUU	96	0	transposon	LINE	baggins
71	UAUGGCUCUCUCAGUCGCUUCCCGGGA	96	0	ncRNA		rRNA
72	ACCCUACUGAUGAUGUGUUGUUGCC	96	no match			
73	AACUGUGAGGGGAUCCUCAA	96	no match			
74	GGUCCUGAGAGAUGGGCGAGCGCCGUUC	95	no match			
75	ACGAGUAAUAUGAGCGCCCAGAG	94	0	transposon	LTR	opus
76	UCUCAUCGUCGUAGAACAAGCCCGAGG	94	0	Su(Ste)		Su(Ste)
77	AGAGCAUAAAGACGACGAAAGUGAC	94	0	transposon	LINE	Doc3
78	UUGGGAACACCGCGUGUUGUUGGCCU	93	0	ncRNA		5SrRNA
79	UCGCCGAAUCCCGGGGCCGAGGGAGC	93	no match			
80	UGCUCUCGAAUGUAUGCCCGAUAUU	93	1	transposon	LINE	R1-element

81	CUCAACUGUGAGGGGAUCCUCAA	92	no match			
82	GAUCCUCAACUGUGAGGGGAUCCUCAA	92	no match			
83	UUCGAGAACAAACGAUACUGUUCGGC	91	0	transposon	LINE	G2
84	ACCGAAUUAAUCGCGAGGCUGCAAG	89	0	transposon	LTR	mdg3
85	AAUUCCGUAAACGAAGCCCAAAGU	89	0	transposon	LTR	Rt1c
86	UGGCUCUCUCAGUCGCUUCCCGGGA	86	0	ncRNA		rRNA
87	UGAUCUCGGAAGCUAAGCAGGGUC	86	no match			
88	AAUACAUGCCGACGGGCGCUGACCCC	83	no match			
89	CAGAUCAACAAAGUACGAUGGGGA	79	0	no annotation		
90	CACACAGAGGAGAGCAGAUGAGAGCA	78	no match			
91	AGAGCAUAAAGACGACGAAAGUGACA	77	0	transposon	LINE	Doc3
92	AAAGGAAUUGACGGAAGGGCACCACC	76	0	ncRNA		rRNA
93	GAGGUAACGAUGGAGUUUGACGGC	75	0	transposon	LTR	flea
94	AAGGAGUCUAACACGUGCGCGAGUC	73	no match			
95	AGCUUCAUCCAAGAGAACUGUGCCCGA	72	0	transposon	LINE	R1-element
96	GUGGAGCCGGGCGUGGAAUGCGAGUG	72	no match			
97	AUGGUUGCAACGGGUAACGGGGAAUC	72	0	ncRNA		rRNA
98	CAAGAUAUGACCAAAGCAAAGACAC	71	0	transposon	LINE	Cr1a
99	GAUUCCGGAGAGGGAGCCUGAGAA	70	0	ncRNA		rRNA
100	UGAGACUACAAACGGAUUGCGGACCG	69	0	no annotation		
101	CAAGAUAUGACCAAAACAAAGACAC	68	0	transposon	DNA	1360
102	CAACCAGAGAUCUAGAGGGCCCAA	66	0	no annotation		
103	UCCUCAACUGUGAGGGGAUCCUCAA	65	no match			
104	UAAACCGUCAUUCCAGCGCAAGUUGCCC	65	0	transposon	LTR	ninja-Dsim-like
105	AAAACCUGAAGAUGAUCCUGAGU	65	0	transposon	LINE	Helena
106	CUGUGAGGGGAUCCUCAA	64	2	coding gene		CG13800-RA|CG13800
107	GUGGCUUUGAGAGGCGCCGUAGUU	64	0	no annotation		
108	UCCAAUAUCACCGCGGAGACUGUCGC	63	0	transposon	LINE	Doc4
109	GGCGGGUGUUGACGCGAUGUGAUU	63	no match			
110	AAAAACCUGAAGAUGAUCCUGAGUC	63	1	transposon	LINE	Helena
111	GGCGGGUGUUUGACGCGAUGUGAUUU	63	no match			
112	AAAACCUGAAGAUGAUCCUGAGUCGU	62	0	transposon	LINE	Helena
113	GCAUUCGAGACCCAACGGCACGUGA	61	0	transposon	LINE	INE-1
114	AUGGCUCUCUCAGUCGCUUCCCGGG	60	0	ncRNA		rRNA
115	UGCACUAAUAUAAGCCCUGACGUCCGA	60	0	transposon	LTR	G4
116	UGGGCCCGAACCCUUCUGGUGUGU	59	0	transposon	LINE	baggins
117	UCUCAUCGUCGUAGAACUAGCCCGA	59	0	Su(Ste)		Su(Ste)
118	CAAGCUAGACUAUGUAUCCCCGGG	58	0	transposon	LINE	Doc
119	ACCCUUGAUAAACGGUUGACUUUCGG	58	0	transposon	LINE	Rt1b
120	GGCUUUGAGAGGCGCCGUAGUUACCAC	58	0	no annotation		
121	ACCAACGGGAUACCUCAUACUCGAG	57	0	transposon	LTR	micropia
122	GUGGCUUUGAGAGGCGCCGUAGU	55	0	no annotation		
123	GGCUUUGAGAGGCGCCGUAGUUACCACC	55	0	no annotation		
124	CAAACCAAUCCCAACCAAACCCGUCAAA	54	0	transposon	LINE	G2
125	GCCCGGCUAGCUCAGUCGGUAGAGCAUGA	53	0	ncRNA		tRNA
126	ACACGGGAAAACUUACCAGGUCCGAA	52	0	ncRNA		rRNA
127	GGUCCUGAGAGAUGGGCGAGCGCCGUU	52	no match			
128	CCAGACGUAAUACUCCACUACCCGG	51	0	transposon	LTR	HMS-Beagle
129	GGGUUCCGGGGGGAGUAUGGUUGC	51	2	ncRNA		rRNA
130	CAAGCUAGACUAUGUAUCCCCGGGU	51	0	transposon	LINE	Doc
131	AACACGGGAAAACUUACCAGGUCCGAA	51	0	ncRNA		rRNA
132	GGCUCAACUGUGAGGGGAUCUCAA	50	no match			
133	AGCUUCAUCCAAGCGAACUGUGCCC	50	0	transposon	LINE	R1-element
134	CACGGGAAAACUUACCAGGUCCGAA	50	0	ncRNA		rRNA
135	AAACCAUUCGUAGACGACCUGCU	49	no match			
136	UCGGCCCGAACCCUUCUGGUGUG	48	0	transposon	LINE	baggins
137	AUACCACCCUGAACGCGCCCGAUCUCGU	47	no match			
138	ACCGAAUUAAUCGCGAGGCUGCA	47	0	transposon	LTR	mdg3
139	AAUACAUGCCGACGGGCGCUGACCCCC	47	no match			
140	ACGGGAAAACUUACCAGGUCCGAAC	46	0	ncRNA		rRNA
141	CAACCAGAGAUCUAGAGGGCCCAAGA	45	0	no annotation		
142	ACAUUCAUAACCGAGCGACUGUUCA	45	0	transposon	LTR	GATE
143	AGAUGUCCGGGGCUGCACGCGCGCUAC	45	1	ncRNA		rRNA
144	ACAUUCAUAACCGAGCGACUGUUCAAUC	44	0	transposon	LTR	GATE
145	CCCAGAAACACAGUGUUUGACGCCAA	44	0	coding gene		CG18743-RA|Hsp70Ab
146	GGAUCUCAACUGUGAGGGGAUCCUCAA	44	no match			
147	UAAACCGUCAUUCCAGCGCAAGUUGCCCC	44	0	transposon	LTR	ninja-Dsim-like
148	CCACAGAGUACGGGGCUUACUGAAGUCA	43	0	transposon	LINE	baggins
149	UGUUUCAUCGUUAGACGGCUCGGGUCG	43	2	AT-chX		AT-chX-B
150	UGCUCUCGAAUGUAUGCCCGAUAAUU	43	no match			
151	UACAUCGAAAAGCGAGUUGACGUGU	43	0	transposon	LTR	Max
152	UCCGAAGAUAACGGGGAGCAGAGC	42	1	transposon	LTR	297
153	GCAGUUAUUACGACGAGAAGAUGGC	42	0	transposon	Unknown	RR50960
154	ACGGGAAAACUUACCAGGUCCGAA	42	0	ncRNA		rRNA
155	CCACCCUGAACGCGCCCGAUCUCGUC	41	no match			
156	ACAACAAACAACACAUACCCGCUGG	41	0	transposon	LTR	GATE
157	GAAACCAUCAGACGCCUCCAGAGGAU	41	1	transposon	LINE	G
158	CACAGAGUACGGGGCUUACUGAA	41	0	transposon	LINE	baggins
159	CGCAAACCCAAAAUGACAAUGCACGA	40	0	no annotation		
160	GCCGAACGUAGUCUCUCGAGAGCGAG	40	0	transposon	LTR	invader1
161	AAGAACAACAGUGCAGACGGCGCCCGC	40	0	transposon	DNA	H
162	CGAGAGAAGACUUUCACCCGAUGA	40	0	coding gene		CG12717-RA|CG12717
163	CCGAGGCCCUGUAAUUGGAAUGAGUAC	40	0	ncRNA		rRNA

164	GACACAGAGGAGAGCAGAUGAGAGCA	40	no match			
165	GUCUGGUUAAUUCCGAUAACGAACGA	39	0	ncRNA		rRNA
166	AAAAACCUGAAGAUGAUCCUGAGUCG	39	1	transposon	LINE	Helena
167	AACUUCGGGAUAAGGAUUGGCUC	39	0	ncRNA		rRNA
168	GCAUUCGAGACCCAACGGCACGUGACA	39	0	transposon	LINE	INE-1
169	GUAACAACGAAGAUGAUUCUGGCCC	39	0	transposon	LTR	invader4
170	CCAGACGUAAUACUCCACUACCCGGA	39	0	transposon	LTR	HMS-Beagle
171	UGUUUCAUCGUUAGACGGCUCGGGU	39	1	AT-chX		AT-chX-B
172	GCCAAUGAAAACUACACCCAGACGGA	38	no match			
173	UGCUCUCGAAUGCUUGCCCGAUAUA	38	0	transposon	LINE	R1-element
174	CACAGAGUACGGGGCUUACUGAAGUCA	38	0	transposon	LINE	baggins
175	AAACCAUUCGUAGACGACCUGCUU	38	no match			
176	UGCUCUCGAAUGUAUGCCCGAUAUUA	37	2	transposon	LINE	R1-element
177	UAACAAGAUGCAUACGAUGUGGC	37	0	transposon	DNA	1360
178	UUCGAGAACAAACGAUACUGUUCGGCC	37	0	transposon	LINE	G2
179	UAACUUCGGGAUAAGGAUUGGCUCU	36	0	ncRNA		rRNA
180	GGGACAUGUGGCGUACGGAAGACCCGC	36	no match			
181	GACAACCAGAGAUCUAGAGGGCCCAA	36	0	no annotation		
182	AAGGAGUCUAACACGUGCGCGAG	36	no match			
183	UCCGAGGAACCAAUUGAUCCAGCUGC	36	0	Su(Ste)		Su(Ste)
184	UCCUCGAUAGUAUAGUGGUUAGUAUCCC	35	0	ncRNA		tRNA
185	UGUUUCAUCGUUAGACGGCUCGGGCGU	35	1	AT-chX		AT-chX-B
186	GCUUUAAUCACGAGCAAUGCGGUGA	35	0	no annotation		
187	UCCGAAGAUAACGAGGAGCAGAGCU	35	0	transposon	LTR	297
188	GAUUCCGGAGAGGGAGCCUGAGA	35	0	ncRNA		rRNA
189	ACGACGAGAAUGAGAUCAGCGCGA	35	0	transposon	LTR	frogger
190	UGAGACUACAAACGGAUUGCGGA	35	0	no annotation		
191	AACUGAAUAAACGAAAUGGAUGACACA	35	0	transposon	LTR	invader4
192	CAAACCAAUCCCAACCAAACCCGUCA	34	0	transposon	LINE	G2
193	AAGAGCAAUGGAUGAUAUUUUAAGAGA	34	0	transposon	LTR	gypsy4
194	AUGGUUGCAACGGGUAACGGGGA	34	0	ncRNA		rRNA
195	AGAGGGCGUGAAACCGUUAAGAGGU	34	no match			
196	CACGGGAAAACUUACCAGGUCCGAAC	33	0	ncRNA		rRNA
197	GGAACACGAAACGAAAGCCCCCUGU	33	0	transposon	LINE	Rt1a
198	CGCGGGCGGCCCGGGUUCGUUUCCC	33	1	ncRNA		tRNA
199	UCCUCGAUAGUAUAGUGGUUAGUAUCCCC	33	0	ncRNA		tRNA
200	CCGAACUUGAGACCGCCCCUGGACGA	33	0	Su(Ste)		Su(Ste)
201	UGCUCUCGAAUGUAUGCCCGAUAUUU	33	2	transposon	LINE	R1-element
202	AUGGCUCUCUCAGUCGCUUCCCGG	33	0	ncRNA		rRNA
203	GUCUGGUUAAUUCCGAUAACGAACGAG	33	0	ncRNA		rRNA
204	UGAGACUACAAACGGAUUGCGGAC	33	0	no annotation		
205	GGAUCAACUGUGAGGGGAUCCUCAA	33	no match			
206	CUCGCCGAAUCCCGGGGCCGAGGGAGC	32	no match			
207	CACUUCGAAACCAGGUGCCAGAGGG	32	0	transposon	LINE	Doc
208	AGAAACCUCCCGUGGAGCAGAAGGGC	32	no match			
209	AUGGUUGCAACGGGUAACGGGGAAU	32	0	ncRNA		rRNA
210	AAAACCUGAAGAUGAUCCUGAGUCGA	31	1	transposon	LINE	Helena
211	GCAUUCAAUACAGCGAGAUGGCCCAU	31	0	transposon	LINE	Rt1a
212	ACAUUCAUAACCGAGCGACUGUUC	31	0	transposon	LTR	GATE
213	AAUUCCGUAAACGAAGCCCAAAGUU	31	0	transposon	LTR	Rt1c
214	UCCUGAGAGAUGGGCGAGCGCCGUU	31	no match			
215	ACCCUACUGAUGAUGUGUUGUUGC	31	no match			
216	CUACAUAUGGUUGAGGGUUGUA	31	0	ncRNA		rRNA
217	GUAACUUCGGGAUAAGGAUUGGCUCUG	30	0	ncRNA		rRNA
218	CCGAGGCCCUGUAAUUGGAAUGAGUA	30	0	ncRNA		rRNA
219	CAACCAGAGAUCUAGAGGGCCCAAG	30	0	no annotation		
220	AAUUCCGAUAACGAACGAGACUCUGGC	30	no match			
221	UGCUUGGACUACAUAUGGUUGAGGGUU	30	0	ncRNA		rRNA
222	UCCGAAGAUAACGAGGAGCAGAGC	30	0	transposon	LTR	297
223	CCGAACUUGAGACCGCCCCUGGACGAC	30	0	Su(Ste)		Su(Ste)
224	ACCAACGGGAUACCUCAUACUCGAGU	30	0	transposon	LTR	micropia
225	GUGGAGCCGGGCGUGGAAUGCGAGU	30	no match			
226	UCGCAAAAGUUGACCGAACUUGA	30	0	ncRNA		rRNA
227	UACACACUCAUCAACCAAAGGACGUA	30	0	no annotation		
228	ACCACCCUGAACGCGCCCGAUCUCG	30	no match			
229	AACCAAAGCUUUUGGGCUCCGGGG	29	0	ncRNA		rRNA
230	GAAACCGUCAUUCCAGCGCAAGUUGCCC	29	1	transposon	LTR	ninja-Dsim-like
231	CCCUUGAUAAACGGUUGACUUUCGG	29	0	transposon	LINE	Rt1b
232	ACAUUCAUAACCGAGCGACUGUUCAAU	29	0	transposon	LTR	GATE
233	UCAACUGUGAGGGAUCCUCAA	29	no match			
234	AGUCGGUCCUGAGAGAUGGGCGAGCGC	29	no match			
235	GCAUUCGAGACCCAACGGCACGUGAU	29	1	transposon	LINE	INE-1
236	CACGCAGUUACCGACCGUUGUGUCUG	29	0	transposon	LINE	Rt1a
237	CCCUGAACGCGCCCGAUCUCGUC	29	no match			
238	AAACCUGAAGAUGAUCCUGAGUCG	29	0	transposon	LINE	Helena
239	GCGCCGGGCCGUACCCAUAUCCGCAGC	29	no match			
240	GCCCGGAUAGCUCAGUCGGUAGAGCAUUG	29	0	ncRNA		tRNA
241	UCGGCCCGAACCCUUCUGGUGUGUUG	28	0	transposon	LINE	baggins
242	AAACCUGAAGAUGAUCCUGAGUC	28	0	transposon	LINE	Helena
243	GGUCCUGAGAGAUGGGCGAGCGCCGUUCC	28	no match			
244	GCAUUCAAUACAGCGAGAUGGCCCA	28	0	transposon	LTR	Quasimodo
245	AGAUGUCCGGGGCUGCACGCGCGCU	28	1	ncRNA		rRNA
246	CCACCCUGAACGCGCCCGAUCUCGU	28	no match			

247	CGCAAACCCAAAAAGACAAUACACGAC	28	0	repeat		trf
248	GCAUUCAAUACAGCGAGAUGGCCC	28	0	transposon	LTR	Quasimodo
249	GGAUCCUCAACUGUGAGGGGAUCUCAA	27	no match			
250	UCUCAUCGUCGUAGAACAAGCCCGG	27	1	Su(Ste)		Su(Ste)
251	AACCUAAUCACGUUCGGUGCCCGA	27	0	transposon	LINE	Rt1a
252	UCAAGAAUAACAGGACCAUCCAAGAGU	27	0	no annotation		
253	GCCCAGCUUCCCGGACAUCUUCUUC	27	0	repeat		trf
254	ACCAGAGAUCUAGAGGGCCCAAGA	27	0	no annotation		
255	UAGAGAGUGAUGCCGAGUAUAAA	27	0	repeat		trf
256	GUGUAACAACUCACCUGCCGAAGC	27	0	ncRNA		rRNA
257	UAUGGCUCUCUCAGUCGCUUCCCGGG	27	0	ncRNA		rRNA
258	AGAACCUGAAACCGUAUGCGUAC	27	no match			
259	GUGAGGUCUCCGGACGUGAUCACUG	27	0	ncRNA		rRNA
260	ACGGAAGACCCGCUCCCCGGCGCC	27	no match			
261	CAAGCUAGACUAUGUAUCCCCGGGUA	27	0	transposon	LINE	Doc
262	CAAGAUAUGACCAAAGCAAAGACA	27	0	transposon	LINE	Cr1a
263	GCUUUAAUCACGAGCAAUGCGGU	26	0	no annotation		
264	CCCGGGGCGUUCUAUAUAGUUAUGUAU	26	0	ncRNA		rRNA
265	CCCGAAAGCAGACGAGGUAACGUAC	26	0	transposon	Low_complexity	F-element
266	UUCAGAGAUUCUCGAGCUUCAGUCGCC	26	0	transposon	LTR	ninja-Dsim-like
267	GGCGAGAUGGGUGACAGCCCCGUACGG	26	no match			
268	GUUCCUCGGGAUCAAGGGCAACGAGUU	25	0	coding gene		CG40635-RA|CG40635
269	GGAGUGCCCUUCGUCCUGGGAAACGGGGC	25	no match			
270	UUCCGGAGAGGGAGCCUGAGAAAC	25	0	ncRNA		rRNA
271	ACAAACAGAAUUCCCGAAUGAGAG	25	0	transposon	LINE	Cr1a
272	CCCUUGAUAAACGGUUGACUUUCG	25	0	transposon	LINE	Rt1b
273	GUGGAAGUUACCGAUGACUUGGGAC	25	0	transposon	LTR	rooA
274	AGAGCAUAAAGACGACGAAAGUGA	25	0	transposon	LINE	Doc3
275	AAUGGGGCGAAGCUACCAUCUGUGGGAU	25	no match			
276	AUGGCUCUCUCAGUCGCUUCCCGGGAA	25	0	ncRNA		rRNA
277	UACGACGAGAACGAGAUCAGCGCGAU	25	0	transposon	LTR	frogger
278	CACAGAGUACGGGGCUUACUGAAGUC	25	0	transposon	LINE	baggins
279	GUUCCUCGGGAUCAAGGGCAACGAGU	25	0	coding gene		CG40635-RA|CG40635
280	AACUCAACUGUGAGGA	25	1	no annotation		
281	GAGAGUGGACGGGAGCGGCGGGGGCG	24	no match			
282	UGUUUCAUCGUUAGACGGCUCGGGUC	24	2	AT-chX		AT-chX-B
283	ACGGAAUGGAAUCGAGCGCUGAGU	24	0	transposon	LINE	X
284	CAGUCGCGAAAUUGGUAUCCCCGAU	24	0	transposon	LTR	diver2
285	ACCCUUGAUAAACGGUUGACUUUCGGC	24	0	transposon	LINE	Rt1b
286	UGGGAACUCUAAUGGGACUGCCGGU	24	no match			
287	UCCGGAGAGGGAGCCUGAGAAACG	24	0	ncRNA		rRNA
288	UUCUCAUCGUCGUAGAACAAGCCCGA	24	0	Su(Ste)		Su(Ste)
289	GUGUAACAACUCACCUGCCGAAGCA	24	0	ncRNA		rRNA
290	AAACCAAAGCUUUUGGGCUCCGGGGGA	23	0	ncRNA		rRNA
291	UUCAACCGGACGGGCCUAGAGUUA	23	0	coding gene		CG40635-RA|CG40635
292	GGCUUUGAGAGGCGCCGUAGUUACCACCC	23	0	no annotation		
293	UGGCUUUGAGAGGCGCCGUAGUUA	23	0	no annotation		
294	AGCAAUUCUAGAGAACGAAAGCGAG	23	0	transposon	LTR	flea
295	ACGAGCGCACGUGUUAGGACCCGAAAGA	23	no match			
296	UCCCCGCGGAAAGACGGAAAGACCCCAUG	23	no match			
297	CAAACCAAUCCCAACCAAACCCGUC	23	0	transposon	LINE	G2
298	CACAGAGUACGGGGCUUACUAAA	23	0	transposon	LINE	baggins
299	CCCAGCUCGGGGAGGUAGUGACGA	23	0	ncRNA		rRNA
300	ACGCGGGCGGCCCGGGUUCGUUUCCC	23	1	ncRNA		tRNA
301	ACCCUACUGAUGAUGUAUUGUUGCC	23	no match			
302	CUGGAUACUAGGUAUCGACGCCGA	23	0	no annotation		
303	GUUGCAAUUAAACAGAUUCCGCAGUGC	23	0	Su(Ste)		Su(Ste)
304	GAGUACUUCAGCCAGACACUGGAC	23	0	Su(Ste)		Su(Ste)
305	GAACCACUUUCAUGCUUGGGAUU	22	0	ncRNA		rRNA
306	ACCGAAUUAAUCGCGAGGCUGCAAGAC	22	0	transposon	LTR	mdg3
307	CGCAAAAGUUGACCGAACUUGAU	22	0	ncRNA		rRNA
308	AACGAGUGUUAUUGUGGGCCGGUACU	22	0	ncRNA		rRNA
309	GCAUUCGAGACCCAACGGCACGUGAUA	22	1	transposon	LINE	INE-1
310	CACAGAGUACGGGGCUUACUAAAGUCA	22	0	transposon	LINE	baggins
311	UGCCGAACGUAGUCUCUCGAGAGC	22	0	transposon	LTR	invader1
312	UGCUCUCGAAUGUAUGCCCGAUAUC	22	1	transposon	LINE	R1-element
313	UCCGAAGAUAAGUGCAGUUGGAAA	22	2	transposon	LTR	ninja-Dsim-like
314	AGCCCGUGGACGGUGUGAGGCCGGU	22	no match			
315	UUCCGAGGAACCAAUUGAUCCAGCUGC	22	0	Su(Ste)		Su(Ste)
316	UGUUUCAUCGUUAGACGGCUCGGGCGGA	21	0	AT-chX		AT-chX-B
317	AAUGAAGCGCGGGUAAACGGCGGGAGU	21	no match			
318	CCCAGCUCGGGGAGGUAGUGACGAAAA	21	0	ncRNA		rRNA
319	CAAACCGAUAUGCUAUCUUGGCUGAU	21	0	transposon	LINE	Doc
320	AAAACCUAGAAGAUGAUCCUGAGUCGC	21	no match			
321	CCGAACUUGAGACCGCCCCUGGA	21	0	Su(Ste)		Su(Ste)
322	CCCGGGGCGUUCUAUAUAGUUAUGUAUA	21	0	ncRNA		rRNA
323	GCCAAUUACAUUAGACGAGACGGCC	21	0	transposon	LTR	opus
324	UGGAAGGGCCAGUGCUUCAUGUC	20	1	no annotation		
325	AACGCGCCCGAUCUCGUCUGAUCU	20	no match			
326	AACUACUGGACCAGAUGAUAGAGGGC	20	0	transposon	Unknown	RR50960
327	GCAUUCGAGACCCAACGGCACGUGAC	20	0	transposon	LINE	INE-1
328	CAACCAGAGAUCUAGAGGGCCCA	20	0	no annotation		
329	AGCUGGGACUGAGGACUGCGACGU	20	no match			

330	CCUACAUGGAGACAUCACCGUUGG	20	0	transposon	LINE	G2
331	ACUUCGGGAUAAGGAUUGGCUCU	20	0	ncRNA		rRNA
332	GGCUUUGAGAGGCGCCGUAGUUA	20	0	no annotation		
333	AGUCGGUCCUGAGAGAUGGGCGAGC	20	no match			
334	UCUCAUCGUCGUAGAACAAGCCCGAU	20	1	Su(Ste)		Su(Ste)
335	GCCAAUUACAUUAGACGAGACGGC	19	0	transposon	LTR	opus
336	AACACGGGAAACCUCACCCGGCCCGGAC	19	no match			
337	GGACCUCAACUGUGAGGGGAUCCUCAA	19	no match			
338	AGAACUGGCGCUGCGGGAUGAA	19	1	ncRNA		rRNA
339	AAGACAAUUAUCCGAUACCGCUUAUC	19	0	transposon	LTR	flea
340	UGUGAGGGGAUCCUCAA	19	2	coding gene		CG13800-RA|CG13800
341	GGCGGGGCGCGGGACAUGUGGCGU	19	no match			
342	UUCAAGGUUACCCCAGGAUGCUGUG	19	0	repeat		trf
343	AGAUCCCCGAAUCCGGAGUGGCGGAG	19	no match			
344	UCCGAUUUCAAGUACCAGGCGAAC	19	0	repeat		RSP
345	UACUAAUUCAUCUCAGCCGGUACG	19	0	transposon	LINE	F
346	CAGAACUUGAAGUCGCCGAGGCAGC	19	0	transposon	LINE	F
347	AAAGGAAUUGACGGAAGGGCACCAC	19	0	ncRNA		rRNA
348	GAAGCCCAAACGUGAACCAGGUUCGU	19	0	transposon	Unknown	RR50960
349	ACGAAAAUGAUCGAGUGGAUAUACAGA	19	0	transposon	LTR	McClintock
350	GCUGAACGUAGUCUCUCGAGAGCGA	19	1	transposon	LTR	invader1
351	AGCUUCAUCCAAGCGAACUGUGCCCGU	19	1	transposon	LINE	R1-element
352	UAGCUAAUUAACACAAUCCCGGGGC	19	0	ncRNA		rRNA
353	UAGAGAGUGAUGCCGAGUAUAAAA	19	0	repeat		trf
354	CGGCUCAUCUCAUCCUGGGGCUGU	19	no match			
355	CUUGUUCCCCGGAUAGUUUAGUUAC	19	0	ncRNA		rRNA
356	GGAGCCGGGCGUGGAAUGCGAGUGCCU	18	no match			
357	UGCUCUCGAAUGUAUGCCCGAUAUAU	18	1	transposon	LINE	R1-element
358	AGACGUGGCGACCCGCUGAAUUU	18	no match			
359	AGGAAUUGACGGAAGGGCACCACC	18	0	ncRNA		rRNA
360	ACCACCCUGAACGCGCCCGAUCUC	18	no match			
361	AAGAGGGCGUGAAACCGUUAAGAGG	18	no match			
362	ACUCGCCGAAUCCCGGGGCCGAGGGAGCG	18	no match			
363	GAUUCCGGAGAGGGAGCCUGAGAAAC	18	0	ncRNA		rRNA
364	UACGACGAGAACGAGAUCAGCGCG	18	0	transposon	LTR	frogger
365	GAGGCAAUCACCGAGAGCAUCAACGGC	18	0	transposon	LINE	F
366	AGUGAGGUCUCCGGACGUGAUCACUG	18	0	ncRNA		rRNA
367	UGGGUUCCGGGGGGAGUAUGGUUGC	18	2	ncRNA		rRNA
368	AGCACACAGAGGAGAGCAGAUGAGAGCA	18	no match			
369	UCAAGAAUAACAGGACCAUCCAAGA	18	0	no annotation		
370	UAUACCGUUACCUUCACCACCUGGC	18	0	transposon	LINE	Doc4
371	UUCGAGAACAAACGAUACUGUUCGG	18	0	transposon	LINE	G2
372	CUGUUAAGAAACCACUCGAGCCGCC	18	0	transposon	LINE	Doc4
373	AGAACUGGCGCUGCGGGAUGAACC	18	1	ncRNA		rRNA
374	AAUACAUGCCGACGGGCGCUGACCCCCU	18	no match			
375	ACCCUUGAUAAACGGUUGACUUUCGU	18	1	transposon	LINE	Rt1b
376	CGCAAACCCAAAAAGACAAUACACGA	18	0	repeat		trf
377	AAAACUGAUAGACAGAUGGACGGAGC	18	0	Su(Ste)		Su(Ste)
378	CGCUAGCUCAUUUCCCGGGAUGUU	18	0	transposon	LINE	Fw2
379	ACCCUGAACGCGCCCGAUCUCGUCU	18	no match			
380	ACCAACGGGAUACCUCAUACUCGAGUA	18	0	transposon	LTR	micropia
381	AUCCUCAACUGUGAGGGAUCCUCAA	18	no match			
382	GAGGUCUCCGGACGUGAUCACUGU	18	0	ncRNA		rRNA
383	UGCACUAAUAUAAGCCCUGACGUCCG	18	0	transposon	LTR	G4
384	AAGGAAUUGACGGAAGGGCACCACC	17	0	ncRNA		rRNA
385	GCAUUCGAGACCCAACGGCACGUGACAC	17	0	transposon	LINE	INE-1
386	UUGUUCCCCGGAUAGUUUAGUUAC	17	0	ncRNA		rRNA
387	ACGAGCGCACGUGUUAGGACCC	17	no match			
388	ACUGUGAGGGGAUCCUCAA	17	no match			
389	GCCUGAUGACCAUAGCGAGUUGGUACCA	17	no match			
390	GCGGGCGGCCCGGGUUCGUUUCCC	17	1	ncRNA		tRNA
391	AAACGAGUGUUAUUGUGGGCCGGUACU	17	0	ncRNA		rRNA
392	CAGAACUGGCGCUGUGGGAUGAA	17	0	ncRNA		rRNA
393	GAAGAAGAGAUGCCCAGUGAGUUU	17	0	transposon	LTR	ninja-Dsim-like
394	CUGGCAUUUAUACAGUGCAUUGGACU	17	0	transposon	LINE	baggins
395	UCUCAUCGUCGUAGAACAAGCCCGU	17	1	Su(Ste)		Su(Ste)
396	UGCUCAUUCCGAACAUACACCGACAA	17	0	transposon	LINE	Doc
397	CGAGCACAACAGAUCUCACAGGUCGGC	17	0	transposon	LTR	gypsy3
398	CAAGAUAUGACCAAAACAAAGACA	17	0	transposon	DNA	1360
399	GCGGGAUUAUUACAGAUGAUGGAUCA	17	0	transposon	DNA	H
400	UGCUCUCGAAUGUAUGCCCCGAUAU	17	no match			
401	CGAACAACGAGACCACGUGUUGCA	17	0	transposon	LTR	HMS-Beagle
402	AGCUUCAUCCAAGCGAACUGUGGCCGA	17	0	transposon	LINE	R1
403	UGAGGGGAUCCUCAA	17	1	no annotation		
404	GUGCGGAGUGCCCUUCGUCCUGGGAA	17	no match			
405	CAGAGCAUUAAUCAAGACCGACCAGG	17	0	transposon	LTR	flea
406	UGCCAGGCCCGUAUAACGUUAAUGAUU	17	0	ncRNA		rRNA
407	AUAAUAAUAAAUAAAAGGUGACCGGC	17	0	AT-chX		AT-chX-A
408	AGAAAGCACCGUUUCCCGUCCGAUC	17	no match			
409	GCCCGCGUCUGAUUAGCUAGUUGGUGA	17	no match			
410	GGAUCAGCUGUGAGGGGAUCCUCAA	17	no match			
411	UCUCAUCGUCGUAGAACAAGCCCGAA	17	0	Su(Ste)		Su(Ste)
412	CAGUUCAUCAAAUCGCCUUACUG	17	1	transposon	LTR	flea

413	ACUUCAUAACCGAGCGACUGUUCAA	16	2	transposon	LTR	GATE
414	GGCUGCAAUUGUUUCCCGUGGGAC	16	0	transposon	LINE	baggins
415	UUCGAUUCCGGAGAGGGAGCCUGAGA	16	0	ncRNA		rRNA
416	AGCUUCAUCCAAGAGAACUGUGCCCG	16	0	transposon	LINE	R1-element
417	AAGCUUAUCCUUUGCUUGAUGAUUCGA	16	0	ncRNA		rRNA
418	CUUGUUCCCCGGAUAGUUUAGUUA	16	0	ncRNA		rRNA
419	AAUUCCGUAAACGAAGCCCAAAGUUC	16	0	transposon	LTR	Rt1c
420	AGCCGGACGGUGGCCAUGGAAGUCGGA	16	no match			
421	UCGGUGGCGAAGUUGGCUCAGAAG	16	1	transposon	LTR	297
422	CCUCAACUGUGAGGGGAUCCUCAA	16	no match			
423	UCGGGAUAAGGAUUGGCUCUGAAG	16	0	ncRNA		rRNA
424	ACCAAAGCUUUUGGGCUCCGGGG	16	0	ncRNA		rRNA
425	ACCGAAUUCAUCGCGAGGCUGCAAGA	16	1	transposon	LTR	mdg3
426	GGGGACGGCGGAGCGAGCGCACGGGGU	16	no match			
427	GUCUGGUUAAUUCCGAUAACGAACGAGA	16	0	ncRNA		rRNA
428	GAGCGCACGUGUUAGGACCCGAAA	16	no match			
429	CAAACCAAUCCCAACCAAACCCGU	16	0	transposon	LINE	G2
430	UCCCCUCUAAGACGUGCAACCGCGAAU	16	0	transposon	LTR	gypsy6
431	ACCAAGAUCAUCAACCCGUGUUA	16	0	transposon	LINE	Cr1a
432	AGCGUCCGACGAAUACCCUGGGG	16	no match			
433	GGGCGAAGCUACCAUCUGUGGGAUU	16	no match			
434	AGGAGUCUAACACGUGCGCGAGUC	16	no match			
435	CGAGAUGUUCAACUACACGGAUUCGCC	16	0	transposon	LTR	3S18
436	UGAGACUACAAACGGAUUGCGGACCGG	16	0	no annotation		
437	CCAGACGUAAUACUCCACUACCCGGAC	16	0	transposon	LTR	HMS-Beagle
438	GGGCUCCGGGGGAAGUAUGGUUGC	16	0	ncRNA		rRNA
439	GCCCGAUGUCCGCGAGAUUGCCGC	16	0	transposon	LTR	opus
440	GUGGCUUUGAGAGGCGCCGUAGUUU	16	1	no annotation		
441	AGAACCAUCACUGGGGCACCGUGG	16	0	transposon	Low_complexity	F-element
442	CAGAUCAACAAAGUACGAUGGGGAA	16	0	no annotation		
443	GCCCCAGUGGCCUAAUGGAUAAGGCAUC	16	0	ncRNA		tRNA
444	AGAGCAUAAAGACGACGAAAGUGACAC	15	1	transposon	LINE	Doc3
445	GAACUGGCAACAAAUUGAGCCCAACA	15	no match			
446	CCAAGAAAAAUCCCAGACCUGAUCGA	15	0	transposon	LINE	Doc
447	CAACUGUGAGGGAUCCUCAA	15	no match			
448	CCAGAUGAUAUUCCGAAAACAGCCGU	15	0	transposon	LINE	Doc4
449	AGAUGUCCGGGGCUGCACGCGCGC	15	1	ncRNA		rRNA
450	AUACCACCCUGAACGCGCCCGAU	15	no match			
451	UGUUUCAUCGUUAGACGACUCGGGC	15	1	AT-chX		AT-chX-B
452	CUUCGGGAUAAGGAUUGGCUCUGAA	15	0	ncRNA		rRNA
453	CAGGAAUCAAGGUGGCUAAGAGGA	15	0	transposon	LTR	ninja-Dsim-like
454	AAAAUUACAAAGACGACGGCCCACC	15	0	transposon	LTR	springer
455	AAUACAUGCCGACGGGCGCUGACCC	15	no match			
456	CACAGAGUACGGGGCUUACUGAAGUCAG	15	0	transposon	LINE	baggins
457	AACCACCCUGAACGCGCCCGAUCUCGUC	15	no match			
458	AACCUGAAGAUGAUCCUGAGUCGC	15	0	transposon	LINE	Helena
459	UUUGCAAAAACGAGAGCGAUAGA	15	0	transposon	LTR	micropia
460	GGCACAAUUACCAGAAGAUGUGGA	15	0	transposon	LINE	Cr1a
461	UGGGACUCAAAGAAACAAUCAGUGGU	15	0	transposon	LTR	DM88
462	CACGGCGAUGGUACUGCACUCGGUAGG	15	no match			
463	GAUGUCGUAUCACCCAGGACGUUCGA	15	0	transposon	LINE	Rt1b
464	AGUGAGGUCUCCGGACGUGAUCACU	15	0	ncRNA		rRNA
465	CGUACACAUAUUUCGCAACGUGAC	15	0	repeat		HETRP_DM
466	GCAUUCGAGACCCAACGGCACGUGACAU	15	1	transposon	LINE	INE-1
467	GCCCGAUGCCGACGCUCAUCAGACCCC	15	no match			
468	CACAGAGUACGGGGCUUACUGAAG	15	0	transposon	LINE	baggins
469	AGAAUUCACCAAGCGUUGGAUUGUUC	15	1	ncRNA		rRNA
470	AACCAAGCGCGGGUAAACGGCGGGAGU	15	no match			
471	AACGCAGGUGUCCUAAGGCGAGCUC	15	no match			
472	AGCACGGGAGACACACGGCGGGUGCU	15	no match			
473	GCCGAACGUAGUCUCUCGAGAGCGAA	15	1	transposon	LTR	invader1
474	AACCAAAGCUUUUGGGCUCCGGGGGA	15	0	ncRNA		rRNA
475	GUUCCUCGGGAUCAAGGGCAACGAG	15	0	coding gene		CG40635-RA|CG40635
476	GCUCCGGGGGAAGUAUGGUUGC	15	0	ncRNA		rRNA
477	CCCAAACGCAGACAUCGAGCGGUCAA	15	0	transposon	Low_complexity	F-element
478	UGCACAUUAAGUUUUGAUCGCCGA	14	0	transposon	LTR	invader1
479	UGAGAUAUGCUAGCAGACUGGAG	14	0	transposon	LINE	G2
480	ACUGUGAGGGGAUCCUCAACUGUGUCAA	14	no match			
481	UAGACAGAAAAUAAGGGGACACGC	14	0	transposon	LTR	opus
482	GGAGAGGUGGCAGAGUGGUCGAAUGU	14	no match			
483	UCUCAUCGUCGUAGAACAAGCCCGAG	14	0	Su(Ste)		Su(Ste)
484	UGCUCCCGAAUGUAUGCCCGAUAU	14	1	transposon	LINE	R1-element
485	AAGGCUAAAUACCGGCACGAGACCGAU	14	no match			
486	ACGGAAUGGAAUCGAGCGCUGAGUAU	14	1	transposon	LINE	X
487	CGUGACCUGAUUCAUCGCAGUUCCAGC	14	0	transposon	LINE	Circe
488	CAUGGCCGUUCUUAGUUCGUGGA	14	0	ncRNA		rRNA
489	GAUUCCGGAGAGGGAGCCUGAGAAA	14	0	ncRNA		rRNA
490	GCCUGAUGACCAUAGCGAGUUGGUAC	14	no match			
491	AAAAAACCUGAAGAUGAUCCUGAGUCGC	14	2	transposon	LINE	Helena
492	CGCGUGGGUUCGGAUCCCAUCCUCGU	14	1	ncRNA		tRNA
493	ACCUGAUUGAUAGCGUAGACGACGACC	14	0	transposon	LINE	Rt1a
494	UGGGAACACCGCGUGUUGUUGGCC	14	0	ncRNA		5SrRNA
495	AAACUUGAAAAUCCAGGUGAUCCGGC	14	0	transposon	LTR	GATE

496	AAAUCUCGCGCCGGGCCGUACCCAU	14	no match			
497	AGAGUCGUUGAAAAGUAGGCGGGCACCG	14	no match			
498	UCCGCACCCUGGGCUCGAUCUUGA	14	0	transposon	LTR	GATE
499	GCUGAGAAGACGGUCGAACUUGACUAUC	14	no match			
500	GGUGACGUGUCCGAGCGGCCGAG	14	no match			
501	GGCAACUGUGAGGGGAUCCUCAA	14	no match			
502	ACAACAAACAACACAUACCCGCUGGC	14	0	transposon	LTR	GATE
503	GUAACUUCGGGAUAAGGAUUGGCUC	14	0	ncRNA		rRNA
504	GCUUUAAUCACGAGCAAUGCGGUG	14	0	no annotation		
505	AGCCGUACCUGUUGGUUUGUCCCAUU	14	0	ncRNA		rRNA
506	UGUGCUAGAAGACGAUUUUCGGACCGA	14	0	transposon	LINE	Cr1a
507	GGAUCCUCAACUGUGAGGGGGAUCCUCAA	14	no match			
508	GCUGCGGGAUGAACCGAACGCCGGGUU	14	no match			
509	UCCCAGCUCGGGGAGGUAGUGACGAA	14	0	ncRNA		rRNA
510	GUAACUUCGGGAUAAGGAUUGGCU	14	0	ncRNA		rRNA
511	UGCCGAGAGCUGAUGACGAGCUUU	14	no match			
512	AAGACAAUUAUCCGAUACCGCUUA	14	0	transposon	LTR	flea
513	AUCCCCAGGACUUGAUCGCAUCGA	14	0	transposon	LINE	G2
514	AAAACCUGAAGAUGAUCCUGAGUCGUC	14	1	transposon	LINE	Helena
515	GAUCUCGGAAGCUAAGCAGGGUC	14	no match			
516	AUGACGGAAGGGCACCACCAGGAGUGG	14	1	ncRNA		rRNA
517	AGGCGCGGGUAACCCGUUGAACCCCAUU	13	no match			
518	UGAGACUACGAACGGAUUGCGGACC	13	1	no annotation		
519	UUCACACUCAAGAUGAUCGUGGCCA	13	0	transposon	LTR	297
520	UGCGCGAAGAAGGAGGACGAGAAGGCA	13	0	transposon	LINE	G2
521	ACGGAAUGGAAUCGAGCGCUGAGUAG	13	0	transposon	LINE	X
522	CCUGCAUGAAGAUGAUUCCGAAGA	13	0	transposon	LINE	Circe
523	CCGAUAACGAACGAGACUCAAAU	13	0	ncRNA		rRNA
524	AAUAGGGACGGUCGGGGGCAUCAGU	13	no match			
525	CGCAAACCCAAAAAGACAAUACACGACA	13	0	repeat		trf
526	GAAACCAUCAGACGCCUCCAGAGGA	13	0	transposon	LINE	G
527	GAGCGCACGUGUUAGGACCCGAAAG	13	no match			
528	AAAAACCUGAAGAUGAUCCUGAGUCGCA	13	1	transposon	LINE	Helena
529	ACGAGCGCACGUGUUAGGACCCGAAAG	13	no match			
530	GCCCAGAUGGCGGAAUUGGUAGAC	13	no match			
531	AAGAGGUAAACGGGUGGGGUCCGCGC	13	no match			
532	GCUGCGGUGUCCGAGUGGUUAAGG	13	0	ncRNA		tRNA
533	UGUUUCAUCGUUAGACGGCUCGGGCA	13	1	AT-chX		AT-chX-B
534	CAAACCGAUAUGCUAUCUUGGCU	13	0	transposon	LINE	Doc
535	UCAAGCGAUAUUCGAGGCCCAACAA	13	0	transposon	LINE	Doc4
536	GAAGCCGGUGAUCGAUAGUUCCU	13	0	repeat		trf
537	CAAAAUGACAAUGCACGACAGAGAGU	13	1	no annotation		
538	UCAGAUAAAAACGUGUUAGUAAA	13	0	no annotation		
539	CUGUUAAGAAACCACUCGAGCCGC	13	0	transposon	LINE	Doc4
540	CCACUUUCAUGCUUGGGAUUGUG	13	0	ncRNA		rRNA
541	GAGGCAAUCACCGAGAGCAUCAACGGCC	13	0	transposon	LINE	F
542	UCAGAUAAAAACGUGUUAGUAAAUA	13	0	no annotation		
543	GCCAGGCCCGUAUAACGUUAAUGAUU	13	0	ncRNA		rRNA
544	UGGCAACAUGUCGAGCUCCCAGAAC	13	no match			
545	CGAGAGUGAGAAUACCUUGGCGGACUA	13	0	coding gene		CG1490-RB|Usp7
546	AUGCCCGAUGUCCGCGAGAUUGCCGUU	13	2	transposon	LTR	opus
547	AUUCAAGGUUACCCCAGGAUGCUGU	13	1	repeat		trf
548	GGUCCUGAGAGAUGGGCGAGCGCCGU	13	no match			
549	UCUCAUCGUCGUAGAACAAGCCCGUA	13	1	Su(Ste)		Su(Ste)
550	UUCAAGGUUACCCCAGGAUGCUG	13	0	repeat		trf
551	UGUUCCCCGGAUAGUUUAGUUACG	13	0	ncRNA		rRNA
552	UCAAGAGACGCUUGAUAGCCUAGA	13	0	transposon	LINE	Rt1b
553	GUGGCAUUGAGAGGCGCCGUAGUU	13	1	no annotation		
554	UGGGCCCGAACCCUUCUGGUGUGUUGG	13	0	transposon	LINE	baggins
555	UGCCGAACGUAGUCUCUCGAGAU	13	1	transposon	LTR	invader1
556	GCCCGGGUAGCUCAGUCGGUAGAGCAUUG	13	0	ncRNA		tRNA
557	GCACACAGAGGAGAGCAGAUGAGAGCUA	13	no match			
558	UGCACCGAAAAUAUAUGCGAGGAGA	13	0	repeat		trf
559	UUAACGAUCACAAAGGCAUCACGCGC	13	0	no annotation		
560	AAACCUGAAGAUGAUCCUGAGUCGCA	13	0	transposon	LINE	Helena
561	GCUGAACCACUUUCAUGCUUGGGAU	13	0	ncRNA		rRNA
562	CGGUCCCGCUCAGAUGCCGCAGCCCGA	13	1	coding gene		CG9536-RA|CG9536
563	GCUUUAAUCACGAGCAAUGCGGUGAACG	13	0	no annotation		
564	GGAUCCUCAA	13	0	transposon	LTR	gypsy3
565	UCCGAUAACGAACGAGACUCAAA	13	0	ncRNA		rRNA
566	UCGUCGUCGUCCAGUUCGUCCAGC	13	0	coding gene		CG3732-RA|CG3732
567	UCUCAUCGUCGUAGAACAAGCCUGA	13	1	Su(Ste)		Su(Ste)
568	CCGGGAUGGACAUACCUCUGGUGUACCA	13	no match			
569	UCGCCGAAUCCCGGGGCCGAGGGAGCGAG	13	no match			
570	CAACCAGAGAUCUAGAGGGCCCAU	13	1	no annotation		
571	GAAUCCCGGGGCCGAGGGAGCGAGACCCGU	13	no match			
572	GAACCACUUUCAUGCUUGGGAUUGU	12	0	ncRNA		rRNA
573	GUGAAACUGCUUAGAGGUUAAGCCCGAU	12	0	ncRNA		rRNA
574	CAAGGUGCAGGUGGACUACAAGGG	12	no match			
575	AUCUCAACUGUGAGGGGAUCCUCAA	12	no match			
576	GCUCGUUCUGGGUAAAUAGAGUUUCUA	12	0	ncRNA		rRNA
577	ACCUGUUACAUUUGCUGAAGCGAC	12	0	transposon	LTR	Dm88
578	UAAUUCCGAUAACGAACGAGACUCA	12	0	ncRNA		rRNA

579	AUGACAGGCACCCGAAAAAGCAAGAG	12	0	transposon	LINE	accord2
580	AUGGUUGCAACGGGUAACGGGGAAUCA	12	0	ncRNA		rRNA
581	CACCCUUAACGCGCCCGAUCUCGU	12	no match			
582	GCACACAGAGGAGAGCAGAUGAGGCA	12	no match			
583	GAUGUCGUAUCACCCAGGACGUUCGACC	12	0	transposon	LINE	Rt1b
584	ACCCUGAGCGCGCCCGAUCUCGU	12	no match			
585	GUGAGGUCUCCGGACGUGAUCACUGU	12	0	ncRNA		rRNA
586	UGGGUGGUGGUGCAUGGCCGUUCU	12	0	ncRNA		rRNA
587	CUCCCACGGAACACCAGAGCAAUCCC	12	0	transposon	LINE	G
588	UGGGCUCCGGGGGAAGUAUGGUCGCA	12	1	ncRNA		rRNA
589	ACCACUUUCAUGCUUGGGAUUGU	12	0	ncRNA		rRNA
590	GCCGAACGUAGUCUCUCGAGAGCG	12	0	transposon	LTR	invader1
591	CACAGAGUACGGGGCUUACUGAAGU	12	0	transposon	LINE	baggins
592	CACAGAGUACGGGGCUUACUGAAGUCAA	12	1	transposon	LINE	baggins
593	UCCGAAGAUAACGAGGAGCAGAGCUGA	12	0	transposon	LTR	297
594	ACGAGUAAUAUGAGCGCCCAGAGUA	12	1	transposon	LTR	opus
595	AGGAGUCUAACACGUGCGCGAGU	12	no match			
596	CAAACCAAUCCCAACCAAACCCGUCAAAC	12	0	transposon	LINE	G2
597	CUCGGGUCAAUCUGUAACUGGUCAAU	12	no match			
598	UGUCUCAUCGUUAGACGGCUCGGGC	12	1	AT-chX		AT-chX-B
599	GAUCAACUGUGAGGGGAUCCUCAA	12	no match			
600	AUGCUCUCGAAUGUAUGCCCGAUA	12	0	transposon	LINE	R1-element
601	CGGGAUGGAGCAGCCCGGUAGCUCGGC	12	no match			
602	AUAUCCGUGAACUACAUCCCAACGACC	12	0	no annotation		
603	UCUCAUCGUCGUAGAACAAGCCCGAGGA	12	0	Su(Ste)		Su(Ste)
604	CCGAAGAUUAUUUGGGAUCCGCUGA	12	0	transposon	LINE	Cr1a
605	CGCUGAAUAAACAUACGCCCGGUCGC	12	0	transposon	LTR	GATE
606	GGAUCUUCAACUGUGAGGGGAUCCUCAA	12	no match			
607	GCAUCGGUGGUUCAGUGGUAGAAUGCU	12	0	ncRNA		tRNA
608	GACAACAAUACGUUAAUGACCUCAA	12	0	transposon	DNA	Ivk
609	AGAACGAUGACGAUCCCACGGCUGGAGC	11	0	transposon	LTR	ninja-Dsim-like
610	CCAGAUGGAUGGUUGGCAGGGAUAACA	11	0	transposon	LINE	G2
611	GCAUUCAAUACAGCGAGAUGGCCUAA	11	0	transposon	LINE	Quasimodo
612	ACUCGCUGAAUCCCGGGGCCGAGGGAG	11	no match			
613	ACAUUCAUAACCGAGCGACUUUCAA	11	no match			
614	UGACCACCCAUACAAAUCCCCUCGC	11	0	transposon	LINE	Doc
615	AGCAAACCAAUCCCAACCAAACCCGU	11	0	transposon	LINE	G2
616	CGCUAGCUCAUUUCCCGGGAUGUUG	11	0	transposon	LINE	Fw2
617	CGCCGAAUCCCGGGGCCGAGGGAGCGAG	11	no match			
618	UAGAAAACAACCAGGACUAUAUUCGGA	11	0	Su(Ste)		Su(Ste)
619	AAACGAGAACUUUGAAGGCCGAAGU	11	no match			
620	CUCCCCCGCCCCACCCCGCGCGCGCCGC	11	no match			
621	UUUCCUGAAAUCCGAGUUAAUAUGGC	11	0	no annotation		
622	UACAAUACAACGCUGACGACUAAA	11	0	no annotation		
623	UUCUUGAGAAUUUGGACGCCGUUUA	11	0	transposon	LINE	Cr1a
624	CACUGCUGUCACCGCUGCCGCCCGU	11	0	transposon	LINE	R1-element
625	ACGGAAUGGAAUCGAGCGCUGAGUA	11	0	transposon	LINE	X
626	GUAGUACGAGAGGAACCGCAGGUACG	11	0	ncRNA		rRNA
627	UGCCGGAGCGAGGAGGAGGACUAG	11	0	Su(Ste)		Su(Ste)
628	AAACCUUCGUAGACGACCUGCUU	11	no match			
629	GGUUAAUUCCGAUAACGAACGAGACU	11	0	ncRNA		rRNA
630	AGAGGGACGGCCGGGGGCAUUCGU	11	no match			
631	AACUACUGGACCAGAUGAUAGAGG	11	0	transposon	Unknown	RR50960
632	CAGAUCAACAAAGUACGAUGGGGAAC	11	0	no annotation		
633	UGGCUCUCUCAGUCGCUUCCCGGGAA	11	0	ncRNA		rRNA
634	GGCCGUCGAGAUGAAGCAGGUGGCCCA	11	0	coding gene		CG10691-RA|l(2)37Cc
635	ACCUUCAUCCAAGCGAACUGUGCCCGA	11	0	transposon	LINE	R1-element
636	GAAAAUCCGGGGGAGAGGGUGU	11	no match			
637	GACACCGGAACAUGUGGCUGCGAG	11	0	transposon	LTR	diver2
638	GAAGCUUAUCCUUUGCUUGAUGAUUCGA	11	0	ncRNA		rRNA
639	ACCAACGGGAUACCUCACACUCGAG	11	1	transposon	LTR	micropia
640	GGACAAAACAACAUCCCGGGAAAUGAGC	11	0	transposon	LINE	baggins
641	ACCAGAGAUCUAGAGGGCCCAAGUUG	11	no match			
642	CUAAACCAUUCGUAGACGACCUGCUU	11	no match			
643	UGUUUCAUCGUUAGACGGCUCGGCC	11	1	AT-chX		AT-chX-B
644	CACCAAGCGUUGGAUUGUUCACC	11	0	ncRNA		rRNA
645	CAAGAUAUGACCAAAACCCAGACAC	11	0	transposon	DNA	1360
646	CACAGAGUACGGGGCUUACUAAAGUCAA	11	0	transposon	LINE	baggins
647	GGAGCCGGGCGUGGGAUGCGAGUG	11	no match			
648	UGAUGCCGUCGAUGUCUUAUGCCAAUG	11	0	coding gene		CG12362-RA|CG12362
649	GCUUUAAUCACGAGCAAUGCGGUGAAC	11	0	no annotation		
650	UCGCCGAAUCCCGGGGCCGAGGGAG	11	no match			
651	CAACGGGAUACCUCAUACUCGAGUAA	11	0	transposon	LTR	micropia
652	ACGGCCUGGGCUACACACGUGCUAC	11	no match			
653	UGCGGGAUGAACCGAACGCCGGGUU	11	no match			
654	AGCUGAGAUGUGAUGACGAGUGA	11	no match			
655	CAAGCUAGACUAUGUAUCCCCGG	11	0	transposon	LINE	Doc
656	UUGGGGGCAUUAGUAUUACGACGCGA	11	0	ncRNA		rRNA
657	AUGUCUAUGAGAUGACCGUAAUGAGC	11	0	transposon	LTR	ninja-Dsim-like
658	AGGCCACCAUCGAAAGGGAAUCCGGUU	11	no match			
659	GCUCCGGGGGAAGUAUGGUUGCAA	11	0	ncRNA		rRNA
660	ACCUUGGGAAUGGCUUCACAGUUC	11	0	ncRNA		alphagamma-element
661	AGCGAAUGAUUAGAGGCCUUAGGGU	11	0	ncRNA		rRNA

662	CAAAUUAUAAGGAUGAUGAUACCGAC	11	0	no annotation		
663	ACCUCGAACAAUAAAGCAUACGCCGGU	11	0	transposon	LTR	GATE
664	UGUUUCAUCGUUAGACGGCUCGGC	11	1	AT-chX		AT-chX-B
665	AAGGAGUCUAACACGUGCGCGA	11	no match			
666	GCGGGAUGAACCGAACGCCGGGUU	11	no match			
667	AACUUCGGGAUAAGGAUUGGCUCUGA	11	0	ncRNA		rRNA
668	AGCCAGUUACCGAAGCUGCGGAU	11	no match			
669	UCCGGAGAGGGAGCCUGAGAAAC	11	0	ncRNA		rRNA
670	AGAUCAAGAAUCCCGUGCCAAGACUGC	11	0	transposon	LINE	Circe
671	GUCCUGGGCUGCACGCGCGCUACAAU	11	0	ncRNA		rRNA
672	UACCAUGAGAUUACACUGGAGGA	11	1	transposon	LINE	Cr1a
673	AGGACGAAAAGACCCCGGGACCUUC	11	no match			
674	GCCUGAUGACCAUAGCGAGUUGGUACC	11	no match			
675	CCCGAAAGCAGAUGAGGUAACGUACC	11	0	transposon	LINE	F
676	AACCCUGAACGCGCCCGAUCUCGUC	11	no match			
677	GCGGCCCGCGUCUGAUUAGCUAGUUGGU	11	no match			
678	UUAGAAGACAUAUCAUACCCAGGAAG	11	0	transposon	LTR	Idefix
679	AACUUCGGGAUAAGGAUUGGCU	11	0	ncRNA		rRNA
680	UCCGAUUGCAAGUAAUAGACGAACUGA	11	0	repeat		RSP
681	UCCAACUGAAGGAGUACGAAUGAUC	11	0	transposon	LINE	R1-element
682	UCGAUUCCGGAGAGGGAGCCUGAGA	11	0	ncRNA		rRNA
683	UUCAGAGAUUCUCGAGCUUCAGUCGC	10	0	transposon	LTR	ninja-Dsim-like
684	AAUUCCGAUAACGAACGAGACUCA	10	0	ncRNA		rRNA
685	GACCAUGAGAUUACACUGGAGGAU	10	0	transposon	LINE	Cr1a
686	CCCAGAUAAAUGAUGAAAUUUCGCGC	10	0	transposon	LINE	INE-1
687	GCCUGAUGACCAUAGCGAGUUGGU	10	no match			
688	UGAGCGAACAAACCCAACAUAGUGAAA	10	2	transposon	LTR	McClintock
689	AACGAGUGUUAUUGUGGGCCGGU	10	0	ncRNA		rRNA
690	GGCUCUCUCAGUCGCUUCCCGGGA	10	0	ncRNA		rRNA
691	UGUUUCAUCGUUAGACGACUCGGGCGG	10	1	AT-chX		AT-chX-B
692	UCUCAUCGUCGUAGAAGAAGCCCGA	10	1	Su(Ste)		Su(Ste)
693	GGUCCUGAGAGAUGGGCGAGCGCCG	10	no match			
694	UCUGCGGGUUCAGCGCCACCUGG	10	no match			
695	AUCCCACGGAACACCCGAGCAAUGC	10	0	transposon	LINE	G2
696	CUCGGCCGAUCGAAAGGGAGUCGGGUU	10	no match			
697	ACUCGCCGAAUCCCGGGGCCGGGGGA	10	no match			
698	CACACUAUCAGACAAUCCUGCGACUCGA	10	0	transposon	LTR	gtwin
699	CUGUAACUCGGGUCAACAACUCGGGUC	10	no match			
700	UGCGCGAAGAAGGAGAACGAGAA	10	0	transposon	LINE	G6
701	GAGAGAAUGAUAAGAACCAUUACGGA	10	0	transposon	LINE	Cr1a
702	ACCGAAUUAAUCGCGAGGCUGCAC	10	1	transposon	LTR	mdg3
703	UGAUCCUCAACUGUGAGGGGAUCCUCAA	10	no match			
704	UCUCAUCAUCGUAGAACAAGCCCGA	10	0	Su(Ste)		Su(Ste)
705	UGAGACUACAAACGGAUUGCGGACCGGA	10	0	no annotation		
706	UUCCGGCACAAUGCGCAUGAAUGCGA	10	0	no annotation		
707	AUGGUGACCACGGGUGACGGGGAA	10	no match			
708	UACUUGUUCCCCGGAUAGUUUAGUUA	10	0	ncRNA		rRNA
709	CCGAACUUGACAUCGCCCCUGGACGACC	10	0	repeat		trf
710	GUAACAACGAAGAUGAUUCUGGCC	10	0	transposon	LTR	invader4
711	UGUGGGAUUAUGACUGAACGCCUC	10	2	ncRNA		rRNA
712	AUUCACCAAGCGUUGGAUUGUUCACCCA	10	0	ncRNA		rRNA
713	ACCGAAUUAAUCGCGAGGCUGCAAGAA	10	1	transposon	LTR	mdg3
714	ACUCGCCGGAUCCCGGGGCCGAGGGAGC	10	no match			
715	AGCGGGGAAAGAAGACCCUGUUGAG	10	1	repeat		LSU-rRNA_Dme
716	UGCUCUCGAAUGUAUGCCCGAUG	10	1	transposon	LINE	R1-element
717	CUACAAUGGCGGUGACAGUGGGCA	10	no match			
718	UGCUCGAUCAAAAGGCCCUGGCA	10	0	transposon	LTR	Dm88
719	AGGGACGUUAAUGAAGAUGCCGUGG	10	0	coding gene		pseudogene
720	CCCGGAAUGUUCUCAGGAAGAUGACUC	10	0	ncRNA		snoRNA
721	GCGGAGUUAAAAGCGUUUACGUUCGA	10	0	transposon	LTR	micropia
722	AGCGGGGAAAGAAGACCCUGUUGAGC	10	1	repeat		LSU-rRNA_Dme
723	UCGCAUCGUCGUAGAACAAGCCCGA	10	0	Su(Ste)		Su(Ste)
724	GUGCCAGGCCCGUAUAACGUUAAUGAUU	10	0	ncRNA		rRNA
725	AACUGAAUAAACGAAAUGGAUGACACAC	10	0	transposon	LTR	invader4
726	UGGGGGCAUUAGUAUUACGACGCGAGA	10	0	ncRNA		rRNA
727	CUCGCCGAAUCCCGGGGCCGAGGGAG	10	no match			
728	GGUUCGAUUCCGGAGAGGGAGCCUGAG	10	0	ncRNA		rRNA
729	CCUAAAACAAUUCAAGAUCUGGCCGG	10	1	transposon	LTR	blood
730	UGUUUCAUCGUUAGACGGCUCGGAC	10	1	AT-chX		AT-chX-B
731	UGAUUAGACAUUCCCGUUUCUCGCGAC	10	0	transposon	LTR	rover
732	GGAUCCUUAACUGUGAGGGGAUCCUCAA	10	no match			
733	AAGCAGAACUGGCGAUGCGGGAUGA	10	2	ncRNA		rRNA
734	ACACAACAACAUCCCGGGAAAUGAGC	10	0	transposon	LINE	Fw2
735	UGGAGCCACAGUAGAUUUACAUUU	10	0	transposon	LTR	gtwin
736	UUCGAUUCCGGAGAGGGAGCCUGAG	10	0	ncRNA		rRNA
737	ACAAACGAUACCACACGAGAACGG	10	0	transposon	LINE	Rt1b
738	AACUACAACACAUCGAGCGACCGCG	10	0	transposon	LINE	Doc3
739	AUAAAUAAAAGGUGACCGGCAACGACGU	10	0	no annotation		
740	AAAGUUGACCGAACUUGAUUAUUUA	10	0	ncRNA		rRNA
741	CCCAGCUCGGGGAGGUAGUGACGAAA	10	0	ncRNA		rRNA
742	UGGGCCCGAACCCUUCUGGUGUGUU	10	0	transposon	LINE	baggins
743	GACAAAGAGAAGCGGUGUGUGAUCGAG	10	0	transposon	LTR	297
744	CACAGAGUACGGGGCUUACUAAC	10	1	transposon	LINE	baggins

745	GGGACAUGUGGCGUACGAAAGACCCGC	10	no match			
746	CUGGAGAAUAACCGAUGGACCAAUG	10	0	transposon	LTR	diver
747	CUCCCGGGGCUACGCCUGUCUGAGCGUCG	10	no match			
748	GCUUCAUCCAAGCGAACUGUGCCCGA	10	0	transposon	LINE	R1-element
749	GCAAGUCGAACGGAGUUAUAUUGU	10	no match			
750	UCCGAGGCCCUGUAAUUGGAAUGAGU	10	0	ncRNA		rRNA
751	AUGGUUGCAACGGGUAACGGGGAA	10	0	ncRNA		rRNA
752	CGAGAGAAGACUUUCACCCGAUGAGA	10	0	coding gene		CG12717-RA|CG12717
753	GGUUAAUUCCGAUAACGAACGAGACUC	10	0	ncRNA		rRNA
754	AAGCCAGUUACCGAAGCUGCGGAU	10	no match			
755	UGCUCCCGAAUGUAUGCCCGAUA	10	1	transposon	LINE	R1-element
756	AUGGCUCUCAGUCGCUUCCCGGGA	10	no match			
757	AGAAUCCGGAUGGUAUUCGACCCUUUA	10	0	coding gene		CG32240-RA|CG32240
758	CACACAACAAAACUAUAACCCAGCUGC	10	0	transposon	LINE	F
759	CAAUCUCAGAUAAAAACCCGAACGU	10	0	transposon	LTR	297
760	GCUCUCGAAUGUAUGCCCGAUAUA	10	0	transposon	LINE	R1-element
761	ACAAACAGAAUUCCCGAAUGAGA	10	0	transposon	LINE	Cr1a
762	UCAACCUGAGAGUUUGAUCCUGGCUC	10	no match			
763	AGUGCCUCGAGGAAAUCCAGACGA	10	0	transposon	LINE	R1-element
764	UAUGGCUCUCUCAGUCGCUUCCCGG	10	0	ncRNA		rRNA
765	CCAGACGUAAUACUCCACUACCCG	10	0	transposon	LTR	HMS-Beagle
766	UGCGGGAUGAACCGAACGCCGGGU	10	no match			
767	GGACAACUGUGAGGGGAUCCUCAA	10	no match			
768	UGUUUCAUCGUUAGACGGCUCAGGC	10	1	AT-chX		AT-chX-B
769	UGCUUGGACUACAUAUGGUUGAGGGU	10	0	ncRNA		rRNA
770	AACUUCGGGAUAAGGAUUGGCUCUG	10	0	ncRNA		rRNA
771	CAGAGAUGUGGCUAGCCUGAUAA	10	0	transposon	LTR	Stalker
772	UGGAACGAAAUUGGCCUGAUUAGCGG	9	0	transposon	LTR	GATE
773	CUGGACAGAAGGCUGACUUGGGG	9	0	transposon	LINE	G2
774	UGAGAUGUACAAUGCUGCCGAGUU	9	0	transposon	LINE	G2
775	CCAGACGUAAUACUCCACUACCCGGACU	9	0	transposon	LTR	HMS-Beagle
776	ACGCCAAGCACCCGAUUCUGAUACC	9	0	transposon	LTR	3S18
777	UAGUACAAUCUAUCAGACUGAGUUG	9	0	transposon	LINE	Fw2
778	CAAACCGAUAUGCUAUCUUGGCUG	9	0	transposon	LINE	Doc
779	ACCACCCUGAACGCGCCCGAUCU	9	no match			
780	GACCAUAUCCGAAGACGCGGCGCUAGUA	9	1	transposon	LINE	Rt1b
781	GACACGUUCAACCAGAUGGGCCUGGAG	9	0	Su(Ste)		Su(Ste)
782	UGGCAAUCAUGUUGGGUGAU	9	0	no annotation		
783	CGCCACUCGAACCGGACACGCCCCU	9	1	coding gene		CG10845-RA|CG10845
784	GAGUACUUCAACCAGACACUGGACGUGA	9	0	no annotation		
785	CAGUUCAUCAAAGCGCCUUACUGG	9	0	transposon	LTR	flea
786	GACCAUAUCCGAAGACGCGGCGC	9	0	transposon	LINE	Rt1b
787	AGUGCGGUAACGCGACCGAUCCCGGA	9	no match			
788	CGAGUAAUAUGAGCGCCCAGAGUGA	9	0	transposon	LTR	opus
789	UACUUGUUCCCCGGAUGGUUUAGUUAC	9	1	ncRNA		rRNA
790	UACGACGAGAACGAGAUCAGCGC	9	0	transposon	LTR	frogger
791	CAAGAAACGACACGCUUACGUCAGA	9	0	transposon	LTR	flea
792	CUGGCAUUGAAACGAUGAUGAUCUCA	9	0	transposon	LINE	Doc2
793	CGAACCGAUUAAGAGUUAAAGUGCA	9	0	no annotation		
794	AAAACCUGAAGAUGAUCCUGAGUCGCAU	9	1	transposon	LINE	Helena
795	AGCUUCAUCCAAGCGAACUGUGCC	9	0	transposon	LINE	R1-element
796	AGGGUUCGAUUCCGGAGAGGGAGC	9	0	ncRNA		rRNA
797	AGAACUGGCGCUGCGGGAUGAAC	9	1	ncRNA		rRNA
798	CCCAGAAACACAGUGUUUGACGC	9	0	coding gene		CG18743-RA|Hsp70Ab
799	CUAAAGUCCAUAUUGGAAAAGCUCGA	9	0	transposon	LTR	Idefix
800	AAACGAGUGUUAUUGUGGGCCGGUA	9	0	ncRNA		rRNA
801	CACGGGAAAACUUACCAGGUCCGA	9	0	ncRNA		rRNA
802	GCCCGGCUAGCUCAGUCGGUAGAGCAU	9	0	ncRNA		tRNA
803	CACAGAGUACGGGGCUUACUCAA	9	1	transposon	LINE	baggins
804	AAUAAUACAAACAGAGUUCCCGAAUGA	9	0	transposon	LINE	Cr1a
805	GAUGACUAUAUCAUUCUCGGGAUUAG	9	0	transposon	LTR	gypsy8
806	AACCGUUGAAGACUACUGCCGCCUA	9	1	transposon	LINE	R1-element
807	CGUUAUAUUGGUUGAACUCUAGAUA	9	0	ncRNA		rRNA
808	GUAACAACGAAGAUGAUUCUGGCCCA	9	0	transposon	LTR	invader4
809	GGCGGCGUCCGGUGAGCUCUCGCUGGCC	9	no match			
810	CUACGAUGAUGACGAGUUCGCCGAUCA	9	0	coding gene		CG3121-RA|CG3121
811	CAACCAGAGAUCUAGAGGGCCCAG	9	1	no annotation		
812	UCUGUGAGCGGGAUUUGAGCCUGAGC	9	0	transposon	LINE	Doc4
813	UGUUUCAUCGUUAGACGGCUCGGU	9	1	AT-chX		AT-chX-B
814	GAGGGACGGCCGGGGGCAUUCGU	9	no match			
815	CCGAGGCCCUGUAAUUGGAAUGAGU	9	0	ncRNA		rRNA
816	AGCACUCGCCGAAUCCCGGGGCCGAGGGA	9	no match			
817	GCAUUCGAGACCCAACGGCACGUCAAU	9	2	transposon	LINE	Rt1b
818	UACACAGGUAAACCAGGAAGGCC	9	0	no annotation		
819	UAAGACUUAAAGAUGAUGAACCCAUU	9	0	transposon	DNA	NOF
820	CGGCGGCGUCCGGUGAGCUCUCGCUGGCC	9	no match			
821	UGUUAUCUAAGGCGACAUGGGUGCA	9	0	transposon	LINE	Cr1a
822	GCACACAGAGGAGAGCAGAUGAGAGCAA	9	no match			
823	CGGGAAAACUUACCAGGUCCGAAC	9	0	ncRNA		rRNA
824	UGCUCCGAAAACAAGGCCCGUCGCUAA	9	0	transposon	LINE	G6
825	GUGCUCGUUCUGGGUAAGUAGAGUUUCU	9	1	ncRNA		rRNA
826	UCCCAGCUCGGGGAGGUAGUGACGA	9	0	ncRNA		rRNA
827	GUGAUGUCAACGUUUGAUCCCCUGG	9	0	transposon	LTR	ninja-Dsim-like

828	AAGAGGGACGGCCGGGGGCAUUC	9	no match			
829	AUGUAAGCGUAUUACCGGUGGAGUU	9	0	ncRNA		rRNA
830	UGCUCUCGAAUGUAUGCCCGAUAUAA	9	1	transposon	LINE	R1-element
831	CGGCAGGGUUCGUGACUGGGGUGA	9	no match			
832	GGCACGGACUUGGGGAAUCCGACUGUCU	9	0	ncRNA		rRNA
833	CACUCUAAAAAAGCUUCGGGACGAGA	9	0	transposon	LTR	invader3
834	ACCAGAGAUCUAGAGGGCCCAAUAA	9	1	no annotation		
835	AGACAAUUGACGGUACCGCACACC	9	0	transposon	LTR	Dm88
836	CCGAACUUGAGACCGCCCCUGGAC	9	0	Su(Ste)		Su(Ste)
837	ACCCUGAACGCGCCUGAUCUCGUC	9	no match			
838	AAAAACCUGAAGAUGAUCCUGAGUCGU	9	1	transposon	LINE	Helena
839	AUUCCGGAGAGGGAGCCUGAGAAAC	9	0	ncRNA		rRNA
840	CCCGUAAACCGCCAUCAGCAACAC	9	0	transposon	Low_complexity	F-element
841	ACACACAGAGGAGAGCAGAUGAGAGCA	9	no match			
842	UAGACAGAAAAUAAGGGGACACGCCGA	9	0	transposon	LTR	opus
843	GUGAGGUCUCCGGACGUGAUCACUGUGAC	9	0	ncRNA		rRNA
844	UGCUCGAUCAAAAGGCCCUGGCAA	9	0	transposon	LTR	Dm88
845	AACUGGCACGGACUUGGGGAAUCC	9	0	ncRNA		rRNA
846	UGGCUCUCUCAGUCGCUUCCCGGG	9	0	ncRNA		rRNA
847	AUCCCUAGAAGAACAUUGAAGCGCU	9	0	Su(Ste)		Su(Ste)
848	UUCAACUUUAACCCAAGAGACGUGACGA	9	0	transposon	DNA	transib4
849	UUGACUAUUAUUUGGAGAACCACAG	9	0	transposon	LTR	opus
850	AGCCCGUGGACGGUGUGAGGCCGGUGGC	9	no match			
851	GCAUUCGAGACCCAACGGCACGUGACU	9	1	transposon	LINE	INE-1
852	AACGAACGAGACCUUAACCUACU	9	no match			
853	GGCUCCGGGGGAAGUAUGGUUGC	9	0	ncRNA		rRNA
854	GGAACACGAAACGAAAGCCCCCUG	9	0	transposon	LINE	Rt1a
855	UACAUCGAGAAUCUCACGCCCAU	9	1	transposon	LINE	Doc3
856	CGAGGAUAGUCCCUUCAUCAUGGGU	9	0	coding gene		CG30393-RA|CG30393
857	ACCAAGAUCAUCAACCCGUGUUAUUC	9	0	transposon	LINE	Cr1a
858	AUACAUGCCGACGGGCGCUGACCCCCU	9	no match			
859	CAACCAGAGAUCUAGAGGGCCCAAGG	9	1	no annotation		
860	GCUCUCGAAUGUAUGCCCGAUAU	9	0	transposon	LINE	R1-element
861	CCGAAAGGCGUAGUCGAUGGGAACAGG	9	no match			
862	UCCAGACGCAGCCCGGCCAGGAGGC	9	0	transposon	LTR	ninja-Dsim-like
863	GCCGAACUUGAGACCGCCCCUGGAC	9	0	Su(Ste)		Su(Ste)
864	CACAGAGUACGGGGCUUACUAGUC	9	2	transposon	LINE	baggins
865	ACGAGUGUUAUUGUGGGCCGGUACU	9	0	ncRNA		rRNA
866	GCAAAGUGAACACGUCGCUAAGCGAAA	9	0	ncRNA		alphagamma-element
867	UCUGACGUGAUUCGGUUGACCUG	9	0	transposon	LINE	Doc
868	AAUGGCUCUCUCAGUCGCUUCCCGGGA	9	1	ncRNA		rRNA
869	UGCUCUCGAAUGUAUGCCCCGAUA	9	no match			
870	AACGAACGAGACUCUGGCAUGCUAA	9	no match			
871	CCGGAUUACCCGAAGUCAUUCCAUC	9	0	transposon	LTR	springer
872	GUAACCAAUGAAGUCCUCCUGAGGUGC	9	0	ncRNA		snRNA
873	CCGAACUUGAGACCGCCCCUGGACCA	9	1	Su(Ste)		Su(Ste)
874	CGAGAUCAAACCGCCGAGCUGCAAC	9	0	repeat		(TTG)n
875	UGCUCUCGAAUGUAUGCCCGAUAA	8	1	transposon	LINE	R1-element
876	UGGCCCAAACUGAUUUCCCUAAAUC	8	0	repeat		SAR_DM
877	ACUUGAGACCGCCCCUGGACGACCC	8	0	Su(Ste)		Su(Ste)
878	UCAUACGGGUAGUACAACUUACAAUUGU	8	0	ncRNA		rRNA
879	UCAACUGUGAGGGGGAUCCUCAA	8	no match			
880	CAAUCUCAGAUAAAAACCCGAACGUGGA	8	0	transposon	LTR	invader3
881	CCGUUCAAAAAUGUUCAUGCAGGAGA	8	0	transposon	LTR	Rt1c
882	AGUUUUAUCCGGUAAAGCGAAU	8	no match			
883	UGCACAGGGAUGCGCGUCUGCCGAU	8	0	transposon	LTR	ninja-Dsim-like
884	AGCAAAGUUAAAGGCCCUGCUGA	8	0	transposon	LTR	gypsy3
885	UGCAAUAGAAACGCGCGAUGAGUUGCCC	8	0	transposon	LTR	Dm88
886	CGAGAGAAGACGUUCACCCGAUGAG	8	1	coding gene		CG12717-RA|CG12717
887	CUCAGCUGUGAGGGGAUCCUCAA	8	no match			
888	UUUCUCAUCGUCGUAGAACAAGCCCGA	8	0	Su(Ste)		Su(Ste)
889	UGACUCUAGACCGGAAGCUCACUUGGC	8	1	transposon	LINE	G
890	CCGGCUCGUAGCACGCAUCCUAGA	8	0	transposon	LINE	G2
891	CGUACUCAUAUGCGCACCCACACACUU	8	0	coding gene		CG8963-RA|CG8963
892	CACAGAGUACGGGGCUUACUAAAGU	8	0	transposon	LINE	baggins
893	UGUGAACGAGUGAACUGGCAACAUU	8	1	Su(Ste)		Su(Ste)
894	CGAAAACCGCAAUUACUUCUGCACCAGCC	8	no match			
895	GCAUUCAAUACAGCGAGAUGGCCCAA	8	0	transposon	LTR	Quasimodo
896	CAUCCUAGAAUCGACCGACCAGC	8	0	transposon	LTR	flea
897	UUUGUAAGGACCUGAGAACUUGUAC	8	0	transposon	LINE	Doc4
898	UUACCAGGUCCGAACAUAAGUGUG	8	0	ncRNA		rRNA
899	ACCAGAGAUCUAGAGGGCCCAAG	8	0	no annotation		
900	UCGCAAAAGUUGACCGAACUUGAU	8	0	ncRNA		rRNA
901	ACGAGUAAUAUCCGCCCGGAGUGA	8	no match			
902	AUAUCCGUGAACUACAUCCCAACGAC	8	0	no annotation		
903	GAAGUCAUUACUGAGAUCAUACGAGA	8	0	transposon	LTR	diver2
904	AACGAGCUCCAGAUGAUGGGCCAGUUC	8	0	coding gene		CG10795-RA|CG10795
905	GUAGUCAUUACCGUGGGAUAUCA	8	0	transposon	LINE	Circe
906	GAAGCCGGUGAUCGAUAGUUCCUCU	8	0	repeat		trf
907	CGCUCACAGACUACCUAAAAGGCGC	8	0	transposon	LINE	Doc4
908	CCCAGCGAUCUCCGUACCCGGUUU	8	no match			
909	AUCGAUAUUAUCCCAGCACUACGAAGC	8	0	no annotation		
910	AACUCGCCGAAUCCCGGGGCCGAGGGAGC	8	no match			

911	CUAGAAUGCAAACAGUCACGUUGCG	8	0	repeat		HETRP_DM
912	UAACCGUCAUUCCAGCGCAAGUUGCCCC	8	1	transposon	LTR	ninja-Dsim-like
913	UGCUCUCGAAUGCUUGCCCGAUAU	8	0	transposon	LINE	R1-element
914	GUGAGGGGAUCCUCAA	8	2	transposon	LINE	baggins
915	AAGAACAACAGUGCAGACGGCGCCCG	8	0	transposon	DNA	H
916	ACGGAAUGGAACCGAGCUCUGAGUAG	8	0	transposon	LINE	X
917	CCUGCAGAGAACGCACCCGCUUGA	8	0	transposon	LINE	G2
918	GGAUCAUCAACUGUGAGGGGAUCCUCAA	8	no match			
919	AAGUGAUCGAUACGCCCCCUCAGU	8	1	transposon	DNA	S
920	AGGUGGGGAGUUUGGCUGGGGCGGCAC	8	no match			
921	AGCUAAUUAACACAAUCCCGGGGC	8	0	ncRNA		rRNA
922	CAGUUCAUCAAAGCGCCUUACUGGA	8	0	transposon	LTR	flea
923	AAGGUGCCGGAAUACACGCUCAUC	8	no match			
924	AGAUGUCCUGGGCUGCACGCGCGCUA	8	0	ncRNA		rRNA
925	GAUGUCGUAUCACCCAGGACGUCCGA	8	0	transposon	LINE	INE-1
926	AGCGUAUUACCGGUGGAGUUCUU	8	0	ncRNA		rRNA
927	UGAAACUGAAGCACGAAUUGUCCA	8	0	transposon	LTR	Dm88
928	UGAGCGAACCCGGGGAACUGAAAC	8	no match			
929	AGCAAACCAAUCCCAACCAAACCCG	8	0	transposon	LINE	G2
930	UAAGACUUAAAGAUGAUGAACCCA	8	0	transposon	DNA	NOF
931	GUGAUGUCAACGUUUGAUCCCCUGGG	8	0	transposon	LTR	ninja-Dsim-like
932	CCGAUAACGAACGAGACUCAAAUAU	8	0	ncRNA		rRNA
933	AACAGCUUAUCACCCCGGAAUUGGU	8	no match			
934	CGCUUUGAUAGAAACCUAUAGAGA	8	0	transposon	LTR	opus
935	CUGGCAUUUAUACAGUGCAUUGGA	8	0	transposon	LINE	baggins
936	CCGAAGAUGACCAAGUUGGCUACCGA	8	0	transposon	LTR	gypsy6
937	AAUGUAGGUAAGGGAAGUCGGC	8	0	ncRNA		rRNA
938	UAGAUGCUAAUCCCGAGGAUGACAU	8	0	transposon	LTR	opus
939	GCCUGAUGACCAUAGCGAGUUGGUA	8	no match			
940	CCCGAUGAGAAGAAUGGCCCGUAAAGC	8	0	transposon	LINE	RR49783
941	GAAGCAGACGGCGGCCCAGAAGCGUC	8	0	coding gene		CG12717-RA|CG12717
942	AAGUCCCGCAACGAGCGCAACCCUCGCCA	8	no match			
943	GAAACGAGAAACUUCAUCCCAGUCGGA	8	0	no annotation		
944	AAGAACAACAACAACCAGAGCAU	8	1	transposon	Unknown	RR50960
945	UUCCGAUAACGAACGAGACUCAAA	8	0	ncRNA		rRNA
946	GAGAUUCCCACUGUCCCUACCUACUAUC	8	2	ncRNA		rRNA
947	UGGGCCCGAACCCUUCUGUGUUGU	8	1	transposon	LINE	baggins
948	ACGGAAGACCCGCUCCCCGGCGCCGCUC	8	no match			
949	UCAGAUAAAAACGUGUUAGUAAAU	8	0	no annotation		
950	ACAUACCAGAUGGAUUUCGCCCGUUU	8	no match			
951	GGAUUCUCAACUGUGAGGGGAUCCUCAA	8	no match			
952	UACGACGAGAACGAGAUCAGCGCGAA	8	1	transposon	LTR	frogger
953	GACGGUACACCUGGACGAUUAGAUCA	8	1	coding gene		CG34069-RA|mt:CoII
954	GCCACAGAGGAGAGCAGAUGAGAGCA	8	no match			
955	CGUAAAACAAUUCAAGAUCUGGCCGGC	8	0	transposon	LTR	blood
956	UUCAGAAUACCCGGGCUUGAACUUC	8	0	repeat		trf
957	GCUGCGGGAUGAACCGAACGCCGGGU	8	no match			
958	CGCAAAAGUUGACCGAACUUGAUU	8	0	ncRNA		rRNA
959	ACUCGCCGAAUCCCGGGGGCCGAGGGAGC	8	no match			
960	UACUCGAAGACACCACGGUGAAAG	8	0	transposon	LTR	ninja-Dsim-like
961	UGCACUAAUAUAAGCCCUGACGUCCGAA	8	0	transposon	LTR	G4
962	CAACAAAGCACCAGGACAUGACGU	8	0	transposon	LINE	X
963	GGUAACGCAGGUGUCCUAAGGCGAGCU	8	no match			
964	GCAAGCUACAAGCGUUCCUGCAGCGAUC	8	0	transposon	Simple_repeat	I
965	GUGAGGUCUCCGGACGUGAUCACU	8	0	ncRNA		rRNA
966	GCUCUCGAAUGUAUGCCCGAUAUAC	8	0	transposon	LINE	R1-element
967	GCAGCCCGGGCGUCCUCCAGCGAUGAA	8	0	transposon	LINE	R1-element
968	AGUCGGUCCUGAGAGAUGGGCGAG	8	no match			
969	GAGGCAAUCACCGAGAGUACCGAGA	8	0	no annotation		
970	CAAGAUAUGACCUAAGCAAAGACAC	8	1	transposon	LINE	Cr1a
971	CGAGAUCAAACCGCCGAGCUGCAA	8	0	repeat		(TTG)n
972	GCCGAACUUGAGACCGCCCCUGGACGA	8	0	Su(Ste)		Su(Ste)
973	CCACACAGAGGAGAGCAGAUGAGAGCA	8	no match			
974	AAGAGGAGAAGGUGGCUUUGAGA	8	0	no annotation		
975	ACGUACUAAAACAAAAACCUCCGCGU	8	0	transposon	LINE	Rt1b
976	CAACAAGACAUCCGAUUAGUUCAUU	8	0	transposon	LINE	Circe
977	GACGCGAUGUGAUUUCUGCCCAGUGCU	8	2	ncRNA		rRNA
978	UGUGCUAGAAGACGAUUUUCGGGUUU	8	1	no annotation		
979	GGGAAUCCUCGAGCCCUGACCUCACC	8	1	coding gene		CG17119-RA|CG17119
980	CUGGUUAAUUCCGAUAACGAACGAG	8	0	ncRNA		rRNA
981	AUUCACCAAGCGUUGGAUUGUUCACC	8	0	ncRNA		rRNA
982	GCCCGUGGACGGUGUGAGGCCGGUGGC	8	no match			
983	GACGCGAUGUGAUUUCUGCCCAGUGCUC	8	2	ncRNA		rRNA
984	GUGGGUGGUGGUGCAUGGCCGUUCUU	8	1	ncRNA		rRNA
985	GGCUGGGUCGGUCGGGCUGGGGCGC	8	no match			
986	AGAACUGGCGAUGCGGGAUGAACCGA	8	no match			
987	CGGACAGAGACGGCGCAUUUUCCA	8	0	transposon	LTR	McClintock
988	GGCAAUGAGUCGUCCUGGGCAUCG	8	0	coding gene		CG11379-RA|CG11379
989	CUGUCUAUUCGCCAGGUCCCGAUAGAA	8	0	transposon	LINE	Cr1a
990	CCGGAGAAGACGUUGAAUGGAUCG	8	0	no annotation		
991	UCUGAAACAAGAAGAGGACGACGA	8	0	transposon	Low_complexity	F-element
992	UGCUAAGGGAUCCAGUUGCGAUA	8	0	transposon	DNA	1360
993	UACGUAAACAACCCCAGGCUGACUCC	8	0	transposon	LTR	Max

994	AGAACGAUGACGAUCCCACGGCUGGA	8	0	transposon	LTR	ninja-Dsim-like
995	AGUUUCAUCCAAGCGAACUGUGCCCGA	8	1	transposon	LINE	R1-element
996	GGACCUUGCUGCGGACCCUGUGGAC	8	0	repeat		trf
997	CAACUCGUUACAAUCGUGUCUACGACC	7	0	transposon	LTR	opus
998	UCUGAGGGGGAUCCUCAA	7	2	coding gene		CG3848-RC|trr
999	AAAACCCGAAGAUGAUCCUGAGUCGC	7	1	transposon	LINE	Helena
1000	ACGGGAAAACUUACCAGGUCCGAACA	7	0	ncRNA		rRNA
1001	GGUCCCGGGACAUGGCGUGGUGGGUAG	7	no match			
1002	ACCUGAGAGGGUGACCGGCCACACUGG	7	no match			
1003	UCUUCACCGUUGACUAGCCCCACACA	7	0	Su(Ste)		Su(Ste)
1004	GCUCGUCCCUAACUCCCUUACUAUUCU	7	no match			
1005	UCCGAAGAUAACGAGGAGCAGAG	7	0	transposon	LTR	297
1006	UCGGGAUAAGGAUUGGCUCUGAA	7	0	ncRNA		rRNA
1007	AACGAGCGCAACCCUUGUCCUUAGUUAC	7	no match			
1008	UCGGUGGCGAAGUUGGCUCAGAG	7	0	transposon	LTR	297
1009	CCCAGAAACACAGUGCUUGACGCCA	7	1	coding gene		CG18743-RA|Hsp70Ab
1010	CACCACAACAAAACGGCGUCGCGG	7	0	transposon	LTR	Dm88
1011	GUCAAGUUCCGUGAGAUCGGCCUGGC	7	0	coding gene		CG12699-RA|CG12699
1012	CUAACACGUGCGCGAGUCGGGGGCU	7	no match			
1013	GAGGUCUCCGGACGUGAUCACUGUGA	7	0	ncRNA		rRNA
1014	ACUGACAAUGACAGGGCAAUUUA	7	0	transposon	LTR	gypsy4
1015	AAACAACACAAUCGAUCCCGAAAU	7	0	transposon	LTR	17.6
1016	GCUGGGUCGGUCGGGCUGGGGCG	7	no match			
1017	AACUGUGAGGGAUCCUCAA	7	no match			
1018	CCGAUAACGAACGAGACUCAAAUA	7	0	ncRNA		rRNA
1019	CCCUGAACGCGCCCGAUCUCGU	7	no match			
1020	CAUGCCGACGGGCGCUGACCCCCU	7	no match			
1021	GUUCCUCGGGAUCAAGGGCAACGA	7	0	coding gene		CG40635-RA|CG40635
1022	ACCACUUUCAUGCUUGGGAUUG	7	0	ncRNA		rRNA
1023	AGAGGUUCGAAGGCGAUCAGAUACC	7	0	ncRNA		rRNA
1024	GAUCCCCGAAUCCGGAGUGGCGGAGAUG	7	no match			
1025	CGGCUCAUCUCAUCCUGGGGCUGUA	7	no match			
1026	UUGUACACACCGCCCGUCGCUACUACC	7	0	ncRNA		rRNA
1027	UGUUUCAUCGUUAGACGGCUCGGGCGC	7	1	AT-chX		AT-chX-B
1028	AAACGAAACUUUGAAGGCCGAAGU	7	no match			
1029	CCACAGAGUACGGGGCUUACUGAAGUUU	7	2	transposon	LINE	baggins
1030	UUCCAGUGUUACUCGAUACGACGACUC	7	0	no annotation		
1031	GCAUCCUAGAAUCGACCGACCAG	7	0	transposon	LTR	flea
1032	AUUCCGGAGAGGGAGCCUGAGAA	7	0	ncRNA		rRNA
1033	AUCUUAUGGUAUUGGGUUUUGAUGC	7	0	repeat		LSU-rRNA_Dme
1034	CGUUUCAUCGUUAGACGGCUCGGGC	7	1	AT-chX		AT-chX-B
1035	CAAUCUACAAAGAGCUGAAAAUCCG	7	0	transposon	Low_complexity	F-element
1036	AAAAACCUGAAGAUGAUCCUGAGU	7	1	transposon	LINE	Helena
1037	GGCACGGACUUGGGGAAUCCGACUGUC	7	0	ncRNA		rRNA
1038	GACAACCAGAGAUCUAGAGGGCCC	7	0	no annotation		
1039	GUGGAGCCGGGCGUGGAAUGCGAGUGC	7	no match			
1040	AGAGGUUCGAAGGCGAUCAGAUACCG	7	0	ncRNA		rRNA
1041	CAGGCCCAUCUGGUUGAACGUGUCCUG	7	0	Su(Ste)		Su(Ste)
1042	AGUGGCAUGUCUCGAGGUUAACCCGUG	7	no match			
1043	ACCAAGAUCAUCAAACCCGUGUUA	7	0	transposon	LINE	Cr1a
1044	CAACUGUGAGGGGGAUCCUCAA	7	no match			
1045	AAAUCUGGAUACUAGGUAUCGACGCCG	7	0	no annotation		
1046	GAGAGGCGCCGUAGUUACCACCCC	7	0	no annotation		
1047	ACGGGAAAACUUACCAGGUCCGA	7	0	ncRNA		rRNA
1048	CAGGGGGAUCUUUGAGUGAUGAAA	7	1	transposon	LINE	baggins
1049	UCCGAUUUCAAGUACCAGACAAAC	7	0	repeat		RSP
1050	GCAUUGAUUACGUCCCUGCCCUUUGUA	7	0	ncRNA		rRNA
1051	GAUGGUGUGCAUCGAUCGCUGUGUGGC	7	0	coding gene		CG9878-RA|Tim10
1052	GGAGAGAUACCCAAGUGGCCAACGGG	7	no match			
1053	GAUUCCGGAGAGGGAGCCUGAG	7	0	ncRNA		rRNA
1054	AUCCCACGGAACACCCGAGCAAUGCCAU	7	1	transposon	LINE	G2
1055	AAAGAGGGCGUGAAACCGUUAAGAGGU	7	no match			
1056	GGAACACGAAACGAAAGCCCCCUGUC	7	0	transposon	LINE	Rt1a
1057	CCGGGGCUACGCCUGCCUGAGCGUCGCUU	7	no match			
1058	ACCGAAUGCUAUAAGUUACAAAGAUG	7	0	transposon	LTR	297
1059	UGAGCAAUAACAGGUCUGUGAUGCCCUU	7	0	ncRNA		rRNA
1060	AGGCGCGGGUAACCCGUUGAACCCCAUUC	7	no match			
1061	AAACCUUCGUAGACGACCUGCU	7	no match			
1062	CAAUCCGAUACAUUUCACGCCGACA	7	0	transposon	LINE	INE-1
1063	AAAGCGAAUGAUUAGAGGCCUUAGG	7	0	ncRNA		rRNA
1064	AGAUUAAGCCAUGCAUGUCUAAGUACA	7	0	ncRNA		rRNA
1065	UGAGAUAGUCGGGCUUGAUUGGGAAA	7	0	ncRNA		rRNA
1066	GACGCGAUGUGAUUUCUGCCCAGUGC	7	2	ncRNA		rRNA
1067	GCCUGAGAAACGGCUACCACAUCUAAG	7	0	ncRNA		rRNA
1068	CACGAAACUGGAACGGGAUUAAGA	7	0	no annotation		
1069	GACACGCAUGGAUCCACCACCUUCAGGA	7	1	coding gene		CG3982-RA|CG3982
1070	CGACUCGUUUUUAAAUUUCGCAACAAGC	7	0	transposon	LINE	Cr1a
1071	UGUUAUCUAAGGCGACAUGGGUGC	7	0	transposon	LINE	Cr1a
1072	ACUCGCCGAAUCCCGGGGGCCGAGGGAG	7	no match			
1073	UGUCUCAUCGUUAGACGGCUCGGG	7	1	AT-chX		AT-chX-B
1074	CUGUUUCAUCGUUAGACGGCUCGGGC	7	0	AT-chX		AT-chX-B
1075	GUCGAGAGAUAUCGGAAGAGUUU	7	0	ncRNA		rRNA
1076	CACUUUCAUGCUUGGGAUUGUGA	7	0	ncRNA		rRNA

1077	AGGACGAUAACGACUAGAACCGCGUA	7	no match			
1078	ACGAGCGCACGUGUUAGUACCCGAA	7	no match			
1079	UUCGGCCCGAACCCUUCUGGUGUG	7	0	transposon	LINE	baggins
1080	GAUUAACGAGAUUCCUACUGUCCCU	7	0	ncRNA		rRNA
1081	AAGACAAUCUGGGAGGUGCCCGA	7	no match			
1082	GCGGAAAGAGAGUAAUUUUGGCC	7	0	transposon	LINE	Cr1a
1083	UGAGAUUACAAACGGAUUGCGGACC	7	1	no annotation		
1084	UCACAUGAACUUGGAAUUCCCAGUAAGU	7	0	ncRNA		rRNA
1085	GGCUCAUCUCAUCCUGGGGCUGU	7	2	no annotation		
1086	GAUUCAAAGACGAGGUUGACGCCGC	7	0	transposon	LTR	gtwin
1087	GGAGGAGCCGGAACUGGGAGCCCAGGA	7	0	coding gene		CG11319-RA|CG11319
1088	CUACGUCCGUUGGAUUAUGCCUGAA	7	0	ncRNA		rRNA
1089	AACUGAAUAAACGAAAUGGAUGA	7	0	transposon	LTR	invader4
1090	GGUAGUAGAACACGUUUUAUUCUGGA	7	0	transposon	LTR	ninja-Dsim-like
1091	GACGUAGAAAAACUUGGAGACGA	7	0	transposon	LTR	diver2
1092	ACGCCCGAUACAUAGUGUCGAAUCG	7	1	coding gene		CG40635-RA|CG40635
1093	UUCAAGGUUACCCCCAGGAUGCUGUGC	7	no match			
1094	CCAUGGACUCAGAUCAGCUUAAGGC	7	0	transposon	LTR	gypsy6
1095	UCGCGAAAGUCGCUACCGAUGUACCCACG	7	0	coding gene		CG8982-RA|Acp26Aa
1096	ACCGAUUAAUCGCGAGGCUGCAA	7	no match			
1097	ACGACUAUGAGAGUGCAGCUGAGGC	7	0	transposon	LTR	297
1098	AGUGAGGUCUCCGGACGUGAUCACUGU	7	0	ncRNA		rRNA
1099	CUCGCAAGAAAACUAAUCCGAGUAUG	7	0	transposon	LINE	Doc
1100	GCUAAACCAUUCGUAGACGACCUGCUU	7	no match			
1101	CAAAGCUUUUGGGCUCCGGGGGAA	7	0	ncRNA		rRNA
1102	AGGGACGUUGAUGAAGAUGCCGUGGU	7	0	coding gene		CG12717-RA|CG12717
1103	CCAGACGUGAUACUCCACUACCCAGA	7	0	transposon	LTR	HMS-Beagle2265
1104	CUGGUAAAGCGAAUGAUUAGAGGCC	7	0	ncRNA		rRNA
1105	UGUUCCAUCGUUAGACGGCUCGGGCGG	7	1	AT-chX		AT-chX-B
1106	AGAGGUGAAAUUCUUGGACCGGCGC	7	2	ncRNA		rRNA
1107	UGACGGAAGGGCACCACCAGGAGU	7	0	ncRNA		rRNA
1108	AAGGUGGAAAUAAAGAAAAGGAUCCUCAA	7	no match			
1109	CGUAACUUCGGGAUAAGGAUUGGCUCU	7	0	ncRNA		rRNA
1110	ACCCUGAACGCGACCGAUCUCGUC	7	no match			
1111	ACGCGACCUCAGAUCAGACGUGGC	7	no match			
1112	GUAUACCUAUACAUUACCGCUAAA	7	0	ncRNA		rRNA
1113	ACGAGCUGGAUAACUUGUUGCCCCACA	7	0	coding gene		CG8251-RA|Pgi
1114	CAACGAGUUCCUCUGCCACAUGCCC	7	1	Su(Ste)		Su(Ste)
1115	CAAGAUAUGACCAAAGCAAAGACACUAG	7	0	transposon	LINE	Cr1a
1116	AACGCUUUCAACACCGCCAGAUGGCCCA	7	1	transposon	LINE	F
1117	AACCAAGCUAGAGUACAGAUCGAAA	7	0	transposon	LINE	jockey
1118	ACCUGAGAAGAAACCGAUGGCGCCC	7	0	coding gene		CG8787-RA|Asx
1119	ACAACCGAUAUAAUUCUAACUGAGCACC	7	0	transposon	LTR	G4
1120	ACAAAGUUAACGCGAGCCAAAUUUC	7	0	transposon	LTR	gypsy12
1121	CUUGGUUAGGUCGAUUCUUUUUGAAU	7	no match			
1122	AAGAAGAUGAAGCGCAACAAGCCGA	7	0	transposon	LTR	ninja-Dsim-like
1123	CAGCUGCCAUACUGCGAUCCACUGAA	7	0	ncRNA		rRNA
1124	CUAGCAAAAAUAUUUGAUCCGUUAGG	7	0	transposon	DNA	HB
1125	UCGGACAAUUAUUUUUUAUGUUGUGGC	7	0	repeat		SAR_DM
1126	UUCAGGGGGAUCUUUGAGUGAUGAA	7	0	transposon	LINE	baggins
1127	CGCGAGAGGUGAAAUUCUUGGACCGUCG	7	0	ncRNA		rRNA
1128	GUGGCUUUGAGAGGCUCCGUAGUUA	7	1	no annotation		
1129	AAACCUGAAGAUGAUCCUGAAGUCU	7	no match			
1130	UAAGGCUGAGAACUGAUGACGAGUGUU	7	no match			
1131	CACCUUGGGAAUGGCUUCACAGUUC	7	0	ncRNA		alphagamma-element
1132	ACUCUAGAUAACCUCGGGCCGAUCGC	7	no match			
1133	CAGAUCGAGAACUGGUUUGCUGAGCGCU	7	0	coding gene		CG17210-RA|scpr-B
1134	CCGCAAGCGUAUCAAUCUGCGUCCCGAC	7	0	coding gene		CG12334-RA|Atg8b
1135	AGCAGGACGGAAAGACCCCGGGACCUUU	7	no match			
1136	UGCUCUCGAAUGUAUGCCCGAUAAU	7	2	transposon	LINE	R1-element
1137	GAGGUCUCCGGACGUGAUCACUGUGAC	7	0	ncRNA		rRNA
1138	CACAGAGUACGGGGCUUACUAAAGUC	7	0	transposon	LINE	baggins
1139	CAACCUGAGAGUUUGAUCCUGGCUC	7	no match			
1140	GACUGCAGAUCGAGAGGUCCCCGGUU	7	0	ncRNA		tRNA
1141	UAUGAAGUAGCCCGUCUUCCGAA	7	1	transposon	LTR	gypsy4
1142	AGGAGUCUAACACGUGCGCGAG	7	no match			
1143	AACCAUUCGUAGACGACCUGCU	7	no match			
1144	AGAUCCCCGAAUCCGGAGUGGCGGAGA	7	no match			
1145	UACGUAAACAACCACAGGCUGACUC	7	1	transposon	LTR	Max
1146	UGAGAUAGAUUCCCGGGAAUUCAAAAG	7	0	no annotation		
1147	AUACCGCCCUAGUUCGAACCAUAAA	7	1	ncRNA		rRNA
1148	CCCAGAAACACAGUGUUUGACGCC	7	0	coding gene		CG18743-RA|Hsp70Ab
1149	GCAUUCGAGACCCAACGGCACGUGAUU	7	2	transposon	LINE	INE-1
1150	ACCAACGGGAUACCUCACGCUCGAGUA	7	2	transposon	LTR	micropia
1151	AGGUAGUGACGACAAUUGCAUCGGUU	7	0	coding gene		CG1059-RA|Karybeta3
1152	GAAUAUUCUGUGCAUGGGAUAAUGA	7	0	ncRNA		rRNA
1153	UGUUUCAUCUUUAGACGGCUCGGGC	7	1	AT-chX		AT-chX-B
1154	CUUAUCCUUUGCUUGAUGAUUCGA	7	0	ncRNA		rRNA
1155	CUCCCACGGAACACCAGAGCAAUCCCACC	7	0	transposon	LINE	G
1156	ACUUCGGGAUAAGGAUUGGCUCUGA	7	0	ncRNA		rRNA
1157	ACCUGAUUGAUAGCGUAGACGACGAC	7	0	transposon	LINE	Rt1a
1158	GAUGACUAUAUCAUUCUCGGGAUUA	7	0	transposon	LTR	gypsy8
1159	GAUCCUCAACUGUGAGGGGAUCUCAA	7	no match			

1160	CACCCUGAACGCGCCCGAUCUCG	7	no match			
1161	ACCGAAUUAAUCGCGAGGCUGCAAGU	7	1	transposon	LTR	mdg3
1162	AGAACACGCAAGAGCAGCUCCAGC	7	0	transposon	LINE	G6
1163	GCGGGAGCCCCGGGGAGAGUUCUCUU	7	no match			
1164	AGAUCAGUGAAAUCAUCUUCCGGGA	7	0	transposon	LTR	micropia
1165	CGAGUGUUAUUGUGGGCCGGUACU	7	0	ncRNA		rRNA
1166	CAGGCAAGGUGUUGUAAGUGGUUGAGC	7	0	ncRNA		rRNA
1167	GGAGAGGUGGCAGAGUGGUCGAAUG	7	no match			
1168	UGAGACUACAAACGGAUUGCGACC	7	2	no annotation		
1169	CGCCCCCCGGCCCCGCGUCCUCCCUC	7	no match			
1170	UAACGAGUUCCUCUGCCGCGUGCC	7	1	coding gene		CG40719-RA|CG40719
1171	GACUGCAGAUCGAGAGGUCCCCGGUUC	7	0	ncRNA		tRNA
1172	AAGAACGAAAGUCGGAGGUUCGAAGA	7	no match			
1173	ACUUCGGGAUAAGGAUUGGCUCUGAA	6	0	ncRNA		rRNA
1174	CAACAAAGCACCUGGACAUGACGUC	6	0	transposon	Simple_repeat	I
1175	AUUCGUGAUGGGGAUCGGGGAUUGC	6	no match			
1176	CUGUGAGGGGACCUCAA	6	2	coding gene		CG2083-RB|CG2083
1177	GCCCUAGUUCUAACCAUAAACGAUGCC	6	0	ncRNA		rRNA
1178	AUACAUGCCGACGGGCGCUGACCCC	6	no match			
1179	GGAUCCUCAACUGUGAGGGAUCCUCAA	6	no match			
1180	CAUGGCUCUCUCAGUCGCUUCCCGGGA	6	1	ncRNA		rRNA
1181	CCCGAAGGCAGAUGAGGUAGCGUAC	6	1	transposon	LINE	FW
1182	CAAACGUAAUGUUACGGUGCCCAA	6	0	ncRNA		rRNA
1183	UUCCUCGGGAUCAAGGGCAACCAGU	6	0	repeat		trf
1184	UAACGAGAGAAGCAUUUUUGCCGC	6	0	transposon	LTR	gypsy12
1185	GCCCUGAAGACCAUUACUCGUCAUU	6	0	coding gene		pseudogene
1186	UUCCAGCGAAUACCGAGUGUUUUGGC	6	1	transposon	LTR	GATE
1187	GCUGCUAGAGAGAACUGAUAAAGUU	6	0	transposon	LTR	flea
1188	UGCUAGAAGACGAUUUUCGGGCCGA	6	0	transposon	LINE	Cr1a
1189	CAACCAAAGAGGUAUUGCACUAGCGA	6	0	transposon	LINE	X
1190	CGCGCGCCAAGGAGCACCCCGUCUUUCUG	6	0	coding gene		CG10340-RA|CG10340
1191	CGCUUCAUCCAAGCGAACUGUGCCCGA	6	1	transposon	LINE	R1-element
1192	AAAACCUAGAAGAUGAUCCUGAGUCG	6	no match			
1193	UAUGCUAAAAGUUAUCCUUACCCAGUC	6	0	transposon	LTR	opus
1194	AUAGGACCCUGAAAGCACCGCAUC	6	no match			
1195	UCUGAUGUUACGCCCGAAGAGAUA	6	0	transposon	Low_complexity	F-element
1196	AACUGAAUAAACGAAAUGGAUGACUA	6	2	transposon	LTR	invader4
1197	CCUGAAGGAAUGAAGUUUUCACCGAC	6	no match			
1198	CUCGACGGUAACCGAUGGUUAGGC	6	0	transposon	LTR	Quasimodo
1199	UGCCGGGGCGAGGAGGAGGACUAGCUG	6	0	Su(Ste)		Su(Ste)
1200	GAUUCCGCAGUGCGUUGAUAUGGCCAG	6	0	Su(Ste)		Su(Ste)
1201	UGCACAGGGAUGCGCGUCUGCCGA	6	0	transposon	LTR	ninja-Dsim-like
1202	GAGAGAUGGGCGAGCGCCGUUCCGAA	6	no match			
1203	CCAGCCUACAUGGAGACAUCACCG	6	0	transposon	LINE	G2
1204	UUUGCAAAAACGAGAGCGAUAGAG	6	0	transposon	LTR	micropia
1205	GACGAAAUAAACAAGCACUCGAGCAGA	6	0	transposon	Simple_repeat	I
1206	CGACUCGUUUUUAAAUUUCGCAAC	6	0	transposon	LINE	Cr1a
1207	GAUCUCAACUGUGAGGGGAUCCUCAA	6	no match			
1208	UGCUCUCGAAUGCUUGCCCGAUAA	6	1	transposon	LINE	R1-element
1209	CAACAAAUCAUGAAAGAGCGCGAGU	6	0	no annotation		
1210	AGAUCAGACGUGGCGACCCGCUGAAU	6	no match			
1211	AGAUGUCCUGGGCUGCACGCGCGCUAC	6	0	ncRNA		rRNA
1212	UAGCUGAUCCAGAUCGAGGUUCUCCGGG	6	0	transposon	LTR	flea
1213	UGACAGUCCAAACCCCGUAACUUGAU	6	1	transposon	LTR	GATE
1214	UGCGCAGAGUCAUGCCAGUACUGGA	6	0	transposon	LINE	G2
1215	GUGGGUGGUGGUGCAUGGCCGUU	6	1	ncRNA		rRNA
1216	UGGCCACACUGUCUCCUCCUGAGA	6	no match			
1217	CAGUCUAUGAAAGGUGUGUUGCCCAGC	6	0	transposon	LTR	GATE
1218	UUGGUUCGAAAGCUAUCGUGCUGA	6	0	transposon	LTR	DM88
1219	AUCCAACUGUGAGGGGAUCCUCAA	6	no match			
1220	GCACACUGAGGAGAGCAGAUGAGAGCA	6	no match			
1221	GCCGUGAUCGUCUAGUGGUUAGGACCCC	6	0	ncRNA		tRNA
1222	GCCGAGAUAUGCUGCGAGCUGCCAA	6	1	transposon	Simple_repeat	I
1223	AGGCCAUUAUACACUAUAGUUAGUGA	6	0	transposon	LTR	roo
1224	AAACGAGUGUUAUUGUGGGCCGGUAC	6	0	ncRNA		rRNA
1225	AACCACCCUGAACGCGCCCGAUCUCGU	6	no match			
1226	UUCAGCGAGAUACUGGACGUGAUCCU	6	0	repeat		trf
1227	UUCCGGAGAGGGAGCCUGAGAAACG	6	0	ncRNA		rRNA
1228	UCCGGAGAGGGAGCCUGAGAAACGG	6	0	ncRNA		rRNA
1229	UUGGGCCCGAACCCUUCUGGUGUG	6	0	transposon	LINE	baggins
1230	CACACUGGAAAAAUCUCCGAUUGG	6	0	transposon	LTR	opus
1231	CAAUUCAUCAAGCGGUGGAUCCUGCGUUU	6	no match			
1232	CUCGGAAGCUAAGCAGGGUCGGGC	6	no match			
1233	CUGGGAAACGGGGCGCGGCCGGAA	6	no match			
1234	CGGACAGAGACGGCGCAUUUUCC	6	0	transposon	LTR	McClintock
1235	AUUCCGAUGCAUGGACCCUCGAAGCA	6	0	coding gene		CG11181-RB|cup
1236	UCCCCGCGGAAAGACGGAAAGACCCCA	6	no match			
1237	AUCCCCUGGACCUGUACUGACGCUCAU	6	no match			
1238	AUAGGGACGGUCGGGGGCAUCAGU	6	no match			
1239	AGGGACGUUAAUGAAGAUGCCGUGGU	6	0	coding gene		pseudogene
1240	AGGACGUUCACACGAGGGUUACUAUGA	6	0	transposon	LTR	opus
1241	CCACAGAGUACGGGGCUUACUGA	6	0	transposon	LINE	baggins
1242	AUGGAACACAUAGGGGCUUCGAGG	6	0	transposon	LTR	HMS-Beagle

1243	GUUGUAUUACGAGAUUGAGGACUGGC	6	0	coding gene		CG14579-RA|CG14579
1244	UGUCCGCCCUUCGUGCCUCAAGAUAA	6	1	transposon	DNA	1360
1245	UGCUCUCGAAUGUAUGCCCGAUGUA	6	1	transposon	LINE	R1-element
1246	GACGAGAUAAACCAGCACUCGAGC	6	0	transposon	LINE	Doc3
1247	UCUUGCAAUAAAGCUCGUCCUCGGGU	6	0	transposon	LINE	G6
1248	AUGGCUCUCUCAGUCGCUUCCCGGGAAA	6	0	ncRNA		rRNA
1249	AAAGAAACGAAGACAAGAACAGCUCGA	6	0	transposon	LTR	invader6
1250	CUGACUAAGAGGGCUGCACAGUU	6	0	transposon	LTR	HMS-Beagle
1251	GGCUCUCUCAGUCGCUUCCCGGGAA	6	0	ncRNA		rRNA
1252	ACCGAAAUGAAUUUGGCUAGCGCUA	6	0	transposon	LTR	GATE
1253	CCCACCUGAAUCUCCGCAACCCAGU	6	0	transposon	LINE	G2
1254	AAACCUGAAGAUGAUCCUGAGUCGU	6	0	transposon	LINE	Helena
1255	CGGAACGAAACGGCCUUUUCUAAU	6	no match			
1256	GCGAUGAACAGAAAAGAAUCAACCGA	6	0	transposon	Unknown	RR50960
1257	AGCGAACCGGGACCAGGCCAGUGCUCU	6	no match			
1258	CCGAAAGAUGGUGAACUAUGCCUGGGCA	6	no match			
1259	AUUCCGAACAUACACCGACAAACA	6	0	transposon	LINE	Doc
1260	AAAAACCUGAAGAUGGUCCUGAGU	6	2	transposon	LINE	Helena
1261	ACAUUCAUAACCGAGCGACUGUUCAU	6	1	transposon	LTR	GATE
1262	UACCGGGGCUAAGCAUAUUACCGA	6	no match			
1263	CGUACACAUAUUUCGCAACGUGACUG	6	0	repeat		HETRP_DM
1264	GAUGCAAACUCCGAAUACCAGCA	6	no match			
1265	UGCUUGGACUACAUAUGGUUGAGGG	6	0	ncRNA		rRNA
1266	GCCCGAAUCGGAUUUCCAUAACUUCCGC	6	0	coding gene		CG31764-RA|vir-1
1267	AAUAGGGAACGUGAGCUGGGUUU	6	no match			
1268	UUCAACCGGACGGGCCUAGAGUUAUUC	6	0	coding gene		CG40635-RA|CG40635
1269	UCUCAUCGUCGUAGAACAAGCCCGC	6	1	Su(Ste)		Su(Ste)
1270	CGCUUGGGAACACCGCGUGUUGUUGGCCU	6	0	ncRNA		5SrRNA
1271	ACUGCUGCUGAGAACUCGACUGA	6	0	transposon	LTR	flea
1272	GGAUAGUCCCGAAGAAACCUAAACC	6	0	transposon	LTR	opus
1273	CUAACCAUAAACGAUGCCAGCUAGC	6	0	ncRNA		rRNA
1274	AUGACAGUUCACGAUUGAUUCCCCA	6	0	transposon	LTR	opus
1275	GCACACAGGGGAGAGCAGAUGAGAGCA	6	no match			
1276	CAAUAAACAACAAAAUCGAUCCCGAA	6	0	transposon	LTR	17.6
1277	UGCUCUCGAAUGUAUGCCCGGUAU	6	1	transposon	LINE	R1-element
1278	AUUGCUUGUACCUGAGUCGUUGCGU	6	0	transposon	LINE	Circe
1279	UCCGGGAACAUGCCACCUCCGUUGU	6	0	coding gene		CG1763-RA|nod
1280	AUACCACCCUGAACGCGCCCGAUCUCGUC	6	no match			
1281	AACUCUAAGCGGUGGAUCACUCG	6	0	ncRNA		rRNA
1282	ACGAAUGAUAAGUUGUAUGCCUCCGG	6	0	no annotation		
1283	CGCAAUCCUGGGCCUCGACGCGAAU	6	0	transposon	LINE	R1-element
1284	ACCAAAGCUUUUGGGCUCCGGGGGA	6	0	ncRNA		rRNA
1285	UGCACUUGAAUCCAAGAGAAACGCA	6	0	transposon	LTR	Idefix
1286	CCCGUAUCAAUUCCACGUUUGGAGC	6	0	transposon	LTR	3S18
1287	GUAGAGCGUCCGACGAAUACCCUGGGG	6	no match			
1288	UAUUACAUAGAUGUAGAUGGGUCGGAG	6	0	no annotation		
1289	CAGACAGAGAGGCUCAUUUACAACCA	6	0	transposon	LTR	McClintock
1290	GACGUCCAGAGCGGUUUCAACGUGG	6	0	coding gene		CG18111-RA|Obp99a
1291	ACUACCCUGAACGCGCCCGAUCUCGUC	6	no match			
1292	GGUGUCCCAAGGCCAGCUCAGUGC	6	0	ncRNA		rRNA
1293	GCACACAGAGGAGAGCAGAUGAGACA	6	no match			
1294	AUGUUUCAUCGUUAGACGGCUCGGGC	6	1	AT-chX		AT-chX-B
1295	CUGAAGUUGAGUGGUGAAUCCGUG	6	0	coding gene		CG41545-RA|CG41545
1296	UACGUAAACAACCCCAGGCUGACUC	6	0	transposon	LTR	Max
1297	UAACGACCAGACAUACGCCAUAAUC	6	0	transposon	LINE	R1
1298	GGCACGGACUUGGGGAAUCCGACUGU	6	0	ncRNA		rRNA
1299	CACUGGAAAAAUCUCCGAUUGGU	6	0	transposon	LTR	opus
1300	CGCAAACCCAAAAGGACAAUACACGA	6	1	repeat		trf
1301	CCGAAUUAAUCGCGAGGCUGCAAGA	6	0	transposon	LTR	mdg3
1302	UAGAUCUUCACCGUUGACUUGCCCC	6	0	Su(Ste)		Su(Ste)
1303	CGACUCGUUUUUAAAUUUCGCAACA	6	0	transposon	LINE	Cr1a
1304	UAGAGAGUGAUGCCGAGUAUAAAAG	6	0	repeat		trf
1305	CACGGGAAAACUUACCAGGUCCGAACA	6	0	ncRNA		rRNA
1306	CCAUCAUUCAAAGCCAGCUGGAUGCCC	6	0	transposon	LINE	G
1307	AGAGAAGACCCUCAGCCUACGAGA	6	0	coding gene		CG41521-RA|CG41521
1308	ACUUGCCGAAUCCCGGGGCCGAGGGAG	6	no match			
1309	CGUUGUGGCCGAUGAAACCCUGGUUC	6	no match			
1310	UGCUCUCGAAUGUAUGCCCCGAUAUAC	6	no match			
1311	AGUGCGGUAACGCGACCGAUCCCGGAGA	6	no match			
1312	CUGGAGGAGGCUAGCUGUAUCCCUCGGC	6	0	transposon	LTR	opus
1313	CGGUUCCUGACGAAAAUAUUUUUUU	6	0	no annotation		
1314	AUCCGUUGAAGACUACUGCCGCA	6	1	transposon	LINE	R1-element
1315	UGCAAUCGAAACGAGAAACUUCAUCCC	6	0	no annotation		
1316	GAGCGCGGGAACGAGAUGUCCGAGGUU	6	1	coding gene		CG5075-RA|CG5075
1317	GACUGCAGAUCGAGAGGUCCCCGG	6	0	ncRNA		tRNA
1318	AUGCCUGAACGCCUCUAAGGUCG	6	0	ncRNA		rRNA
1319	AUUCCCCAUGAACGAGGAAUUCCCAGU	6	no match			
1320	GUAGGGAGUUUGACUGGGGCGGUA	6	0	ncRNA		rRNA
1321	GAAACCAUCAGACGCCUCCAGAGGAA	6	0	transposon	LINE	G
1322	AAAAUUACAAAGACGACGGCCCA	6	0	transposon	LTR	springer
1323	UACAAUGAUCUGAUCGGGUUUUUC	6	0	transposon	LINE	Cr1a
1324	UGUGCUAGAAGACGAUUUUCGGACCG	6	0	transposon	LINE	Cr1a
1325	ACCCUGAACGCGCCCGAUCUCGUCUGA	6	no match			

1326	CGAGAUCAAACCGCCGAGCUGCAACAAC	6	0	repeat		(TTG)n
1327	UGCACUAAUAUAAGCCCUGACGUCC	6	0	transposon	LTR	G4
1328	UCCCUAGAAGAACAUUGAAGCGC	6	0	Su(Ste)		Su(Ste)
1329	AGGGAAAGGAGCCGAGGCGUCGAC	6	0	transposon	LTR	ninja-Dsim-like
1330	UCUCAUCGUCGUAGAACUAGCCCG	6	0	Su(Ste)		Su(Ste)
1331	AGUUCGGUCCUGAGUGCGAGCUCC	6	no match			
1332	CACGAUGAUCGAUGACGAACUACGAA	6	0	transposon	LTR	ninja-Dsim-like
1333	GGACAAAACAACAUCCCGGGAAAUGA	6	0	transposon	LINE	baggins
1334	AGGACGGUGGCCAUGGAAGUCGGAAUC	6	no match			
1335	AGCAGACAAAGGACAACGAGAGA	6	0	transposon	LINE	X
1336	GAAGAAGAGAUGCCCAGUGAGUUUGC	6	0	transposon	LTR	ninja-Dsim-like
1337	AGCAGACUCCAUCGGCAGCUAUG	6	0	coding gene		CG9936-RC|skd
1338	GAGCGCACGUGUUAGGACCCGAAAGAU	6	no match			
1339	AAGCGAAUGAUUAGAGGCCUUAGGGU	6	0	ncRNA		rRNA
1340	GGAGCCGGGCGUGGAAUGCGAGUGCCUA	6	no match			
1341	UGCUUCAUCGUUAGACGGCUCGGGC	6	1	AT-chX		AT-chX-B
1342	AUCGCUGAUGAGACACCGGGACC	6	0	transposon	LTR	accord
1343	ACCAAACCAAGGCAAACUCCGAAUACCG	6	no match			
1344	CCGUGGGUGGUGGUGCAUGGCCGUU	6	2	ncRNA		rRNA
1345	AGGUAGGGAGUUUGACUGGGGCGGUAC	6	0	ncRNA		rRNA
1346	AGCGAAUUAAUCGCGAGGCUGCAA	6	1	transposon	LTR	mdg3
1347	UCUGAAACAAGAAGAGGACGACG	6	0	transposon	Low_complexity	F-element
1348	AACGAAAGUGGACAGCGAGCGAAAGC	6	0	transposon	LTR	diver2
1349	CCGAACUUGAGACCGCCCCUGGAUUU	6	no match			
1350	ACGGGUGACGGGGAAUCAGGGUU	6	1	ncRNA		rRNA
1351	AGCUUAUCCUUUGCUUGAUGAUUCGA	6	0	ncRNA		rRNA
1352	GAGAAACUGACAAAGGCCCUUGA	6	0	transposon	LTR	gypsy8
1353	GGAGAAGCCGGCGGGAGCCCCGGGGAGA	6	no match			
1354	CCCCCAAUGAGAUGGCCUUUGGCGAGA	6	0	transposon	LTR	Quasimodo
1355	UGGAACGAAAUUGGCCUGAUUAGCGGA	6	0	transposon	LTR	GATE
1356	GCAGAGAGAUAUGGUAGAUGGG	6	0	ncRNA		rRNA
1357	GACCAUAAAGAAGCCGUCGAGA	6	0	ncRNA		rRNA
1358	GGAGCGGUGGCCGAGUGGUCGAAGG	6	no match			
1359	UGGACAACCCUGGGGAAAGUACAC	6	0	transposon	LINE	Fw2
1360	UCCGAUUUCAAGUACCAGGCGAA	6	0	repeat		RSP
1361	AUUCACCAAGCGUUGGAUUGUUCACCC	6	0	ncRNA		rRNA
1362	AAUACAUGCCGACGGGCGCUGACCCCCUU	6	no match			
1363	CUUGUUCCCCGGAUAGUUUAGUUACG	6	0	ncRNA		rRNA
1364	CGGGAUAAGGAUUGGCUCUGAAG	6	0	ncRNA		rRNA
1365	CUGGAUACUAGGUAUCGACGCCGAU	6	0	no annotation		
1366	UACAAACAGAAUUCCCGAAUGAGAG	6	0	transposon	LINE	Cr1a
1367	ACAAACAGUACACCUACUAGCCCAUGU	6	0	transposon	LINE	baggins
1368	AGGGGAUCGAAGAUGAUCAGAUACCGUC	6	no match			
1369	CCCUUGAUAAACGGUUGACUUUCGGC	6	0	transposon	LINE	Rt1b
1370	GAAGCUGUCAGUAAACUCACAAU	6	0	transposon	LINE	baggins
1371	UAUAGCUAAUUAACACAAUCCCGGGGC	6	0	ncRNA		rRNA
1372	GGCGAGCGGACGGGGAACAGCCCA	6	no match			
1373	GCGAAUGAUUAGAGGCCUUAGGGU	6	0	ncRNA		rRNA
1374	UACAUCGAAAAGCGAGUUGACGUG	6	0	transposon	LTR	Max
1375	UAUGGUUCUCUCAGUCGCUUCCCGGGA	6	1	ncRNA		rRNA
1376	GUUCGAUUCCGGAGAGGGAGCCUGAGA	6	0	ncRNA		rRNA
1377	AGGUUCGAGUCCUGCAGUUGUCGCC	6	no match			
1378	CCGAACUUGAGACCGCCCCUGGACAAC	6	0	Su(Ste)		Su(Ste)
1379	UGACGCCUUGCGUGUUACGGUUGUUUC	6	0	ncRNA		rRNA
1380	GGAACUGGGAAGGUGGGACUUGAGU	6	0	transposon	LINE	G2
1381	AAUAGGGACGGUCGGGGGCAUC	6	no match			
1382	AGAGCAUAAGGACGACGAAAGUGA	6	1	transposon	LINE	Doc3
1383	AUCCCACGGAACACCCGAGCAAUGCC	6	0	transposon	LINE	G2
1384	CAGGGACACAGCACUAUGGAGCAG	6	1	transposon	LINE	BS
1385	UGAGAUAGAUUUCCGUGAGUUCAAAA	6	0	no annotation		
1386	GACGCGAGAGGUGAAAUUCUUGGACC	6	0	ncRNA		rRNA
1387	AAGGGAACGGGCUUGGCGGAAU	6	no match			
1388	UGUGCUAGAAGACGAUUUUCGGGCCGA	6	0	transposon	LINE	Cr1a
1389	GUCUGAUGACCAAGGUUGUGCUCGAA	6	0	coding gene		CG13434-RA|Nnf1a
1390	ACAAAACAUACUAAAACACCGUUGA	6	0	transposon	LINE	RR49783
1391	UGGUUAAUUCCGAUAACGAACGAGA	6	0	ncRNA		rRNA
1392	AAAGCCUAGAGGUGAACCCGAUACAA	6	0	transposon	LTR	Quasimodo
1393	CGCCGGCGGUGGUAGGUCUGAGU	6	0	coding gene		CG5467-RA|CG5467
1394	UCCGAACGUACCACGGUGCCCCAGUGA	6	0	transposon	Low_complexity	F-element
1395	ACCCUACUGAUGAAUGUUACCGC	6	no match			
1396	UCUCAUCGUAGUAGAACAAGCCCGA	6	1	Su(Ste)		Su(Ste)
1397	AGGGGCGGUCUCAAGUUCGGCAG	6	0	Su(Ste)		Su(Ste)
1398	GCCGAACUUGACGACCCACGAUAAUCCC	6	no match			
1399	AAAGAUAAUAUUCCGAUACAUUUUCC	6	0	transposon	LINE	Circe
1400	CUGGCGAAUAAAAGUAAACGAGCAAA	6	0	transposon	Low_complexity	F-element
1401	UAGUUGCUUUCCGAGCUCGAUGGUC	6	0	no annotation		
1402	UCGAGAUGGAGAGGCUUCAGAGUGA	6	0	transposon	LTR	ninja-Dsim-like
1403	AAACCUGAAGAUGAUCCUGAGUCGUC	6	1	transposon	LINE	Helena
1404	AUGCAAAUAAAUAGCCAUUGCCGAGGGA	6	0	coding gene		CG32199-RA|CG32199
1405	UGCGCGAAGAAGGAGGACGAGAA	6	0	transposon	LINE	G2
1406	AAGGGCUGGGUCGGUCGGGCUGGGGC	6	no match			
1407	AAAUCCCGGAAGCACAUUUGCAACGC	6	0	transposon	LTR	opus
1408	GGAGAAGCCGGCGGGAGCCCCGGGGA	6	no match			

1409	GCCCCAAAGAUUUUCCCCGGCACGU	6	0	Su(Ste)		Su(Ste)
1410	GAUCGGGGUCCGGGACAGCGCAUGGCA	6	no match			
1411	AUGAGGCGCUCGGUGUCGGUGA	6	no match			
1412	GCAUUCAAUACACCGAGAUGGCCCAA	6	0	transposon	LINE	Quasimodo
1413	GGAACUGUGAGGGGAUCCUCAA	6	no match			
1414	UCUCUAAUCUAUGGGUGGUGGUGCAUG	6	1	ncRNA		rRNA
1415	CCGAUUGGUAUUAGCGAGUUUAUC	6	0	transposon	LTR	opus
1416	AACCCUGAACGCGCCCGAUCUCGU	6	no match			
1417	CCAAACUAAAGCCGACGAGAAGCGU	5	0	transposon	LTR	invader6
1418	GCGUGUCCGAUCAGACGAGAUGGCCA	5	0	transposon	LINE	F
1419	GGGAAAGAAGACCCUGUUGAGCUUGA	5	1	repeat		LSU-rRNA_Dme
1420	GUCGUGUUGAUCGGUACUUGCUGC	5	0	transposon	LTR	invader4
1421	CCAAAAGAUAAAAACGUUUACGAGCA	5	0	transposon	LTR	mdg3
1422	CAUCGAAGGAACGAUCGCAAGGCAGU	5	1	transposon	LTR	flea
1423	AUGCUCUCGAAUGUAUGCCCGAUAU	5	0	transposon	LINE	R1-element
1424	AUGGCUCUCUCAGUCGCUUCCCGGGUA	5	1	ncRNA		rRNA
1425	GUAUACCUAUACAUUACCGCUAAAGU	5	0	ncRNA		rRNA
1426	CCAGACGUAAUACUCCACUACCCGGUU	5	2	transposon	LTR	HMS-Beagle
1427	ACGAGCCGGUGACCAUAGAAGUGCCCGU	5	1	coding gene		CG16742-RA|CG16742
1428	UGAUUAGACAUUCCCGUUUCUCGCGA	5	0	transposon	LTR	rover
1429	UCCGAUAACGAACGAGACUCAAAUAUAU	5	0	ncRNA		rRNA
1430	GACGCGAUGUGAUUUCUGCCCAGU	5	2	ncRNA		rRNA
1431	GAGAGAGGACGGGAGCGGCGGGGGC	5	no match			
1432	GAGCCGUACCUGUUGGUUUGUCCCAUU	5	0	ncRNA		rRNA
1433	ACGAGAGGUCCCGGGUUCAUUUCCCG	5	2	ncRNA		tRNA
1434	AUCUUAUGGUAUUGGGUUUUGAUGCA	5	0	repeat		LSU-rRNA_Dme
1435	AAGUGUUAUUGUGGGCCGGUACU	5	1	ncRNA		rRNA
1436	CUCGCCGAAUCCCGGGGCCGAGGGA	5	no match			
1437	AUGACAAGAAACCCCGAUGUAUUCA	5	no match			
1438	GCACCGUGUACGAUUACUUCUAUGGCU	5	0	coding gene		CG2100-RA|CG2100
1439	UUGGCUAUUUCUAAUAUAUUUUCAA	5	0	transposon	DNA	1360
1440	CCAACUGUGAGGGGAUCCUCAA	5	no match			
1441	GGGUUCCGGGGGAGUAUGGUUGC	5	no match			
1442	GCGUCCGGGACUCAACUACGUGGACUA	5	0	coding gene		CG17686-RB|DIP1
1443	UGCAAACGAACGAGAGACGGCAACC	5	0	transposon	LINE	I
1444	ACCCUGAACGCGCUCGAUCUCGU	5	no match			
1445	UGAGACUAAAAACGGAUUGCGGACC	5	1	no annotation		
1446	CCUAUCUUGAGCGGCGAGGACCUUA	5	0	transposon	LINE	Cr1a
1447	CAAGGAGCACCUUCGACCGAUCUG	5	no match			
1448	UAGCUAAUUAACACAAUCCCGGGGCG	5	0	ncRNA		rRNA
1449	GUUUGACUGGGGCGGUACAUCU	5	0	ncRNA		rRNA
1450	AACACGGGAAAACUUACCAGGUCCGAAC	5	0	ncRNA		rRNA
1451	GAUGUAGAGAAACGGUUGACAUUUAU	5	0	transposon	LTR	invader3
1452	UGUGCUAGAAGACGAUUUUCGGGCC	5	0	transposon	LINE	Cr1a
1453	CCGGGAGACACCAAUGCUGGCCAAG	5	0	coding gene		pseudogene
1454	UAUGGCUCUCUCAGUCGCUUCCCGGGAA	5	0	ncRNA		rRNA
1455	UUCGAUAGUACAGGAGGAGUUGCUAGA	5	0	transposon	LINE	Doc3
1456	CAUAUCCGAGGCCCUGUAAUUGGAAU	5	0	ncRNA		rRNA
1457	ACCCUGAACGCGCCCGAUCUCGUCUG	5	no match			
1458	AACAAUUACAACCGAGGCAACAAUA	5	0	transposon	LTR	Stalker2
1459	GCACCGAGAGAUUCGCCUACGCCGGC	5	0	coding gene		CG17210-RA|scpr-B
1460	UGUGUCCCGGGACCCACCAAAUUGUGAG	5	0	transposon	LINE	baggins
1461	CCCAGCGCGCCACCCGGCUCCGUAGCUGUAA	5	no match			
1462	CAGUUCAUCAAAGCGCCUUACUGGAGU	5	1	transposon	LTR	flea
1463	UGUUACAUCGUUAGACGGCUCGGGC	5	1	AT-chX		AT-chX-B
1464	AUGUGAGGGGAUCCUCAA	5	2	no annotation		
1465	UUCGGGAUAAGGAUUGGCUCUGAAG	5	0	ncRNA		rRNA
1466	UGCCAGGCCCGUAUAACGUUAAUGAU	5	0	ncRNA		rRNA
1467	UGCUCUCGAAUGUAUGCCCGAUAC	5	1	transposon	LINE	R1-element
1468	UACCUUGGGAAUGGCUUCACAGUU	5	1	ncRNA		alphagamma-element
1469	UGCGACCGCCCCUGGACGACCCA	5	1	Su(Ste)		Su(Ste)
1470	UGCUAAGGGAUCCAGUUGCGAUAA	5	0	transposon	DNA	1360
1471	CUCGAGAAUACAAUUCCAGUGGCUGAU	5	1	no annotation		
1472	CCAGCUUGGACCGAACACUUUGAAC	5	0	Su(Ste)		Su(Ste)
1473	CAAGACCAGAACCAAUGGACGUGGAC	5	0	transposon	LINE	HMS-Beagle2
1474	AUAGAACCUGAAACCGUAUGCGUAC	5	no match			
1475	CUAACUGUGAGGGGAUCCUCAA	5	no match			
1476	ACCCUUGAUAAACGGUUGACUUUCA	5	0	transposon	LINE	BS
1477	GGUCAACUGUGAGGGGAUCCUCAA	5	no match			
1478	GUAAAUCUUGAGAUCGGGCGUUC	5	no match			
1479	CCGAAGAUGACCAAGUUGGCAACCGC	5	2	transposon	LTR	gypsy6
1480	GGACACAACAACAUCCCGGGAAAUGA	5	0	transposon	LINE	Fw2
1481	UAACGACCAGACAUACGGCAUUAU	5	1	transposon	LINE	R1-element
1482	UCCGUCGAGGUGGUGGUGCGAUC	5	0	transposon	LINE	R1
1483	GCCGGGCGUGGAAUGCGAGUGCCU	5	no match			
1484	AAGAGGGACGGCCGGGGCAUUCG	5	no match			
1485	GCAAUCAAUACAGCGAGAUGGCCCAAA	5	2	transposon	LTR	Quasimodo
1486	CUGGACGUGAUCCUGAAGCCGGUGA	5	0	repeat		trf
1487	GAACUUGAGACCGCCCCUGGACGA	5	0	Su(Ste)		Su(Ste)
1488	AUCUGGUAAAGCGAAUGAUUAGA	5	0	ncRNA		rRNA
1489	UAUUAUUAAGAUGGGCUUGUAGGU	5	0	transposon	LINE	baggins
1490	GUCUGGUUAAUUCCGAUAACGAACG	5	0	ncRNA		rRNA
1491	UGUUACAAAAAAAGCUUCGUCUCGA	5	2	transposon	LTR	gypsy3

1492	UGUUUCAUCGUUAGACGACUCGGGCGU	5	2	AT-chX		AT-chX-B
1493	UAGCAAGUUAAUUGAACCUGAUCA	5	0	transposon	Unknown	RR50960
1494	ACCGAUUAAUCGCGAGGCUGCAAGA	5	no match			
1495	ACACCCUGAACGCGCCCGAUCUCGUC	5	no match			
1496	AGAACGAUGACGAUCCCACGGCUGGAG	5	0	transposon	LTR	ninja-Dsim-like
1497	GCACACAGAGGAGAGCAGAUGAGGGCA	5	no match			
1498	CGAGAAAGUACAAGAGGGUCGAU	5	0	transposon	LTR	frogger
1499	GGAUCCUCAACUGUGAGGGGACCCUCAA	5	no match			
1500	CAAGAUAUGACCAAAGCAAAGAC	5	0	transposon	LINE	Cr1a
1501	GAGCCGGGCGUGGAAUGCGAGUGCCU	5	no match			
1502	GACUAGCUGGCCGAGGAUUUGAUAU	5	0	Su(Ste)		Su(Ste)
1503	AUUGAGCGAACCCGGGGAACUGA	5	no match			
1504	UAAUAAAACAACGACCCGCUACCGGA	5	0	transposon	LTR	opus
1505	UACAUCGAAAAGCGAGUUGACGUA	5	0	transposon	LTR	invader4
1506	AUUGGCCCCUAUCCGAGAUGCGCA	5	0	no annotation		
1507	UUACCGUACGAUCAGGACAACCGU	5	0	transposon	LTR	mdg1
1508	AAAACCUGAAGAUGAUCUUGAGUCGC	5	1	transposon	LINE	Helena
1509	GCACCUGGGAAACCCAACCCUCAUGGAG	5	1	transposon	LTR	ninja-Dsim-like
1510	AUACAAACGAGGGACUAACGGAGC	5	0	coding gene		CG6141-RA|RpL9
1511	UAGUAAAUCACACACCUCUAUCGA	5	0	no annotation		
1512	GACACCGAAACAUGUGGCUGCGAG	5	0	transposon	LTR	diver2
1513	AGAACAACAGUGCAGACGGCGCCCGC	5	0	transposon	DNA	H
1514	GAAGCUGCGCGCUCGGUCGAAGAGGACGA	5	no match			
1515	GACGGACCGCAAUGCCCGCUCACA	5	0	coding gene		CG3715-RA|Shc
1516	AACAUAUGCAUUCCACAAGCCGA	5	0	transposon	LINE	Doc
1517	CAAGCAUUCGACAAAGUCUGGCUCGA	5	0	transposon	Low_complexity	F-element
1518	UGCAACGGGUAACGGGGAAUCAGGGUU	5	0	ncRNA		rRNA
1519	AACUCAAAGAAAAAGACGAUGACU	5	0	Su(Ste)		Su(Ste)
1520	UAAUUCCGAUAACGAACGAGACC	5	1	ncRNA		rRNA
1521	UCGAGAAGCGAACCCGGCGAACU	5	no match			
1522	CCGUAACUUCGGGAUAAGGAUUGCCU	5	1	ncRNA		rRNA
1523	UGUGAUGCCCUUAGAUGUCCUGG	5	0	ncRNA		rRNA
1524	GCAUCGGUGGUUCAGUGGUAGAAUGC	5	0	ncRNA		tRNA
1525	CAAGCUAGACUAUGUAUCCCGGGUAA	5	2	transposon	LINE	Doc
1526	ACCAAGAUGAUCAACUCGUGUUAUUC	5	0	transposon	LINE	Cr1a
1527	CAGGUGGACUACAAGGGCGAGAGC	5	no match			
1528	CAAGAUAUGACCAAAGCAAAGACACU	5	0	transposon	LINE	Cr1a
1529	GGGCUCCGGGGGAAGUAUGGUUGCA	5	0	ncRNA		rRNA
1530	GAUACUUUUAUCAGAGGCCUUCGAGG	5	0	transposon	LTR	HMS-Beagle
1531	ACCCUGCACGCGCCCGAUCUCGUC	5	no match			
1532	CAAACUAAUCACACCCACGCUUUGGA	5	0	transposon	LINE	INE-1
1533	AAACUGUUCACAUGAACUUGGAAUUCCC	5	0	ncRNA		rRNA
1534	ACGGGAAACCUCACCCGGCCCGGACAC	5	no match			
1535	GGUUAAUUCCGAUAACGAACGAGA	5	0	ncRNA		rRNA
1536	AUAGAUAUUAGAGAAUAUGUUGA	5	0	transposon	LTR	invader4
1537	UUCGCAAAAGUUGACCGAACUUGAU	5	0	ncRNA		rRNA
1538	AAUGAUGAUGAGCGAUAUUCAACU	5	no match			
1539	CCGAUGAACCUGAAUAUCCGUUAUGGA	5	0	ncRNA		rRNA
1540	ACCGAUAUAAUUCUAACUGAGCACCC	5	0	transposon	LTR	G4
1541	GGCGGGUGUUGACGCGAUGUGAU	5	no match			
1542	GGAGCCGGGCGUGGAAUGCGAGUGCC	5	no match			
1543	CAUCCUAGAAUCGACCGACCAGCUGU	5	0	transposon	LTR	invader3
1544	UGCUCUCGAAUGUAUGCCCGAUACU	5	2	transposon	LINE	R1-element
1545	UGCGCGAAGAAGGAGGACGAGA	5	0	transposon	LINE	G2
1546	UUCAAGGUUACCCCAGGAUGCUGUU	5	1	repeat		trf
1547	GCCGAACUUGAGACCGCCCCUGGA	5	0	Su(Ste)		Su(Ste)
1548	CUCUCCGGGAGACACCAAUGCUGGCC	5	0	coding gene		pseudogene
1549	UUAAUAACUAUAAGGCAUUUUCCGAA	5	0	transposon	DNA	hopper2
1550	AUGAACACAUAGGGGCUUCGAGG	5	2	transposon	LTR	HMS-Beagle
1551	GGGUUCCGGGGGGAUUAUGGUUG	5	no match			
1552	UAACUGCGAAGUCGAUCAGGUCCGGC	5	0	transposon	LINE	F
1553	GGACUCAUAUCCGAGGCCCUGUAAUUGG	5	0	ncRNA		rRNA
1554	GACCAUGAGAUUACACUGGAGGA	5	0	transposon	LINE	Cr1a
1555	AACGAGAGAAGCAUUUUUGCCGCA	5	0	transposon	LTR	gypsy12
1556	UUUUCUUUGAAGAUGGCCCACAUAA	5	0	repeat		trf
1557	CGAAUAGUGAUUGCGAACCGCCCGA	5	0	transposon	DNA	S
1558	GACUUCCAACUGGCCGAUAUGAA	5	0	coding gene		CG9074-RA|Mst57Da
1559	GGCACAAUACUUGUUCGAGCGAA	5	0	ncRNA		rRNA
1560	GGCUAGCUCAUUUCCCGGGAUGUU	5	1	transposon	LINE	Fw2
1561	CACAGAGUACGGGGCUUACUGA	5	0	transposon	LINE	baggins
1562	GGACUCAUAUCCGAGGCCCUGUAAUU	5	0	ncRNA		rRNA
1563	AGAGAUGACAAUGUGGCACACGC	5	0	transposon	LTR	Max
1564	ACCGUUUACACAUUUAUAUUUUUGUU	5	0	transposon	DNA	1360
1565	CCAGAACGUGUAUUGCUACCAACGGA	5	0	transposon	DNA	H
1566	GCAUUCAUAACGAGGUUGCGCAUUGC	5	0	transposon	LINE	Doc4
1567	GCAGAUCCAGAACGUGUAUUGCUACC	5	0	transposon	DNA	H
1568	CGCCGAAUCCCGGGGCCGAGGGAGCGAGA	5	no match			
1569	UAGUACAUAUUAGGGUAGUUUUUGC	5	0	transposon	DNA	1360
1570	UGCUCUCGAAUGUAUGCCCCGAUAUA	5	no match			
1571	ACCUGAUUGAUAGCGUAGACGACGA	5	0	transposon	LINE	Rt1a
1572	GUGGAUGAGAUCUCGUCACCCGUU	5	0	transposon	LINE	X
1573	UACAAACACACUUUGGAGACUUA	5	0	transposon	LTR	invader1
1574	AUGAUUCAUCGUUAGACGGCUCGGGC	5	2	AT-chX		AT-chX-B

1575	GAAACAAUCAGUGGUUUCCCUAAGC	5	0	transposon	LTR	DM88
1576	GACGAAAUAAACAAGCACUCGAGCA	5	0	transposon	Simple_repeat	I
1577	UUGAAACUGAAGCACGAAUUGUCCA	5	0	transposon	LTR	Dm88
1578	GGAUCCUCAACUGGAGGGAAUCCUCAA	5	no match			
1579	UACAUUCAGUGGCAAGGUUAACCGA	5	no match			
1580	UGUUUCAUCGUUAGACGGCUCGGGA	5	1	AT-chX		AT-chX-B
1581	AAAACCUGAAGAUGAUCCUGAG	5	0	transposon	LINE	Helena
1582	CCACAGAGUACGGGGCUUACUAAA	5	0	transposon	LINE	baggins
1583	UCAUAUAUCGAGCAACUCUCUCCC	5	0	transposon	LINE	Doc
1584	AGAGCCAUCGGACACCAUUGAGA	5	0	repeat		trf
1585	CAGAGCGAGCAGGACUGGGCGGCGG	5	no match			
1586	AGUCGAGAGUGGACGGGAGCGGC	5	no match			
1587	UGCCGGAGCGAGGAGGAGGACUAGC	5	0	Su(Ste)		Su(Ste)
1588	GACUUGAUCUCGGUGUUCAGUACACA	5	0	repeat		LSU-rRNA_Dme
1589	CGCGGCUAGACGGAAAGACCCCGU	5	no match			
1590	UGUUAUAUUAUCCUCGAGAAUGGA	5	1	transposon	LTR	Idefix
1591	CCUCAUCGUCGUAGAACAAGCCCGA	5	1	Su(Ste)		Su(Ste)
1592	GAGACCCGUCGCCGCGCUCUCCCCCCUC	5	no match			
1593	UCCGAAAGAGAAAAAUAUUCAUCGGC	5	0	transposon	LINE	HMS-Beagle2
1594	UACGACGAGAACGAAAUCAGCGCGAU	5	1	transposon	LTR	frogger
1595	GGCACAAUUACCAGAAGAUGUUGAC	5	0	transposon	LTR	roo
1596	CAAUGCUUAACGGAACACGAUCGAG	5	0	transposon	Unknown	RR50960
1597	CAAGCCAGAGAAUCCGGAGAGCG	5	no match			
1598	CGCUAGCUCAUUUCCCGGGAUGUUGU	5	0	transposon	LINE	Fw2
1599	CGCUCUUCACCACAGAGUCUCCGA	5	0	transposon	LTR	GATE
1600	CAAGCUAGACUAUGUAUCCCCGGGUAU	5	1	transposon	LINE	Doc
1601	UUAUGGCUCUCUCAGUCGCUUCCCGGGA	5	0	ncRNA		rRNA
1602	AACGAGCGCAACCCUUGUCCUUAGUUA	5	no match			
1603	GCCAGAGAACCCUCUAAUUGGGCCCGAA	5	0	transposon	LINE	baggins
1604	AAGGCCGGCGGGGAAGGUGGCUCGGG	5	no match			
1605	UGAGACUACAAACGGAUUGCGGACCGC	5	1	no annotation		
1606	GUUCCUCGGGAUCAAGGGCAACCAGU	5	0	repeat		trf
1607	UGUUUCAUCGUUAGACGGCUCGGA	5	1	AT-chX		AT-chX-B
1608	AGCGUAGUCGAUGGGAAACGGGU	5	no match			
1609	GAUGUCCUGGGCUGCACGCGCGCUAC	5	0	ncRNA		rRNA
1610	AACGAGUGAACUGGCAACAUUUCGA	5	1	Su(Ste)		Su(Ste)
1611	UGCUCUCGAACGCUUGCCCGAUAU	5	0	transposon	LINE	R1-element
1612	AGCGUAUUACCGGUGGAGUUCUUAUAU	5	0	ncRNA		rRNA
1613	GAUAGUCGGGCUUGAUUGGGAAA	5	0	ncRNA		rRNA
1614	CUACGAGGAAGCAGUGGAGGAUUGCU	5	0	transposon	LTR	ninja-Dsim-like
1615	GCACACAGAGGAGAGCAGAUGAGAGA	5	no match			
1616	CGGGAAAACUUACCAGGUCCGAA	5	0	ncRNA		rRNA
1617	AUUAUAAGGAUGAUGAUACCGAU	5	0	no annotation		
1618	ACGAGCGCACGUGUUAGGACCACGGAC	5	no match			
1619	UGAGACCGUACCCGAAACCGACACA	5	no match			
1620	UCCGAGGAACCCGUUGAUCCAGCU	5	1	no annotation		
1621	CGACUCGUUUUUAAAUUUCGCAACAAG	5	0	transposon	LINE	Cr1a
1622	GGCGGCGACGACCCAUUCGAACGUC	5	no match			
1623	GUCGGCGUCCGGUGAGCUCUCGCUGGCCC	5	no match			
1624	GACUGCAGAUCGAGAGGUCCCCGGU	5	0	ncRNA		tRNA
1625	GAUAAACAUAGGCAAGGACUCGU	5	0	transposon	LTR	GATE
1626	CGGGAUAAGGAUUGGCUCUGAA	5	0	ncRNA		rRNA
1627	CCGAACUUGAGAUCGCCCCUGGACGACC	5	0	coding gene		CG42398-RA|SteXh:CG42398
1628	ACUCGCCGAAUCCCGGGGCCGAGGGG	5	no match			
1629	ACUCGCCGAAUCCCGGGGCCGAGGGAGU	5	no match			
1630	UGGGUUCCGGGGGAGUAUGGUUGC	5	no match			
1631	CCGGUCCCGGGACAUGGCGUGGUGGGUA	5	no match			
1632	CGAAUCAACAAAUUGCUCCGAUAU	5	1	coding gene		CG41545-RA|CG41545
1633	CAACCGUGAGGGGAUCCUCAA	5	no match			
1634	GGAGAUGACCACGUUGAUAGGCU	5	no match			
1635	GGAACAAGUAGCUGCCCAGAGGCGAC	5	0	transposon	LINE	R1
1636	AAAACCCGAAGAUGAUCCUGAGUC	5	1	transposon	LINE	Helena
1637	CGCCCUAGUUCUAACCAUAAACGAUGCC	5	0	ncRNA		rRNA
1638	AGAGAACGCACCCGCUUGACCUUAC	5	0	transposon	LINE	G2
1639	CACGAUGACAUAGGACAUUACGGCA	5	0	transposon	LTR	flea
1640	AAACCUGAAGAUGAUCCUGACUCG	5	1	transposon	LINE	Helena
1641	CGCCAGCUAAGGUCCCUAAAAUUGGCU	5	no match			
1642	AACUUCGGGAUAAGGAUUGGCUCUA	5	1	ncRNA		rRNA
1643	CUGGCUCUCUCAGUCGCUUCCCGGGA	5	1	ncRNA		rRNA
1644	CUGUGAGAGAGUAGCACGACGCCAGC	5	no match			
1645	UGCACUAAUAUUAUCCCUGACGUCAGA	5	0	transposon	LTR	G4
1646	CAGUACUUUACUACUCUUGACCUC	5	0	transposon	LTR	gypsy12
1647	CACAGAGGAGAGCAGAUGAGAGCA	5	no match			
1648	AGAGAGAAGACCCUCAGCCUACGAG	5	0	coding gene		CG41521-RA|CG41521
1649	UAUGCUUCUUACUAGGGUACUUCCCGA	5	0	transposon	LINE	G2
1650	CAAUCUUAGAUAAAAACCCGAACGUG	5	0	transposon	LTR	gypsy4
1651	GCUGAGAAGACGGUCGAACUUGAC	5	no match			
1652	AAUCGCUCGAUGUCGUGAGGUGACCGC	5	0	no annotation		
1653	CCGAUCAACUUCUUUAUCAGCGAU	5	0	coding gene		CG10602-RA|CG10602
1654	UACAAUCUCCGAGCAAGACCCCGA	5	0	transposon	LINE	Circe
1655	AACUUCGAUAGCGAAAUGACUCUGU	5	0	transposon	LINE	R1-element
1656	UCCGAGGCCCUGUAAUUGGAAUGAGUAC	5	0	ncRNA		rRNA
1657	UGAGACUACAAACGGAUUGCGGUACC	5	no match			

1658	AGGUUCGAAUCCUGUCUUCCCGACCA	5	no match			
1659	GCGCCGCGGAGCCUCGGUUGGCCUCGGA	5	no match			
1660	ACUCGCCGAAUCCCGGGGCCGAGGGAGCGA	5	no match			
1661	UCUCAACUGUGAGGGGAUCCUCAA	5	no match			
1662	UAAGUUACGAACCCUCUUCUUGCGA	5	0	transposon	LTR	opus
1663	ACUGUCCCUACCUACUAUCCAGCCA	5	no match			
1664	AUAACCUUGAGACUGAGGGAGGGGC	5	0	transposon	LTR	opus
1665	AAACCAUUCGUAGACGACCUGCUUC	5	no match			
1666	GCAGACAACAUUUACGGUUUCCAUG	5	0	transposon	LINE	Cr1a
1667	AACCCAGACCGUCAGCUAAGGUCCC	5	no match			
1668	GGCCGCCCGCGAUCCGCCCAAGUGCCGU	5	0	coding gene		CG4691-RA|CG4691
1669	GUGUCCUAAGGCGAGCUCAGGGAGGAC	5	no match			
1670	AGCCCGUGGACGGUGUGAGGCCGGUAGC	5	no match			
1671	CAAAGGCAUGCCCGAUGUCCGCGAGA	5	0	transposon	LTR	opus
1672	CUCCCGGGGCUACGCCUGUCUGAGCGUCGCU	5	no match			
1673	GCUGAGAAGACGGUCGAACUUGACU	5	no match			
1674	UGAGAUAUGCUAGCAGACUGGUGA	5	1	transposon	LINE	G2
1675	GCAUUCGAGACCCAACGGCACGUGAAU	5	2	transposon	LINE	INE-1
1676	ACCCUACUGAUGAUGUGUUGUUGCCA	5	no match			
1677	UAGGCAAACUAGGUGGGUCCCGGGACA	5	0	transposon	LINE	baggins
1678	CUGCCAUCGAUGCUGACCCCCUACCCGC	5	0	transposon	LINE	R1-element
1679	GGAUCCUCAACUGUAAGGGGAUCCUCAA	5	no match			
1680	UUGGACCGUCGUAAGACUAACUUAA	5	0	ncRNA		rRNA
1681	UGAGAAGACGGUCGAACUUGACU	5	no match			
1682	UACUCUGGAAGGACGACCACGUUGU	5	0	transposon	LTR	ninja-Dsim-like
1683	UUCGGCCCGAACCCUUCUGGUGU	5	0	transposon	LINE	baggins
1684	AAUGAAACGACAGUGUCCCGCGGA	5	0	transposon	LTR	springer
1685	ACCGAAUUAAUCGCGAGGCUGCAAGACA	5	0	transposon	LTR	mdg3
1686	GACUGGGGCGGUACAUCUCUCAAAU	5	0	ncRNA		rRNA
1687	AAUAAUACAAACAGAGUUCCCGAAUGAA	5	0	transposon	LTR	copia
1688	UGCUCUCGAAUGUAUGCCCGAUAUACG	5	0	transposon	LINE	R1-element
1689	GGUGGUGCAUGGCCGUUCUUAGUUCG	5	0	ncRNA		rRNA
1690	GGAUCCUCAACUGUGACGGGAUCCUCAA	5	no match			
1691	GCGGAAAGAGAGUAAUUUUGGC	5	0	transposon	LINE	Cr1a
1692	UACGUCAACAUAUCAGGGCAGCCCGA	5	0	transposon	LINE	Doc4
1693	UAGCUAAUUAACACAAUCCCGGGG	5	0	ncRNA		rRNA
1694	CAAGCCAAUCUGGACGUACGGAA	5	0	transposon	LINE	G2
1695	GGACUCCGAUCCUAGAGCGCGGCGUGGC	5	0	coding gene		CG30178-RA|CG30178
1696	ACCCUUGAUAAACGGUUGACUUU	5	0	transposon	LINE	Rt1b
1697	AUCAAUCGCUGAUGAGACACCGGGA	5	0	transposon	LTR	accord
1698	CGCUUCCAGAACAAGUGCCUGAGGCUAA	5	0	transposon	LINE	G
1699	UCAAAUAGAAAUGGACUCAGAGU	5	0	transposon	LTR	rover
1700	UGCAAGAUAAAGACCCUUCUACCUGCGC	5	0	transposon	LINE	Helena
1701	CAUGGUUGCAACGGGUAACGGGGAAU	5	0	ncRNA		rRNA
1702	AAACCGGAGGAAGGUGGGGAUGACGUC	5	no match			
1703	AGGUGUCCCAAGGCCAGCUCAGUGC	5	0	ncRNA		rRNA
1704	UACACACUCAUCAACCAAAGGACG	5	0	no annotation		
1705	GGGGAACUGAGGAGCGCAUUUCAA	5	0	transposon	LTR	flea
1706	CGCUUCAGGAACCUAUUUCGGGCUGAAA	5	1	transposon	DNA	invader2
1707	CCGUACUUUACUUUUGCAACAAAA	5	0	transposon	DNA	1360
1708	CGAGGCCCUGUAAUUGGAAUGAGUACA	5	0	ncRNA		rRNA
1709	CGACCAAUAAAACGAUUAAGACCAGC	5	0	transposon	LTR	McClintock
1710	ACACGGGAAAACUUACCAGGUCCGAAC	5	0	ncRNA		rRNA
1711	UGUUUCAUCGUUAGACGGCUCGGGGC	5	2	AT-chX		AT-chX-B
1712	GACCAUAUCCGAAGACGCGGCGCUAGU	5	1	transposon	LINE	F
1713	AACUACUGGACCAGAUGAUAGA	5	0	transposon	Unknown	RR50960
1714	AAGAAGCGCUACGGAGAAUAUUACUC	5	0	coding gene		CG3354-RA|Mst77F
1715	CAAGAUAUGACCAGAGCAAAGCAA	5	0	Su(Ste)		Su(Ste)
1716	UACAAUCUCCGAGCACGACCCCGA	5	0	transposon	LINE	Circe
1717	GGUUCCAUGGUGUAAUGGUUAGCAC	5	0	ncRNA		tRNA
1718	CCGAACUUGAGACCGCCCCUGGACG	5	0	Su(Ste)		Su(Ste)
1719	UCCGAGCUAAUAUGGCACCAAGCUGA	5	1	no annotation		
1720	CGUUCAAUAAUGACAGAGGUACAUCU	5	0	transposon	LINE	Rt1a
1721	AUCUGUGGGAUUAUGACUGAACGC	5	no match			
1722	UCGGAGGACAUGGUGCCAAAUGUGU	5	0	coding gene		CG32164-RA|CG32164
1723	UAUGCGAGCACCCGUUCCCAGGUU	5	0	transposon	LTR	opus
1724	GAAAGAGUCAACAGAAGCCUAGG	5	0	transposon	LTR	flea
1725	CAGGAAUCAAGGUGGCUAAGAGGACC	5	0	transposon	LTR	ninja-Dsim-like
1726	GCUUCAUCCAAGAGAACUGUGCCCGA	5	0	transposon	LINE	R1-element
1727	AGCGUCCGACGAAUACCCUGGGGG	5	no match			
1728	GGGCAAAUCAUCGCCGGCGACGGGAA	5	0	coding gene		CG33936-RC|CG33936
1729	GCAUUCAAUACAGCGAGAUGGCC	5	0	transposon	LTR	Quasimodo
1730	GCCCGUAGAAAACCUUUGGAGUGC	5	0	transposon	LTR	HMS-Beagle
1731	UCGAGAUCAAGAUGUCAAAAGGGG	5	0	transposon	LTR	flea
1732	ACCCUGAACGCGCCCGAUCUCGUA	5	no match			
1733	CCAGACUUGUCAACCUUCGAUGG	5	0	transposon	LTR	gypsy2
1734	CCGGAUCGAGGACAGUGUAUGGUGGG	5	no match			
1735	GCAUUCGAGACCCAACGGCACGUGAUAU	5	2	transposon	LINE	INE-1
1736	ACCCGUUGAACCCCAUUCGUGAU	5	no match			
1737	AACGAACGAGACCUUAACCUACA	5	no match			
1738	ACCAGAGAUCUAGAGGGCCCAAUAAA	5	1	no annotation		
1739	ACUUGUUCCCCGGAUAGUUUAGUUAC	5	0	ncRNA		rRNA
1740	GCAUUCGCGUCGAGGCCCAGGAUU	5	0	transposon	LINE	R1-element

1741	ACUUGUUCCGUGACUGCGUACCUU	5	no match			
1742	CAAGAUAUGACCAAAACAAAGCAACG	5	0	Su(Ste)		Su(Ste)
1743	ACCGGGACCAGGCCAGUGCUCUUGUGA	5	no match			
1744	ACCAAGAUCAUCAACUCGUGUUA	5	0	transposon	LINE	Cr1a
1745	CACCAAGCGUUGGAUUGUUCACCCAU	4	0	ncRNA		rRNA
1746	AAAGGAAUUGACGGAAGGGCAC	4	0	ncRNA		rRNA
1747	CUGGCUCUCUCAGUCGCUUCCCGGGAA	4	1	ncRNA		rRNA
1748	AAGAACGAAAGUCGGAGGUUCGAAGAC	4	no match			
1749	UGGACAGAUAGAAAGAUUUCACUCCA	4	0	transposon	LTR	HMS-Beagle
1750	UAGCGACGUCGCUUUUUGAUCCUUCGAU	4	0	ncRNA		rRNA
1751	ACGUGGGAACGUGCCUUUAGGUUCGGA	4	no match			
1752	UCAACUGUGAGGAUCCUCAA	4	no match			
1753	CUGGUAAAGCGAAUGAUUAGAGGCCUU	4	0	ncRNA		rRNA
1754	UCCGAUCUCCUGAGUCGGACUCCGAUAU	4	2	transposon	LINE	R1
1755	CCUGACCAGAAGAACCGUAAGGCA	4	0	transposon	LTR	micropia
1756	AGAAUCAACCGAAACUGGCAGUGAGA	4	0	transposon	Unknown	RR50960
1757	CUGUGAGGUGUGAGGAUCCUCAA	4	no match			
1758	CCCGAUUUCAAGUACCAGGCGAACAGA	4	0	repeat		RSP
1759	GGGAACUGGGAAGGUGGGACUUGAGUU	4	0	transposon	LINE	G2
1760	AGCUAAACGAAAGGACUAUCCCGG	4	0	transposon	LTR	gypsy3
1761	CCUGGCGUAGAGGUUAACCGUGAU	4	0	ncRNA		snRNA
1762	GAGCGUCCGACGAAUACCCUGGGG	4	no match			
1763	UUACCGAGAAGAGGAACAGAUUAG	4	0	transposon	LTR	copia
1764	GCACACAGAGGAGACAGAUGAGAGCA	4	no match			
1765	AAAGAGGAGAAGGUGGCUUUGAGA	4	0	no annotation		
1766	UCUCAUCGUCGUAGAACAAGCCCAA	4	1	Su(Ste)		Su(Ste)
1767	AGCUUCAUCCAAGAGAACUGUGCCCGAU	4	1	transposon	LINE	R1-element
1768	UGUUUCAUCGUUAGACGGCACGGGC	4	1	AT-chX		AT-chX-B
1769	UCUCAUCGUCGUAGAACAAGCCCCGA	4	2	Su(Ste)		Su(Ste)
1770	ACCUGAUUGAUAGCGUAGACGAC	4	0	transposon	LINE	Rt1a
1771	GAAUCCUCAACUGUGAGGGGAUCCUCAA	4	no match			
1772	ACGGGAAAAGAAAAUUUGGUUGCUGA	4	1	transposon	LTR	gypsy3
1773	GGUUCCGGGGGGAGUAUGGUUGC	4	2	ncRNA		rRNA
1774	GGGGAACUGAGGAGCGCAUUUCAAA	4	0	transposon	LTR	flea
1775	UGAUCCAAUACCAUCCCGUCAAGGUGCC	4	0	coding gene		CG8209-RA|CG8209
1776	UCGAUUCCGGAGAGGGAGCCUGAG	4	0	ncRNA		rRNA
1777	CAGACACGAACGACAUUUUGAUUA	4	0	transposon	LTR	Idefix
1778	GAAGAAAUGAAUGCCUUCCUGAGCGA	4	0	transposon	LTR	Rt1c
1779	UACAUCGAGAAUCUCACGCCCACAG	4	0	transposon	LINE	Doc3
1780	CGAGAUAGUACGAAUCGGUUACGA	4	0	transposon	LINE	I
1781	ACCUGACGUAACAUUAUCCCAAACU	4	0	transposon	LINE	Circe
1782	GACCGGCAACGACGUUCCACACGC	4	0	no annotation		
1783	AUACCACCCUGAACGCGCCCGAUCUCG	4	no match			
1784	UGCAAUAUGAUGAACGUCAUCCGA	4	0	transposon	LTR	Rt1c
1785	AAAACAAAGCAUUGUGAUGGCCCU	4	0	ncRNA		rRNA
1786	AACCCUGAAGCAAUUGAUGAUGUGA	4	no match			
1787	GCGACCUCAGAUCAGACGUGGCGACC	4	no match			
1788	GUAACCCAAACGUCGAGCAUGAAGGC	4	0	transposon	LINE	R1-element
1789	GGAGAUGACAAUGUAGUGAACGCCA	4	0	transposon	LTR	invader4
1790	AUUUAUGUUAGUUACUUGUUCCCCGG	4	0	ncRNA		rRNA
1791	AACACAGCCCAACUGCGGACCCAUUAA	4	1	coding gene		CG14034-RB|CG14034
1792	ACGGAAUGGAACCGAGCUCUGAGU	4	0	transposon	LINE	X
1793	GUUCGUUUCCCGGCCGACGCACCA	4	2	ncRNA		tRNA
1794	UGUUCAUCGUUAGACGGCUCGGGCG	4	2	AT-chX		AT-chX-B
1795	GGUCGCGGGUUCGUUCCCCUCAUGGGC	4	1	ncRNA		tRNA
1796	GUCCGCGAGAUUGCCGCAAACUUAGAA	4	0	transposon	LTR	opus
1797	CGGCUCCGGGACGGCUGGGAAGGCCC	4	no match			
1798	CUGCUUCUGGGUCGGGGUUUCGUACGU	4	no match			
1799	CUGGGUAAGUCGGGACCUAAGGCGAGGC	4	no match			
1800	CCGAACUUGAGAUCGGCCCUGGACGAC	4	1	coding gene		CG42398-RA|SteXh:CG42398
1801	ACAUUCAUAACCGAGCGAUUGUUCAA	4	0	transposon	LTR	GATE
1802	GUCGUGUUGAUCGGUACUUGCUGCGC	4	0	transposon	LTR	invader4
1803	UCAUCGAGAUAUUAAGGACUGGGGAA	4	0	transposon	LTR	blood
1804	GAUAUUCUUAACCCGACAUCAGAAGUGG	4	0	transposon	LTR	micropia
1805	CCAAGUGAUAUCGUCGGCGGAUGC	4	0	no annotation		
1806	AGAGAGGCGAUGAAGGACGUGGCAAGC	4	no match			
1807	GUCGGUGGUGUAAUGGUUAGCAUAGUU	4	0	ncRNA		tRNA
1808	UACAUCGAAAAGCGAGGUGACGUGU	4	1	transposon	LTR	Max
1809	GCCCCUGAUGAAGAAGAGAUGCCCAG	4	0	transposon	LTR	ninja-Dsim-like
1810	UGACAUGAGACGUCUGGCCAUUCAA	4	0	transposon	LTR	3S18
1811	AGCUUCAUCCAAGCGAACUGUGGCC	4	0	transposon	LINE	R1
1812	GUGCACAUAUACCGAGGCCAACGAUC	4	no match			
1813	GAGGGAGACAAACAUGUUCGAGC	4	0	transposon	Unknown	RR50960
1814	CGCGAGAGGUGAAAUUCUUGGACCGUC	4	0	ncRNA		rRNA
1815	ACCCUUGAUAAACGGUUGACUUUCGGU	4	1	transposon	LINE	Rt1b
1816	UAGAACCUGAAACCGUAUGCGUAC	4	no match			
1817	ACCAAGCGUUGGAUUGUUCACCCAUGC	4	0	ncRNA		rRNA
1818	UACAUCGAGAAUCUCACGCCCACAGGC	4	0	transposon	LINE	Doc3
1819	UAGUGAGGUCUCCGGACGUGAUCAA	4	1	ncRNA		rRNA
1820	GGAUCCUCAACUGUGAGGGGAUCCUCUAA	4	no match			
1821	AACCACUUUCAUGCUUGGGAUUG	4	0	ncRNA		rRNA
1822	UUCGCUAGUCCGUCCGAUUCUUGAGUA	4	0	transposon	LINE	Cr1a
1823	CAAACGAACGAGAGACGGCAACC	4	0	transposon	LINE	I

1824	AUCGAUAUUAUCCCAGCACUACGA	4	0	no annotation		
1825	CGUGUAUCAUUUCCUAGCCAUUAUACG	4	0	repeat		LSU-rRNA_Dme
1826	UUAUGAAUCGAGUAAAAACUGCAG	4	0	transposon	LTR	diver2
1827	AACUGAAUGUGGAGACGUCGGCGCGA	4	no match			
1828	AGAGCAUAAAGACGACGAAAGUGACC	4	1	transposon	LINE	Doc3
1829	AUCAGGGUUCGAUUCCGGAGAGGGAGC	4	0	ncRNA		rRNA
1830	GCUCUCUCAGUCGCUUCCCGGGAAAC	4	0	ncRNA		rRNA
1831	GGAUCGAAGAUGAUCAGAUACCGUCGU	4	no match			
1832	CCGGGACCCACCUAAUUUUGAGUUUGC	4	1	transposon	LINE	baggins
1833	CGUCGGCGCGAGCCCUGGGAGGAGUU	4	no match			
1834	ACAUAUACUGAUAAGACAUUCCCGU	4	1	transposon	LTR	rover
1835	AACGAGAAUAUACACAGGGGCUUA	4	0	transposon	LINE	Doc3
1836	ACAAACAUCAAGAAAACCGACGACAC	4	0	transposon	LTR	INE-1
1837	CGAGCGAACCGGGACCAGGCCAGU	4	no match			
1838	UACGAAUUCAACGAGGUAUGGC	4	0	repeat		SAR_DM
1839	UCCGAAGACAAGUAUACCGUGGCACACC	4	0	transposon	LTR	micropia
1840	CUUGAAAUGAUGAUGACUAAGGUCGC	4	0	no annotation		
1841	UAAGAGGGACGGCCGGGGGCAUUCGU	4	no match			
1842	CCAGACGUAAUACUCCACUACCCGGG	4	1	transposon	LTR	HMS-Beagle
1843	AGCUUCAUCCAAGCGAACUGUGCCCU	4	1	transposon	LINE	R1-element
1844	AACAAGAUGCAUACGAUGUGGCUCCAG	4	0	transposon	DNA	1360
1845	GCCGAGCGCAAAAGUUGUACGAUACC	4	0	transposon	LTR	opus
1846	AGUCCCGCAACGAGCGCAACCCUCGCC	4	no match			
1847	CCCAGAAACACAGUGUUUGACGCCAC	4	1	coding gene		CG18743-RA|Hsp70Ab
1848	UACAUCGAGAAUCUCACGCCCACAGG	4	0	transposon	LINE	Doc3
1849	GCGGACCAUGCUAUGCCUGUGGACC	4	0	coding gene		CG17956-RA|Mst87F
1850	AUCCCACGGAACACCCGAGCAAUGCCAC	4	0	transposon	LINE	G2
1851	AACACAAUCGAUCCCGAAAUUUUGA	4	0	transposon	LTR	17.6
1852	AAGUACGAGAGACACCCUAACAAACC	4	0	transposon	LTR	opus
1853	CAACCGACUACAAAGAUGCCAUACGA	4	0	transposon	Unknown	RR50960
1854	GGCUCAACUGUGAGGGGAUCCUCAA	4	no match			
1855	AGAACGAUGACGAUCCCACGGCU	4	0	transposon	LTR	ninja-Dsim-like
1856	UUUGAACAUCGAGAAUACUGCACAGC	4	0	transposon	LINE	Doc
1857	ACACCUGACGUUUCGACCAAUUUAAUU	4	1	transposon	LTR	diver2
1858	GAAGAAGAGAUGCCCAGUGAGUUUG	4	0	transposon	LTR	ninja-Dsim-like
1859	ACCGAAUUAAUCGCGAGGCUG	4	0	transposon	LTR	mdg3
1860	CCCGCAAUAUGGCAAUCGCUCGAU	4	0	coding gene		CG40609-RA|CG40609
1861	UUCCGAUAACGAACGAGACUCAA	4	0	ncRNA		rRNA
1862	CGAAAUGAACCGUGCAUCGACAA	4	0	coding gene		CG1394-RA|CG1394
1863	UAGACAUUUACUCAUCCGAAGGCAAA	4	0	transposon	LTR	McClintock
1864	CGGGGUGGGGGACGAGCGCUGGGUU	4	no match			
1865	AGUGAGGCCCUCGGAUCGGCACCGCCGGG	4	no match			
1866	AUGUCUAUGAGAUGACCGUAAUGA	4	0	transposon	LTR	ninja-Dsim-like
1867	UCCCACGGAACACCCGAGCAAUGCC	4	0	transposon	LINE	G2
1868	AAAGAGAGUAAUUUUGGCCGUCAC	4	0	transposon	LINE	Cr1a
1869	GCGUAUAACAACUACUUCUCCGUGA	4	1	transposon	LTR	accord
1870	GAAGAAGAAAAAGAUGAUAUCGAGUC	4	0	transposon	DNA	HB
1871	AAACCGAUAUGCUAUCUUGGCUGA	4	0	transposon	LINE	Doc
1872	AGAGCAAUGGAUGACAUAUUGAGAGA	4	0	transposon	LTR	gypsy4
1873	CCACAGAGUACGGGGCUUACUGAG	4	1	transposon	LINE	baggins
1874	CCCCGUACGUGAAAUGACCUGUGUG	4	no match			
1875	CAAACCGAGAGAAGACUUUCACCCGA	4	0	coding gene		CG12717-RA|CG12717
1876	GGAUCCUCAACUGUGAGGGGAUUCUCAA	4	no match			
1877	UGUUCCAUCGUUAGACGGCUCGGGC	4	1	AT-chX		AT-chX-B
1878	ACUGUCCCUACCUACUAUCCAGCGAAAC	4	2	ncRNA		rRNA
1879	ACCCUUGAUAAACGUAUGACUUUCGG	4	0	transposon	LINE	BS
1880	UCAACCUGAGAGUUUGAUCCUGGCU	4	no match			
1881	AGAAGCGAUAUCCGAUGCCGAUUGU	4	0	transposon	LTR	diver2
1882	UGAACCAGAAAACAAGCCUCCUAGA	4	0	transposon	Low_complexity	F-element
1883	CAUUUAUGUUAGUUACUUGUUCCCCGG	4	0	ncRNA		rRNA
1884	ACACGAUUAACGAGUCAAUUUAUA	4	0	transposon	LINE	Cr1a
1885	AGCGUCGGGCGCGGUUAGUACUUA	4	0	ncRNA		5SrRNA
1886	AAUAGAAGCAGUUUGGGGGCAUUAGU	4	0	ncRNA		rRNA
1887	GUGCAUGGCCGUUCUUAGUUCGUGGA	4	0	ncRNA		rRNA
1888	AGAGCGUCCGACGAAUACCCUGGGG	4	no match			
1889	ACCUCUAGAGCCACGGCGAAAUGUGCC	4	no match			
1890	UGCUCUCGAAUGUAUGCCGAUAU	4	no match			
1891	CAGGCACCCGAAAAAGCAAGAGCUCGG	4	0	transposon	LINE	accord2
1892	AUCGCAUGGUGAACGAGGCUGAGA	4	0	coding gene		CG31359-RA|Hsp70Bb
1893	AAGACCGUAAACGAAAAGCUAAGA	4	0	transposon	LTR	Idefix
1894	CAGACACGAACGAAAUUUUGAUUAUU	4	0	transposon	LTR	Idefix
1895	CGCAAACCCAAAAUGACAAUGCACGACA	4	0	no annotation		
1896	GCACAUCCUGAGCUGCGUUUUCCUGA	4	0	no annotation		
1897	UGCUCGGGAGACACACGGCGGGUGCUA	4	no match			
1898	AUGCAAGGGAACGUGAGCUGGG	4	0	ncRNA		rRNA
1899	UAAUUCCGAUAACGAACGAGACUCAAA	4	0	ncRNA		rRNA
1900	GCCGAAGUCAUUGUGCGCCAGAGUAA	4	0	transposon	Simple_repeat	I
1901	AACGAACGAGACCUUAACCUACUAAAU	4	no match			
1902	AAAAUAAUCAAAGACAAUCUCAGCCCGA	4	0	transposon	Low_complexity	F-element
1903	UCAGUGGUAGAAUGCUCGCCUGCCACGC	4	0	ncRNA		tRNA
1904	UUAGGGCUGACCCUGGACAGAAGGC	4	0	transposon	LINE	G2
1905	GGAUCCAACUGUGAGGGGAUCCUCAA	4	no match			
1906	UCAGCCCUUGCCCGCAAAACCAAAUU	4	0	coding gene		CG34102-RA|BG642163

1907	GGAUCAACUGUGAGGGCAUCCUCAA	4	no match			
1908	UUCGAUUUCAAGUACCAGGCGAA	4	0	repeat		RSP
1909	UUAUGGCUCUCUCAGUCGCUUCCCGGG	4	0	ncRNA		rRNA
1910	CUGACCUGUAUUCGGAUAAUGGAA	4	0	transposon	LTR	3S18
1911	UUGUUCCCCGGAUAGUUUAGUUACGU	4	0	ncRNA		rRNA
1912	GAGCGCACGUGUUAGGACCCGAAAGA	4	no match			
1913	AGCAACAACGACGAUUUCAGCAAU	4	1	transposon	LINE	BS
1914	CGCAAACCCAAAAUGACAAUGCACG	4	0	no annotation		
1915	UAGCAGCACAAGGUCACUACCGAGAUCC	4	0	transposon	LTR	Quasimodo
1916	CAGACCGAUAUGCUAUCUUGGCUGA	4	1	transposon	LINE	Doc
1917	CGGGAUAAGGAUUGGCUCUGAAGAUUG	4	0	ncRNA		rRNA
1918	ACGAACGAGACUCUGGCAUGCUA	4	no match			
1919	GGCAAAUUAGAUCCGUAACUUCGGGA	4	0	ncRNA		rRNA
1920	GAGAGGCGCCGUAGUUACCACCC	4	0	no annotation		
1921	ACCCAGACCGUCAGCUAAGGUCCCAA	4	no match			
1922	CGGGACAUGGUGUCGGACCGACUUGGGC	4	1	transposon	Simple_repeat	I
1923	GACGAAAUAAACAAGCACUCGAGCAG	4	0	transposon	Simple_repeat	I
1924	UGCCGGGGCGAGGAGGAGGACUAGCU	4	0	Su(Ste)		Su(Ste)
1925	UUCACCAAGCGUUGGAUUGUUCACCC	4	0	ncRNA		rRNA
1926	AACGAGUAAUAUGAGCGCCCAGAGUGA	4	1	transposon	LTR	opus
1927	AAACUGAAUAAACGAAAUGGAUGAC	4	0	transposon	LTR	invader4
1928	UUGGGCCCGAACCCUUCUGGUGUGUU	4	0	transposon	LINE	baggins
1929	GUACUUCGGGAUAAGGAUUGGCUCU	4	2	ncRNA		rRNA
1930	UGCACCGAAAAUAUUUGCGAGGAGA	4	0	repeat		trf
1931	UACAUCGAAAAGCGAGUUGACGUAU	4	0	transposon	LTR	invader4
1932	UGGUUGCAACGGGUAACGGGGA	4	0	ncRNA		rRNA
1933	GGCGCUAAACCAUUCGUAUACGAC	4	no match			
1934	UACGUCCGUUGGAUUAUGCCUGA	4	0	ncRNA		rRNA
1935	CAGUCUAUGAAAGGUGUGUCGCCCAGC	4	1	transposon	LTR	GATE
1936	CGACAAGAUAGACAGGCGAAACCGCA	4	0	transposon	LTR	micropia
1937	ACGAUGCCAGCUAGCAAUUGGGUGUA	4	0	ncRNA		rRNA
1938	AGCCGGACGGUGGCCAUGGAAGUCGGAAU	4	no match			
1939	AACGAGUAAUAUGAGCGCCCAGAGU	4	1	transposon	LTR	opus
1940	UAGGAACGAAAGAGGCGAGGACAACGA	4	0	coding gene		CG40271-RA|CG40271
1941	GGAUCCCAAGCUGACUCCCGCCGGCCA	4	0	coding gene		CG13340-RA|CG13340
1942	CCCGUAAACCGCCAUCAGCAACACACCC	4	0	transposon	Low_complexity	F-element
1943	CGAGAUGACAACCUUGGACCUGACCA	4	0	transposon	LTR	micropia
1944	CAAAUUGACACCGUAACUUCGGAAGAAGG	4	no match			
1945	ACCGAUAUAAUUCUAACUGAGCACC	4	0	transposon	LTR	G4
1946	GCGGACGAGGCGUUCCGAGCUGCCCA	4	0	coding gene		CG4836-RA|CG4836
1947	CCCGAAAGCAGACGAGUUAACGUACC	4	1	transposon	Low_complexity	F-element
1948	AAUAGGGACGGUCGGGGGCAUCA	4	no match			
1949	ACCAAUGAGACCCCUACUUUACCAUA	4	0	transposon	DNA	1360
1950	AACUUAAUAAACGAAAUGGAUGACA	4	1	transposon	LTR	invader4
1951	UCCAAUAUCACCGCGGAGACUGUCG	4	0	transposon	LINE	Doc4
1952	UGCGUCCCGCAGAUUAGUGAGCCUUGU	4	0	transposon	LTR	opus
1953	UGCAAUAGAAACGCGCGAUGAGUU	4	0	transposon	LTR	Dm88
1954	AUGGCUCUCUCAGUCGCUUCCCGC	4	1	ncRNA		rRNA
1955	ACUGAAUAAACGAAAUGGAUGAC	4	0	transposon	LTR	invader4
1956	GUGGAAAUGAAGAAAGUACAUCGAAAU	4	no match			
1957	AAUCGCUCGAUGUCGUGAGGUGACC	4	0	no annotation		
1958	AAAGAGGGACGGCCGGGGGCAUUCGU	4	no match			
1959	GCUCGUUCUGGGUAAAUAGAGUUUCU	4	0	ncRNA		rRNA
1960	UGUUUCAUCGUUAGACGGCUCGGGCUUACUAA	4	no match			
1961	CGCCGAAUCCCGGGGCCGAGGGAGCGA	4	no match			
1962	UGCGAAAUAAUUGCCAGAGUCACCGG	4	0	transposon	LTR	Quasimodo
1963	CUGCCCAGUGCUCUGAAUGUCAAAG	4	0	ncRNA		rRNA
1964	GAAGACGUUCAUUACGGACAUGGGA	4	0	transposon	LINE	F
1965	UCAGACGAGAUGGCCAACAGAGUG	4	0	transposon	LINE	F
1966	UCCGAACGUACCACGGUCCCCCAGUG	4	1	transposon	Low_complexity	F-element
1967	UGUUCAUCGUUAGACGGCUCGGGC	4	2	AT-chX		AT-chX-B
1968	GUCGAGAGAUAUCGGAAGAGUUUU	4	0	ncRNA		rRNA
1969	CACCAAGCGUUGGAUUGUUCACCCAUG	4	0	ncRNA		rRNA
1970	CAUGGUUGCAACGGGUAACGGGGAAUC	4	0	ncRNA		rRNA
1971	AAUUCCGAUAACGAACGAGACUCAA	4	0	ncRNA		rRNA
1972	AACUGGGAAGGUGGGACUUGAGUUUG	4	0	transposon	LINE	G2
1973	CACCGCCCGUCGCUACUACCGAUUG	4	0	ncRNA		rRNA
1974	UUUCAUAAUAAGUUCCCCGACGUGCC	4	0	Su(Ste)		Su(Ste)
1975	CAAACCGAUAUGCCAUCUUGGCUGA	4	1	transposon	LINE	Doc
1976	UGUAUACCUAUACAUUACCGCUAA	4	0	ncRNA		rRNA
1977	CGCAAACCCAAAAAGACAAUACACG	4	0	repeat		trf
1978	UUGUACACACCGCCCGUCGCUACU	4	0	ncRNA		rRNA
1979	CUAAGGAGGGAUGAACCGGGCCAG	4	0	ncRNA		7SLRNA-RA
1980	GGGAAACCUCACCCGGCCCGGACACGGA	4	no match			
1981	ACCCUUGAUAAACGGUUGACUUUU	4	1	transposon	LINE	Rt1b
1982	AAAUACAUGCCGACGGGCGCUGACCCC	4	no match			
1983	UGUGAAAAGAUCCCGUUUGAAUAUGACC	4	0	transposon	DNA	S
1984	GCACACAGAGGAAGCAGAUGAGACA	4	no match			
1985	CAAGAUAUGACCAAAACAAAGCAACU	4	1	Su(Ste)		Su(Ste)
1986	UUAUAAGGAUGAUGAUACCGACUAA	4	0	no annotation		
1987	CAAGCAUUCGACAGAGUCUGGCUCGA	4	0	transposon	LINE	F
1988	GUAACAACGAAGAUGAUUCUGGCCCAG	4	0	transposon	LTR	invader4
1989	GAGAAAAUUAUGAAGCCACUUGGGA	4	0	transposon	LINE	Cr1a

1990	GCCGAACGUAGUCUCUCGAGAGC	4	0	transposon	LTR	invader1
1991	GACCAAAUCAACACAUCCGACCAG	4	0	no annotation		
1992	ACGAGCGCACGUGUUAGGACCCGAUAG	4	no match			
1993	CGACUCAGUAUCCCGUUUAUUUGG	4	0	transposon	LINE	Rt1b
1994	UCCAAGCUGGGCGAGUCCUCCGACGA	4	0	coding gene		CG3213-RA|CG3213
1995	CACCAAGCGUUGGAUUGUUCACCC	4	0	ncRNA		rRNA
1996	CCGUCGCUCGAUGAUCGUCCCAGACU	4	no match			
1997	AUACCGCCCUAGUUCUAACCAUAA	4	0	ncRNA		rRNA
1998	CAGACUCGGAACGCCUCAUUGGU	4	0	coding gene		CG31359-RA|Hsp70Bb
1999	UCGUCAAAAAGAAGAACGGAACCGACC	4	0	transposon	LTR	micropia
2000	UACCCGGAUCCCGCCUCGCUAUUCUA	4	1	coding gene		CG7056-RA|CG7056
2001	GGCACAAUUACCAGAAGAUGUGGAU	4	0	transposon	LINE	Cr1a
2002	UAAAGGAAUUGACGGAAGGGCACCACC	4	0	ncRNA		rRNA
2003	UGCUCUCGAAUGUAUGCCCGAUAUCU	4	2	transposon	LINE	R1-element
2004	UGUCACGCGGGAGACCGGGGUUCAA	4	0	ncRNA		tRNA
2005	GUAAUUGGAAUGAGUACACUUUAA	4	0	ncRNA		rRNA
2006	AGCAAUAGAAGUAUUGGUAGGCCGA	4	0	transposon	LTR	rooA
2007	AUGCCCGAUGUCCGCGAGAUUGCCGC	4	0	transposon	LTR	opus
2008	ACAGAGAGAAGACUCACCCGACGAGA	4	0	no annotation		
2009	AUCGCAUGGAGGCAACCGACAGCCA	4	0	transposon	LTR	GATE
2010	GACUCUGAAUCCUGCGAUCCGA	4	0	ncRNA		tRNA
2011	GCCCUAAUGAUGAUGACCUGCAUU	4	1	coding gene		CG40600-RA|CG40600
2012	UGCCGAACGUAGUCUCUCGAGAGCGA	4	0	transposon	LTR	invader1
2013	AGUAUCUAGGACUACCAUGGUUGCAA	4	0	ncRNA		rRNA
2014	UAAGACUUAAAGAUGAUGAACCCAU	4	0	transposon	DNA	NOF
2015	UCCGGACAAACACACAAGACCCGAUC	4	0	transposon	LTR	opus
2016	UACGAUACGGAAGAUGGCCCAAA	4	0	transposon	LINE	F
2017	CACAGAGUACGGGGCUUACUGAAGUCAC	4	1	transposon	LINE	baggins
2018	CGGGCGCGGUUAGUACUUAGAUGG	4	0	ncRNA		5SrRNA
2019	CCAGCAUAUACCCGAGAUCAUGGA	4	no match			
2020	GACUACGAUAAUAGGUGACUUACUGA	4	0	transposon	LTR	flea
2021	UAUGCUAAAAGUUAUCCUUACCCAGU	4	0	transposon	LTR	opus
2022	UACGACGAGAACGAGAUCAGCGCA	4	1	transposon	LTR	frogger
2023	UCCGAAGACAAGUAUACCGUGGCA	4	0	transposon	LTR	micropia
2024	ACUCGCCGAAUCCCGGGGCCGAGGGGGC	4	no match			
2025	AUCCAGUACACGGAGCUACCUGCGUU	4	0	transposon	LINE	G2
2026	UCCCAUAUUGUCUAGUGGUUAGGAUACC	4	0	ncRNA		tRNA
2027	UGCUCUCGAAUAUAUGCCCGAUAU	4	1	transposon	LINE	R1-element
2028	GAGGUAACGAUGGAGUUUGACGG	4	0	transposon	LTR	flea
2029	GUGUCAUUGAGAGGCGCCGUAGUUA	4	2	no annotation		
2030	UACAACCCGAUCCAUUCUAAAUAAUA	4	1	no annotation		
2031	CCAGACGUAAUACUCCACUACCCGA	4	1	transposon	LTR	HMS-Beagle
2032	CAGACCUUCCUAUCACACCCUCCCGA	4	0	transposon	LINE	G
2033	AGGGAGUUUGACUGGGGCGGUACAUC	4	0	ncRNA		rRNA
2034	CUGAUGGCAGUGUACUCAGUGGCAGA	4	0	ncRNA		snoRNA
2035	AUGAGAACGAUGACGAUCCCACGGC	4	0	transposon	LTR	ninja-Dsim-like
2036	CGCGGGCGGCCCGGGUUCGUUUCC	4	1	ncRNA		tRNA
2037	UACCGGAGGUGGCAUUGCGAACUUUA	4	1	coding gene		CG8322-RA|ATPCL
2038	GCUAGUUCCGAAAAAAUCAUUACCGA	4	0	transposon	LTR	Stalker2
2039	UGCUCUCGAAUGCUUGCCCGAUAUAC	4	0	transposon	LINE	R1-element
2040	CAUGUCAUAACAACUAAACACGAAGACCC	4	0	transposon	LTR	Quasimodo
2041	GGGUAAACCGCUGAACCACUUUCAUGCU	4	0	ncRNA		rRNA
2042	ACGAGUAAUAUGAGCGCCCAGGGU	4	1	transposon	LTR	opus
2043	UACGACGAGGACGAGAUCAGCGCGC	4	2	transposon	LTR	frogger
2044	UUUCCCGAAAUCCGAGUGAAUAUGGCAC	4	0	no annotation		
2045	GACUAGCUGGCCGAGUAUUUGAUGU	4	0	Su(Ste)		Su(Ste)
2046	UGGGUCGGGGUUUCGUACGUAGC	4	no match			
2047	UGCCAGGCCCGUAUAACGUUAAUGA	4	0	ncRNA		rRNA
2048	UAAAUCAUGACAAGGCCCAAACAAGA	4	0	transposon	LTR	springer
2049	AAGCCCGUGGACGGUGUGAGGCCGGU	4	no match			
2050	ACAACAAACAACACAUACCCGCUGGCC	4	0	transposon	LTR	GATE
2051	GCCGAACUUGAGACCGCCCCUGGACGAC	4	0	Su(Ste)		Su(Ste)
2052	UGAUCGAGCAUAUUCCAUCUUUCGGAA	4	0	transposon	LTR	Dm88
2053	UAUGAGAUGGCGUACAGGAGGCU	4	0	transposon	LTR	ninja-Dsim-like
2054	CUGCUCCGAAAACAAGGCCCGUCGU	4	1	transposon	LINE	G6
2055	UGCACAGAUAAGGCGGGUCCAAGU	4	0	transposon	LINE	TART-A
2056	GAUACCACCCUGAACGCGCCCGAUCUCGU	4	no match			
2057	CACAGAGUACGGGGCUUACUGAAA	4	1	transposon	LINE	baggins
2058	UGACAGACAUCCGAAGAAGCAGGAGC	4	0	transposon	LTR	blood
2059	CUGUUAAGAAACCACUCGAGCCGCCU	4	0	transposon	LINE	Doc4
2060	AGAUGUAACGGGGCUCAAGACGU	4	no match			
2061	AACGAGAAACUUUAUCCCAGUCGGA	4	1	no annotation		
2062	UUAUAAGGAUGAUGAUACCGAUUAAA	4	0	no annotation		
2063	AACUGAAUAAACGAAAUGGAUGACAA	4	1	transposon	LTR	invader4
2064	UGCUCUCGAAUGUAUGCCCGAUAUAUG	4	1	transposon	LINE	R1-element
2065	GGCUCUCUCAGUCGCUUCCCGGGAAAC	4	0	ncRNA		rRNA
2066	UGGUUCAUCGUUAGACGGCUCGGGC	4	1	AT-chX		AT-chX-B
2067	UGAGACUACAAACGGAUUGCGGCCC	4	1	no annotation		
2068	CAAGCGUUGGAUUGUUCACCCAUGCAA	4	0	ncRNA		rRNA
2069	ACUCGCCGAAUCCCGGGGCCGAGGGGA	4	no match			
2070	GCAUUCGAGACCCAACGACACACAUGA	4	0	transposon	LINE	Rt1b
2071	UAGCUAAUUAACACAAUCCCGGGGCGU	4	0	ncRNA		rRNA
2072	UCUCAUCGUCGUAGAACAGGCCCGA	4	1	Su(Ste)		Su(Ste)

2073	UUGUCCGACAAUUCCCGAAGCAAUGCCA	4	0	transposon	LTR	diver2
2074	UGAGACUACAAACGGGUUGCGGACC	4	1	no annotation		
2075	GCACGCAGAGGAGAGCAGAUGAGAGCA	4	no match			
2076	CCCUGAUGAAGAAGAGAUGCCCAGUGA	4	0	transposon	LTR	ninja-Dsim-like
2077	UACCAACCCAGAAUAUGACCCAGCA	4	0	transposon	LTR	McClintock
2078	AGCCGUCGAGAGAUAUCGGAAGA	4	0	ncRNA		rRNA
2079	UCCCACGGAACACCCGAGCAAUGC	4	0	transposon	LINE	G2
2080	UCCGAUAACGAACGAGACUCAAAUA	4	0	ncRNA		rRNA
2081	UUCAAGCGACGAGUUAAUGAGUCAGA	4	0	no annotation		
2082	GGAUCCUAACUCCUCAACUGUGAGGUCAA	4	no match			
2083	CCGCCUGGGGAGUACGGCCGCAAGGUUGA	4	no match			
2084	CUUGGUUCAAAGUGUUCGGUCCCAA	4	0	Su(Ste)		Su(Ste)
2085	UCGCUCGAUGUCGUGAGGUGACCG	4	0	no annotation		
2086	GCUGCGGCUGAUCCCCCGGGCCUUGCACCC	4	no match			
2087	AAUUCCAUGAAAAAGACAUUGGCCA	4	0	coding gene		CG41322-RA|CG41322
2088	GCGCUUGUGACGGCCCGUGAAAAUCC	4	no match			
2089	ACAUCAAUCGCUGAAAAGGCACCGGGA	4	0	transposon	LTR	accord
2090	CCCCGGAUAGUUUAGUUACUUAGCCAAU	4	1	ncRNA		rRNA
2091	AGCGAAACCACAGCCAAGGGAACGGGCU	4	0	ncRNA		rRNA
2092	CGAAGCUAAAACGCUGGAAGGCUCG	4	0	transposon	LTR	opus
2093	UCCGAAGUGACAGACCGAAUCGCAAUU	4	0	transposon	LTR	Quasimodo
2094	AUUCUGGUUGAUCCUGCCAGUAGUU	4	0	ncRNA		rRNA
2095	CAAUCAUCCAAGGCUGAGAUCCGG	4	0	transposon	LTR	mdg3
2096	AGGCGUAGUCGAUGGAUAACCAGUUG	4	no match			
2097	AGAUCCCCGAAUCCGGAGUGGCGGA	4	no match			
2098	AUGGAACACAUAGGGGCUUCGAG	4	0	transposon	LTR	HMS-Beagle
2099	ACCCAGGUGAUUACCACUAUUAACGAA	4	0	Su(Ste)		Su(Ste)
2100	ACCGAAAGCAGACGAGGUAACGUACC	4	1	transposon	Low_complexity	F-element
2101	CGACCAAUAAAACGAUUAAGACCAG	4	0	transposon	LTR	McClintock
2102	AAAACCUGAAGAUGAUCCUGAGUCGCG	4	1	transposon	LINE	Helena
2103	CCACAGAGUACGGGGCUUACUGAA	4	0	transposon	LINE	baggins
2104	AAGGGCUGGGUCGGUCGGGCUGGGGCGC	4	no match			
2105	AAUUACAUUAGACGAGACGGCCCUAC	4	0	transposon	LTR	opus
2106	CCAAGGUGAAGAGUCUCUAGUCGAUA	4	0	ncRNA		rRNA
2107	CAUUCAUAACCGAGCGACUGUUCAA	4	0	transposon	LTR	GATE
2108	GUGGGUUCGAUUCCUUCCAGCCGCACC	4	no match			
2109	UCCGAAGAUGACCAAGUUGGCAACCGC	4	2	transposon	LTR	gypsy6
2110	ACUCGCCGAAUCCCGGGGCCUAGGGA	4	no match			
2111	AGCUUCAUCCAAGCGAACUGUGCCCAA	4	1	transposon	LINE	R1-element
2112	GAAGAAGAAAUCGAAUUACAGCUCAA	4	0	transposon	LTR	297
2113	UCCCGGGGCUACGCCUGUCUGAGCGUCGCU	4	no match			
2114	UAGACGACUUAAUGGGAACGGGG	4	no match			
2115	GCACACAGAGGAGAGCAGAUGAGUGCA	4	no match			
2116	UAGUGUCCCGGGACCCACCAAAUUGUAU	4	no match			
2117	GGCGAAGGCGACAUACUGGCUCAUUA	4	no match			
2118	UAAUUCCGAUAACGAACGAGACUCAA	4	0	ncRNA		rRNA
2119	UGCCCUCGAAUGUAUGCCCGAUAUAC	4	1	transposon	LINE	R1-element
2120	AACUCCAGCAGAGACCGUACCCGAAACC	4	no match			
2121	ACCGAUGGCGAAGGCAGCCCCCUGGGAU	4	no match			
2122	UACUCGAAGACACCACGGUGAAAGUGG	4	0	transposon	LTR	ninja-Dsim-like
2123	GGUUCGAGUCCUGCAGCUGUCGCC	4	no match			
2124	AGGGACGGUCGGGGGCAUCAGUAUUC	4	no match			
2125	UGAGGUCUCCGGACGUGAUCACUG	4	0	ncRNA		rRNA
2126	ACAACCAUAAACACGAAACAGAAUUGC	4	0	transposon	Unknown	RR50960
2127	UCAAGCCAAAGAGGGAGCCCGGU	4	1	transposon	LTR	flea
2128	ACCAAUACUUAUGCACACCGCAUU	4	0	transposon	LTR	Dm88
2129	UGCACAUCGAUAUUUUCUCGACGGA	4	0	transposon	LTR	gypsy3
2130	GAGAGUGGACGGGAGCGGCGGGGGC	4	no match			
2131	CAGAACUUGAAGUCGCCGAGGCAGCCA	4	0	transposon	LINE	F
2132	GCGAUGGUCAGAAAGUGAUGUUGACGC	4	no match			
2133	ACUCCAUCUCAGAGCGGAGAAUUUGGAC	4	no match			
2134	AACCAUAAACGAUGCCAGCUAGCA	4	0	ncRNA		rRNA
2135	CCUAAAGUGUAACGGAGGCGCGCGA	4	no match			
2136	CGAACUCAAGCGUUCUGCCCUUCGU	4	0	coding gene		CG5366-RA|CG5366
2137	GGUGACGGCCAAAAUUACUCUCUUU	4	0	transposon	LINE	Cr1a
2138	UGGAGGCAUCGACGGUGGAGAUCGAGG	4	0	coding gene		CG1403-RA|Sep1
2139	AGAGCUAAACUCGAACACGAUUUCGG	4	0	transposon	LTR	micropia
2140	UCGAGAAAAAUUGACGAAUUGCUG	4	0	transposon	LTR	invader2
2141	CUUGUUCGAGCGAACAGUGGUAUGACG	4	0	ncRNA		rRNA
2142	ACCCUGAACGUGCCCGAUCUCGUC	4	no match			
2143	CCCGAAAGCAGACGAGGUAGCGUACC	4	0	transposon	LINE	F
2144	CGAACCGAUUAAAAAUUAAAGUGCA	4	0	no annotation		
2145	CGGAGUCGGGUUGCUUGGGAAU	4	no match			
2146	CUCCGGGGGAAGUAUGGUUGCAAAGCU	4	0	ncRNA		rRNA
2147	GAUAACACUUAUAUAUCCCCCUUUGC	4	0	no annotation		
2148	ACUCGCCUAAUCCCGGGGCCGAGGGAG	4	no match			
2149	GACACGUUCAACCAGAUGGGCCUG	4	0	Su(Ste)		Su(Ste)
2150	CCAGACGUGAUACUCCACUACCCAGAU	4	0	transposon	LTR	HMS-Beagle2265
2151	CCCAGCUCGGGGAGGUAGUGACGAA	4	0	ncRNA		rRNA
2152	ACCGUAUAGACCGUACCCGAAACCGACA	4	no match			
2153	GAAACAAUCAGUGGUUUCCCUAAG	4	0	transposon	LTR	DM88
2154	GAAGCCGGUGAUCGAUAGUUCCUC	4	0	repeat		trf
2155	GACUCAUAUCCGAGGCCCUGUAAU	4	0	ncRNA		rRNA

2156	GCGUCAGAAAACGUUACGCCGACGC	4	0	repeat		trf
2157	CCCGAAAGCAGAUGAGGUAGCGUACC	4	0	transposon	LINE	FW
2158	CAAGCCAAUCUGGACGUACGGAAUU	4	0	transposon	LINE	G2
2159	UCGCUCGAUGUCGUGAGGUGACCGC	4	0	no annotation		
2160	GGCUAGCUCAUUUCCCGGGAUGUUG	4	1	transposon	LINE	Fw2
2161	GUAGAGCGUCCGACGAAUACCCUGGGGG	4	no match			
2162	UAAACCAUUCGUAGACGACCUGCUU	4	no match			
2163	UGGGAACUGGGAAGGUGGGACUUGAGU	4	0	transposon	LINE	G2
2164	CGGGCUUGAUUGGGAAACAAUAACA	4	0	ncRNA		rRNA
2165	GGAUACUAAGCAGGGACGUUGAUGA	4	0	coding gene		CG12717-RA|CG12717
2166	UUGGGCCCGAACCCUUCUGGUGUGU	4	0	transposon	LINE	baggins
2167	GAUGGCCCUAGCGGGUGUUGACACA	4	0	ncRNA		rRNA
2168	GGAGUCUAACACGUGCGCGAGU	4	no match			
2169	UUCGAUUCCGGGCUUGCGCACCA	4	no match			
2170	GCCCAGAAGACUCACCGGGAAGCA	4	0	transposon	LINE	G2
2171	UGCCGAGAUAUGCUGCGAGUUGCCAAA	4	1	transposon	Simple_repeat	I
2172	CAACCAGAGAUCUAGAGGGCCCUAUA	4	2	no annotation		
2173	CCCCAGGAGGACAACCAUCUUCUCACC	4	0	no annotation		
2174	AUCGAAUGGAGUGCCAGAACCCGUUGG	4	0	transposon	LINE	R1-element
2175	GUUCGAGUCCUGCAGUUGUCGCCA	4	no match			
2176	AUGGCUCUCUCAGCCGCUUCCCGGGA	4	1	ncRNA		rRNA
2177	AACGAGAAUAUACACAGGGGCU	4	0	transposon	LINE	Doc3
2178	AACGGCCUCUAGUGCAGAUCUUGGU	4	no match			
2179	UGACCACACACUUCUUCCAGGACAUC	4	no match			
2180	GUGACUUCAACGCCAAACACUCCUGG	4	0	transposon	LINE	G2
2181	AACCUGAGAGUUUGAUCCUGGCUC	4	no match			
2182	AGCGAGAUAAAGCGAAAACCGGCA	4	0	transposon	LINE	G6
2183	GCCGAACGUAGUCUCUCGAGAGCGG	4	1	transposon	LTR	invader1
2184	CCGAAUUAAUCGCGAGGCUGCAAGAC	4	0	transposon	LTR	mdg3
2185	GCGUCGGUGGUGUAAUGGUCAGC	4	0	ncRNA		tRNA
2186	UGCAGGGAAAAUUUAGCUUUAGCUAAA	4	1	transposon	LTR	ninja-Dsim-like
2187	GGGAACACCGCGUGUUGUUGGCCU	4	0	ncRNA		5SrRNA
2188	AAAACCUGAAGAUGAUCCUGAGUCGCU	4	1	transposon	LINE	Helena
2189	ACCACAGAGUACGGGGCUUACUAA	4	0	transposon	LINE	baggins
2190	ACCCUACUGAUGAUGUGUUGUUG	4	no match			
2191	CUGCCGAACGUAGACUGGUCGCGAG	4	0	transposon	LTR	invader1
2192	AAUAGUGAAAUCCUGGUACAGAACCGA	4	0	transposon	LINE	Circe
2193	UGCUCUCGAGUGUAUGCCCGAUAU	4	1	transposon	LINE	R1-element
2194	UCCCCGCGGAAAGACGGAAAGACCCCAU	4	no match			
2195	UAUAGCUAAUUAACACAAUCCCGGGG	4	0	ncRNA		rRNA
2196	ACUACUACGACGGUGGACUGGUGGA	4	0	coding gene		CG6022-RA|Cchl
2197	GAGUCCAAGAGAUUUGCUCCCGAGGA	4	0	coding gene		CG31359-RA|Hsp70Bb
2198	CGGACUCUAAGACGGUGCUGAGA	4	0	transposon	LTR	ninja-Dsim-like
2199	CCCCGUACGUGAAAUGACCUGUGUGG	4	no match			
2200	UCGGCCCGAACCCUUCUGGUGUGAU	4	1	transposon	LINE	baggins
2201	AUGACAUACAGUACCAUGAAGGUGA	4	0	transposon	LTR	McClintock
2202	GCGAGGCAUGUCUGUCUUUUUCGGC	4	0	transposon	LTR	invader1
2203	UAACCGUCAUUCCAGCGCAAGUUGCCC	4	1	transposon	LTR	ninja-Dsim-like
2204	GGAUCCUCAACUGUGAGGGGCAA	4	no match			
2205	UGAUAGUUCAAAGACCCCAUAUG	4	0	transposon	LINE	Cr1a
2206	GCAUUCGAGACCCAACGGCACGUGAA	4	1	transposon	LINE	INE-1
2207	UACGGCCUGGGCUACACACGUGCUAC	4	no match			
2208	CCGGCUAGCUCAGUCGGUAGAGCAUGA	4	0	ncRNA		tRNA
2209	CGUGGGUGGUGGUGCAUGGCCGUU	4	2	ncRNA		rRNA
2210	AACCGAUAUGGUUGGACCACAAUCUUU	4	0	ncRNA		7SLRNA-RA
2211	AAAUCUCGCGCCGGGCCGUACCCAUA	4	no match			
2212	AUCGAGAGUAUGCUACACACCUGGGG	4	0	transposon	LINE	F
2213	UGCAAGGUUACCCCAGGUGCUGUGCCU	4	no match			
2214	CCGGUGCGGAGUGCCCUUCGUCCUGGGA	4	no match			
2215	GCGCGAACCAUAACGAGAGUAGUGA	4	0	transposon	LTR	Dm88
2216	AACUGAACAAACGAAAUGGAUGACA	4	1	transposon	LTR	invader4
2217	ACUAAUCGCGAAAUCGAGCGGCUGU	4	1	transposon	LINE	Doc
2218	AUGACAGUUCACGAUUGAUUCCCC	4	0	transposon	LTR	opus
2219	CUUUCGCAGAGAGAUAUGGUAGAU	4	0	ncRNA		rRNA
2220	CGGGCUCGCUCAAAAUCGAUGUGC	4	0	coding gene		CG9660-RA|toc
2221	UCCUUCUGAAGAGUUGUUGGCACG	4	0	transposon	LTR	GATE
2222	UACAAGCAAGACCACCACACCCGACGC	4	0	transposon	LTR	gypsy12
2223	UUCCUCGGGAUCAAGGGCAACGAG	4	0	coding gene		CG40635-RA|CG40635
2224	ACACGGGAAACCUCACCCGGCCCGGAC	4	no match			
2225	AGCCUGAGAGGUGAUGCGUAGCC	4	no match			
2226	GAACUGAGAACCGUUGGACCAUGU	4	0	coding gene		CG17376-RA|CG17376
2227	ACUCUCCGAAUCCCGGGGCCGAGGGAG	4	no match			
2228	ACCGAGAGGUCCGGGUAAACCGCUGAA	4	0	ncRNA		rRNA
2229	UUCGAGAACAAACGAUACUGUUCGGCCG	4	0	transposon	LINE	G2
2230	CGUUCAAGGAUCACGCGAGCUACGCCAUA	4	2	transposon	LINE	Rt1b
2231	AGAAGCGAUAUCCGAUGCCGAUU	4	0	transposon	LTR	diver2
2232	UGCCGAGAUAUGCUGCGAGUUGCCAA	4	0	transposon	Simple_repeat	I
2233	UUCCGAGGAAUCUGGGUUCAAGAUUU	4	no match			
2234	CCCUGAUGGAAGACCGAAACAGUUCUGA	4	0	ncRNA		rRNA
2235	ACCCUGAACAAUACACUACACCGCAGC	4	0	transposon	LTR	mdg3
2236	CGGACAUACUAGUUGCAAGUGGCGCGA	4	1	transposon	LINE	TART
2237	UUCAACCGGACGGGCCUAGAGUUAU	4	0	coding gene		CG40635-RA|CG40635
2238	UGUUUCGUCGUUAGACGGCUCGGGC	4	1	AT-chX		AT-chX-B

2239	UUCCGAGGAACCAAUUGAUCCAGCUG	4	0	Su(Ste)		Su(Ste)
2240	CAAACAGUACACCUACUAGCCCAUGU	4	0	transposon	LINE	baggins
2241	CAAUUCUGCAACUCGAGCGACGCCGU	4	0	transposon	LTR	Quasimodo
2242	AUCAUAUUAUCCCAGCACUACGAA	4	no match			
2243	AAUUCGGCCCGAACCCUUCUGGUG	4	0	transposon	LINE	baggins
2244	AACCGAUAUGGUUGGACCACAAUCUUUU	4	0	ncRNA		7SLRNA-RA
2245	ACCGAAAAUAUAAACGAACACUUAGA	4	0	transposon	Unknown	RR50960
2246	GAGCGGCCCGUCCCCCUCCGAGA	4	no match			
2247	GUGUUUCAUCGUUAGACGGCUCGGGC	4	1	AT-chX		AT-chX-B
2248	GAGAACGACGAGGUCCUGGUUGCC	4	0	coding gene		CG8415-RA|RpS23
2249	GGUGCAAGCUCCGAAUUGAAGCCCCAGU	4	no match			
2250	CCCUAGCUCAUUUCCCGGGAUGUUUAU	4	2	transposon	LINE	baggins
2251	UACCAUAGGGAUAACUGGCUUGUGGC	4	0	ncRNA		rRNA
2252	UAGAGAGUGAUGCCGAGUAUAAAAGU	4	0	repeat		trf
2253	GUCAUGCGAGGCGAACCGGGUGAAC	4	no match			
2254	GAUUAACGAGAUUCCUACUGUCC	4	0	ncRNA		rRNA
2255	GGAGAGGUGGCAGAGUGGUCGAAUGUA	4	no match			
2256	GCAUACGAGACCCAACGGCACGUGACA	4	1	transposon	LINE	INE-1
2257	GUAUCCUCAACUGUGAGGGGAUCCUCAA	4	no match			
2258	AUCUGUGGGAUUAUGACUGAAC	4	no match			
2259	CUGUAACUCGGGUCACUGUAACUCGGG	4	no match			
2260	CCCGAAGAAACCUAAACCAAACA	4	1	transposon	LTR	opus
2261	UAAAGAGACAAUUGACGGUACCGC	4	0	transposon	LTR	Dm88
2262	GACUUGAUCUCGGUGUUCAGUAC	4	0	repeat		LSU-rRNA_Dme
2263	GGUUCCAUGGUGUAAUGGUUAGCA	4	0	ncRNA		tRNA
2264	UGUGCUAGAAGACGAUUUUAGGCCCGA	4	0	transposon	DNA	1360
2265	AUCGGGGACUCGCCAAUCUCUUGCU	4	0	transposon	LTR	springer
2266	CCCAAAAGCAGACGAGGUAACGUACC	4	1	transposon	Low_complexity	F-element
2267	CAGGAGUUGACGACACCGGUGCGA	4	1	transposon	LTR	diver
2268	UGUUUCAUCGUUAGACGGCUCGGGCAU	4	2	AT-chX		AT-chX-B
2269	AGAGCGAACCCGGGGAACUGAAACA	4	no match			
2270	UUCCGGAGAGGGAGCCUGAGAAACGG	4	0	ncRNA		rRNA
2271	AAUCAUCCUAAUGAGUCGAACGCGU	3	0	transposon	LTR	gypsy
2272	CAAGAUAUGACCAAAACAAAGACACU	3	0	transposon	DNA	1360
2273	AGUACUGGGAUGACGUCAUAUUUUG	3	0	transposon	LTR	Max
2274	GCACACAGAGGAGAGCAGAUAGAGCA	3	no match			
2275	GGAUCCUCAACUGUGAGGGGACCUCAA	3	no match			
2276	CAGGGUUCGAUUCCGGAGAGGGAGC	3	0	ncRNA		rRNA
2277	UAACUUCGGGAUAAGGAUUGGCUCUGA	3	0	ncRNA		rRNA
2278	CGCUGGCAAGCAACUGGAAGACGGAC	3	1	repeat		trf
2279	GAGGCAAUCACCGAGAGCAUCAACGGUC	3	0	transposon	LINE	Rt1b
2280	UUGGGUUCCGGGGGGAGUAUGGUUGC	3	2	ncRNA		rRNA
2281	AAACCAAAGCUUUUGGGCUCCGGGGGGA	3	1	ncRNA		rRNA
2282	CGUCGAGAGAUAUCGGAAGAGUU	3	0	ncRNA		rRNA
2283	UGUAUCAUCGUUAGACGGCUCGGC	3	2	AT-chX		AT-chX-B
2284	AAACUGAAUAAACGAAAUGGAUGACA	3	0	transposon	LTR	invader4
2285	GCGAAAAACAAUACAGACCGCCUUGGCAA	3	1	transposon	LINE	Circe
2286	CAAAGCAAUAUUGAAACCAUGAAGAGU	3	0	transposon	LINE	Fw3
2287	CCGAUAACGAACGAGACUCUGGC	3	no match			
2288	AGGGCAUAAAGACGACGAAAGUGACA	3	1	transposon	LINE	Doc3
2289	AAGCUAACUACAGACGGGGUGA	3	0	transposon	LINE	R1
2290	GCAUGGCCGUUCUUAGUUCGUGGA	3	0	ncRNA		rRNA
2291	GCAUUCAUAACGAGGUUGCGCAUUU	3	1	transposon	LINE	Doc4
2292	GCACACAGAGGAGAGCAGGUGAGAGCA	3	no match			
2293	CUCAGUCAAAUGGCAUCGUGGAGA	3	0	transposon	LINE	Doc4
2294	UGCUGACGGGGCCCGAAAACACGA	3	0	transposon	LTR	opus
2295	GAAACCGGAACAUGUGGCUGCGAGU	3	1	transposon	LINE	accord2
2296	CUUGUUCCCCGGAUAGUUUAGUUACGU	3	0	ncRNA		rRNA
2297	UGUCCCAAGGCCAGCUCAGUGCGGACA	3	0	ncRNA		rRNA
2298	CCCCAUCCAAAGCCACAGUAUAGCGCC	3	0	transposon	LTR	GATE
2299	CAUGGUUGCAACGGGUAACGGGGA	3	0	ncRNA		rRNA
2300	GGACAAACAAGACGACACCACGA	3	0	no annotation		
2301	AUCCCCAGGACUGGAUCGCAUCGA	3	0	transposon	LINE	G2
2302	GCAUUCAAUACACCGAGAUGGCCCAAU	3	0	transposon	LINE	Quasimodo
2303	ACUCGCCGAAUCCCGGGGCCAGGGA	3	no match			
2304	CCGUACCUGUUGGUUUGUCCCAUU	3	0	ncRNA		rRNA
2305	AUUGAUAUCACGGGCAACGUCGA	3	0	transposon	LTR	gypsy6
2306	UACAUCGAAAAGCGAGUUGACGU	3	0	transposon	LTR	Max
2307	GACGAACUUAUAAGUGAAUUGUUGGC	3	0	transposon	LINE	Cr1a
2308	CAAUCUCAGAUAAAAACCCGAACGUG	3	0	transposon	LTR	invader3
2309	AGGACGAGGCCGAGGCAGGAGCA	3	0	coding gene		CG3788-RA|CG3788
2310	UUACGACGCGAGAGGUGAAAUUCU	3	0	ncRNA		rRNA
2311	AUGGUUGCAACGGGUAACGGGG	3	0	ncRNA		rRNA
2312	GUUUGACUGGGGCGGUACAUCUCUC	3	0	ncRNA		rRNA
2313	GAUGUCUAAGCCCAAACUGGUGUCC	3	0	transposon	LINE	F
2314	AUAGAAGAAAUUGCCGUUACAACUGU	3	0	transposon	LTR	G4
2315	CAGGACCAUAUCCGAAGACGCGGCGC	3	0	transposon	LINE	Rt1b
2316	UCAAGAAUAACAGGACCAUCCAAG	3	0	no annotation		
2317	GCCCGUUAUAUUUGGCUCGUCAAACGU	3	1	transposon	LTR	diver2
2318	UGAGAACGAUGACGAUCCCACGGCU	3	0	transposon	LTR	ninja-Dsim-like
2319	CCAGAUGAUAUUCCGAAAACAGCCUU	3	1	transposon	LINE	Doc4
2320	UUCAACUUUAACCCAAGAGACGUGA	3	0	transposon	DNA	transib4
2321	UAAUCAAGUAACCGAGAAACGUGUU	3	0	transposon	LTR	invader2

2322	GGGGAACUGAGGAGCGCAUUUCA	3	0	transposon	LTR	flea
2323	AUCGAAGUUAUCGACAAUCAGUCAUC	3	0	transposon	LTR	flea
2324	AAGACGGCCACUGAUUCGGAGA	3	0	coding gene		CG3061-RA|CG3061
2325	GUGGCUUUGAGAGGCGCCGUAGUUAU	3	1	no annotation		
2326	AGAGGGCGUGAAACCGUUAAGAGG	3	no match			
2327	AAAUAAUCGGAUGAGACCUUGGAGGAC	3	1	ncRNA		rRNA
2328	UGACGCCUUGCGUGUUACGGUUGUUU	3	0	ncRNA		rRNA
2329	CCCGCACGAAUGGCGUAACGAUUUCUC	3	no match			
2330	UACGAGAGGACCGGGAUGGACGCACCC	3	no match			
2331	AAGAGGGACGGCCGGGGCAUUCGU	3	no match			
2332	GACUCUUAGCGGUGGAUCACUCGGCUC	3	2	ncRNA		rRNA
2333	ACUCCGGAUGUGCUGACCCCUGCGAUUU	3	no match			
2334	CUGAAGUUGAGUGGUGAAUCCGU	3	0	coding gene		CG41545-RA|CG41545
2335	UGCUGUUUACUUCGAGGACAAGGUGA	3	0	transposon	LTR	flea
2336	CAGCCAAGAGUUCCCGUGGUAGGCAGU	3	0	ncRNA		7SLRNA-RA
2337	UUGGGAACACCGCGUGCUGUUGGCCU	3	1	ncRNA		5SrRNA
2338	AGUUAACGAUCACAAAGGCAUCACGC	3	0	no annotation		
2339	GCCGCGGAGCCUCGGUUGGCCUCGGA	3	no match			
2340	AGAUGUCCUGGGCUGCACGCGCGC	3	0	ncRNA		rRNA
2341	ACUCGCCGAAUCCCGGGCCGAGGGAGC	3	no match			
2342	CAGCUGCCAUACUGCGAUCCACUGA	3	0	ncRNA		rRNA
2343	GUUCGAUUCCGGAGAGGGAGCCUGAG	3	0	ncRNA		rRNA
2344	ACCUCAGAUCAGACGUGGCGACCCGCU	3	no match			
2345	GGAUCCUCAUCUGUGAGGCUCAA	3	no match			
2346	CAAGAUAUGACCAAAACAAAGACACC	3	1	transposon	DNA	1360
2347	UCUCAUCGUCGUAGAACAGCCCGA	3	no match			
2348	AACUGUGAGGAUCCUCAA	3	1	no annotation		
2349	AGCUAUGGUGAUGGUGAUGGUGCA	3	no match			
2350	CGAGAGGUCCGGGUAAACCGCUGAA	3	0	ncRNA		rRNA
2351	UCAGCACCAUGGACGAGAUCUCCUCCGGG	3	no match			
2352	AAUAGAAGCAGUUUGGGGGCAUUAGUA	3	0	ncRNA		rRNA
2353	ACCAAGCGUUGGAUUGUUCACCCACU	3	2	ncRNA		rRNA
2354	GACAGCGAUACCUGGGGCCUUGU	3	2	transposon	LINE	F
2355	GCAAGUCUGGUGCCAGCAGCCGCGGUA	3	0	ncRNA		rRNA
2356	UCUGUCUCCGGGCGGAGGUGGGCGGC	3	no match			
2357	CCAUCCUAAAACUGAUAGACAGCUGG	3	1	Su(Ste)		Su(Ste)
2358	GCCUUACGAAACGAAACCCGAUCAA	3	0	transposon	LTR	gypsy
2359	GACUCAUAUCCGAGGCCCUGUAAUU	3	0	ncRNA		rRNA
2360	ACUCACAAUCUCAGAUAAAAACCCGA	3	1	transposon	LTR	invader3
2361	CGGGUAGCUCAGUCGGUAGAGCAUUG	3	0	ncRNA		tRNA
2362	AAAACCUGAAGAUGAUCCUGAUUCG	3	1	transposon	LINE	Helena
2363	CCAGAUGAUAUUCCGAAAACAGCCGUC	3	0	transposon	LINE	Doc4
2364	AGUGAGGUCUCCGGACGUGAUCAC	3	0	ncRNA		rRNA
2365	AAGAGUGGCGCAGUGGAAGCGUGCU	3	1	ncRNA		tRNA
2366	UUCGAUAACAACGUCAACCUGAGGCAU	3	0	transposon	LTR	diver2
2367	AGCGUCGAACGAUUUCUGCCCACCU	3	0	transposon	LTR	Quasimodo
2368	GCGGCGAUGAGAAGCUGACCACAGCCGGC	3	0	coding gene		CG4546-RA|CG4546
2369	AGCUAGACUAUGUAUCCCCGGGUAA	3	0	transposon	LINE	Doc
2370	ACCCUUAACGCGCCCGAUCUCGU	3	no match			
2371	AGGAUCCUCAACUGUGAGGGGAUCCUCAA	3	no match			
2372	ACGGAUGCCGUGUAUCGUUUGACCUU	3	no match			
2373	GAAGAAGAGAUGCCCAGUGAGUU	3	0	transposon	LTR	ninja-Dsim-like
2374	AGCUGCCAUACUGCGAUCCACUGAAA	3	1	ncRNA		rRNA
2375	UACGAAUUCAACGAGGUAUGGCAUU	3	0	repeat		SAR_DM
2376	AAGGCGCUUGAGAGAACUCGGGU	3	no match			
2377	AUGCCUUAGACAGUCACAGUGCCCUCAC	3	0	coding gene		CG31682-RA|CG31682
2378	AAACUGUGAGGGGAUCCUCAA	3	no match			
2379	UACCCUGGUAGUCCACGCCGUAAA	3	no match			
2380	CUGGGGGAUCUUUGAGUGAUGAAG	3	1	transposon	LINE	baggins
2381	AGCAGCUACUGGAGACCGGCGUCC	3	0	coding gene		CG18568-RA|CG18568
2382	UAAUAGAAGCAGUUUGGGGGCAUUAGU	3	0	ncRNA		rRNA
2383	AAACGAGAACUUUGAAGGCCGAA	3	no match			
2384	UCCAACGAGAGAGCGGAGAGCCAGC	3	0	transposon	Unknown	RR50960
2385	CAACAAAGCACCAGGACAUGACGUCA	3	0	transposon	LINE	X
2386	AGGCGAGAAAGCAGCGUUCUGGA	3	0	coding gene		CG4918-RA|RpLP2
2387	UCAACCUGAGAGUUUGAUCCUGGCUCA	3	no match			
2388	UCUCACCUCGUAGAACAAGCCCGA	3	2	no annotation		
2389	UGGUUGCAACGGGUAACGGGGAAUC	3	0	ncRNA		rRNA
2390	UAAACCUGAAGAGAGCCCUGAAUCGU	3	0	transposon	LINE	Rt1b
2391	UACCGCCCUAGUUCUAACCAUAA	3	0	ncRNA		rRNA
2392	AGCCCCUGGACUAGAUGGUAUUCC	3	0	transposon	LINE	F
2393	CGUUAUAUUGGUUGAACUCUAGAU	3	0	ncRNA		rRNA
2394	CUUGACCAGCAUUUUCUUACAAUAUUGU	3	0	coding gene		CG10859-RA|CG10859
2395	UAGACAGAGAUAUUACCAGAAGCCACAA	3	0	transposon	LTR	gypsy4
2396	UUGAAUCGUGUCCCAAUAUCUCCUGUGA	3	0	coding gene		CG42398-RA|SteXh:CG42398
2397	UUUGUAAGGACCUGAGAACUUGUA	3	0	transposon	LINE	Doc4
2398	ACUCGAAGGACGCGAAAAGGUAGCCACU	3	1	coding gene		CG8137-RA|Spn2
2399	AUCUGUGGGAUUAUGACUGAACG	3	no match			
2400	UAGGCAAACUAGGUGGGUCCCGGGA	3	0	transposon	LINE	baggins
2401	UACGAGAGACACCCUAACAAACCAAA	3	0	transposon	LTR	opus
2402	CGGCUUGGAAGGCUGGCGGGGAAGGUGGC	3	no match			
2403	UCGACCAAUAAAACGAUUAAGACCAGC	3	0	transposon	LTR	McClintock
2404	AAACGAGUGUUAUUGUGGGCCGGU	3	0	ncRNA		rRNA

2405	AUAGAAAGUAAGCCCCAUGUAAUCGAA	3	0	transposon	LTR	micropia
2406	UCCGUGGGUGGUGGUGCAUGGCCGUUC	3	2	ncRNA		rRNA
2407	AAACCCAAGGCACCCCCAAGGCCCCG	3	no match			
2408	GGGCGGAGGCCGGCGGCGGGUGGGU	3	no match			
2409	GAUCGAGGCCCAGCCCGUGGACGGUGU	3	no match			
2410	CAACUGUGAGGGGAUCCUAUGA	3	no match			
2411	CAGAGAUCUAGAGGGCCCAAGAAA	3	0	no annotation		
2412	UGCCAGCGAGGAGGAGGACUA	3	2	coding gene		CG7776-RA|E(Pc)
2413	AGCAAUUCUAGAGAACGAAAGCGA	3	0	transposon	LTR	flea
2414	ACUCGCCGAAUCCCGGGGCCGAGGGAGUC	3	no match			
2415	GAGCGCGACGGCAAGCCCUAUUGCAAA	3	0	coding gene		CG31624-RA|CG31624
2416	GAAGCUUGACGACGAAGUAAAGGUGGC	3	0	coding gene		CG17603-RA|Taf1
2417	AGGAUCUUCAACUGUGAGGGGGAUCCUCAA	3	no match			
2418	UAUAGCGACAACCAGAGUUCCCAGU	3	0	transposon	LTR	DM88
2419	CCAGAUAAAAAUAAGAUUGUCAUACGA	3	0	transposon	LTR	Idefix
2420	UGGAGCCGGGCGUGGAAUGCGAGUGCCU	3	no match			
2421	UGGGUUCCGGGGGGGAGUAUGGUUGC	3	no match			
2422	ACCAGAACGAGAAGGUAGCCUAGA	3	0	transposon	LTR	Idefix
2423	GGUCCUCAACUGUGAGGGGAUCCUCAA	3	no match			
2424	UCCCGUAAACCGCCAUCAGCAACACC	3	1	transposon	Low_complexity	F-element
2425	UUCCGAUAACGAACGAGACUCAAAUA	3	0	ncRNA		rRNA
2426	UAUUCCGAUAAACGCCCAUUACACU	3	0	transposon	LTR	gypsy3
2427	UACGACGAGAACGAAAUCAGCGCGA	3	1	transposon	LTR	frogger
2428	AACUGAAUAAACGAAAUCGAUGACA	3	1	transposon	LTR	invader4
2429	UCCUCAACUGUGAGGGAUCCUCAA	3	no match			
2430	UCCCGUAAACCGCCACCAGCAACACAU	3	1	transposon	LINE	F
2431	CGCAUGAUUAAGAUGGAUGUCAUU	3	0	transposon	LINE	Circe
2432	CACAGAGUACGGGGCUUACUAAAGUCAU	3	1	transposon	LINE	baggins
2433	UGCAACUCGAGCGACGCCGUCUCUGCCA	3	0	transposon	LTR	Quasimodo
2434	UAAGAUAUGCUAGCAGACUGGAGA	3	1	transposon	LINE	G2
2435	ACAUUCAGUGGCAAGGUUAACCGA	3	no match			
2436	UGGAGUUCGAGGAGUGGCCCGUUUA	3	no match			
2437	CGAGAUGACAACCUUGGACCUGACC	3	0	transposon	LTR	micropia
2438	ACUACUUCCAUCCUAAAACUGAUAGAC	3	0	Su(Ste)		Su(Ste)
2439	ACGAGUAAUAUGAGCGCCCAGAGUGAUA	3	0	transposon	LTR	opus
2440	CUCAUUCCGAACAUACACCGACAAA	3	0	transposon	LINE	Doc
2441	GUAGGGAGUUUGACUGGGGCGGUAC	3	0	ncRNA		rRNA
2442	ACGGAAGACCCGCUCCCCGGCGCCGCU	3	no match			
2443	CACGAUGAUCGAUGACGAGCUACGAAG	3	0	transposon	LTR	ninja-Dsim-like
2444	AUCCCACGGAACACCCGAGCAAUGCCA	3	0	transposon	LINE	G2
2445	CUCGCCGAAUCCCGGGGGCCGAGGGAGC	3	no match			
2446	UUGUCGCCGUCGUUCACCACCUGG	3	no match			
2447	GGCUUUGUGAGGCGCCGUAGUUUU	3	no match			
2448	CUGCUUCUGGGUCGGGGUUUCAUACGU	3	no match			
2449	UGCUCUCGAAUGUAUGCCCGAAA	3	1	transposon	LINE	R1-element
2450	AUUGAAGGAAUCCCAGCACCAGCGU	3	0	transposon	LTR	flea
2451	CCCCAUCCAAAGCCACAGUACAGCGCAC	3	0	transposon	LTR	GATE
2452	AUGUCGGCUCUUCCUAUCAUUGUGAA	3	0	ncRNA		rRNA
2453	AAAGGAGUCUAACACGUGCGCGAGU	3	no match			
2454	AACGAACGAGACUCUGGCAUGCUAAC	3	no match			
2455	AGAGAUCAGAGAGUGGAUUCGAG	3	0	transposon	LTR	Quasimodo
2456	GCGUUCAACACCGCCAGAUGGCCCGU	3	1	transposon	LINE	F
2457	UCCGAAGACAACGAGGAGCAGAGC	3	0	transposon	LTR	1731
2458	UGAUACCGAGAAUGACAAUGGCUG	3	0	transposon	LTR	gypsy3
2459	ACGAGUGUUAUUGUGGGCCGGUACUAU	3	0	ncRNA		rRNA
2460	CCAACAGAUACCAGAUAAGAAUAAGAU	3	0	transposon	LTR	Idefix
2461	UACAUAAGAAAGCCGACAUCCCGACA	3	0	transposon	LTR	GATE
2462	UUUGCAAAAAACGAGAGCGAUAGAGC	3	no match			
2463	ACUCCGGAUGUGCUGACCCCUGC	3	no match			
2464	ACUGGGAAGGUGGGACUUGAGUUU	3	0	transposon	LINE	G2
2465	CGUACUCUAAUAGUUCUAAUGCCCACC	3	0	transposon	LTR	diver2
2466	UGCUCCGAAAACAAGGCCCGUCGCU	3	0	transposon	LINE	G6
2467	GCACACAGAGGGAGCAGAUGAGAGCA	3	no match			
2468	UAAUACAUAUCGAUCAUUACGGCCCAG	3	0	transposon	LTR	flea
2469	AAUAGAACCUGAAACCGUAUGCGUAC	3	no match			
2470	UCGAGGGCAUCCAGCUGGCUUUGA	3	0	transposon	LINE	G
2471	ACGUGGGCCUCUCCACCCUAUAGU	3	no match			
2472	UGCUUCAUCGUUAGACGGCUCGGGCG	3	1	AT-chX		AT-chX-B
2473	CCGAUUGUCUCCAGGCCACAUAC	3	0	transposon	LINE	accord2
2474	CCUCGUGUCCACACUGAUAACUCCG	3	0	no annotation		
2475	AAUACAUGCCGACGGGCGCUGACCCCCC	3	no match			
2476	AUGCCCGAUGUCCGCGAGAUUGCCGU	3	1	transposon	LTR	opus
2477	GACCGGCUCCGGGACGGCUGGGAAGGCCC	3	no match			
2478	UACAUGCCGACGGGCGCUGACCCCC	3	no match			
2479	CAGUCGCGAAAUUGGUAUCCCCG	3	0	transposon	LTR	diver2
2480	CGGCCUGGUGGUUCUAGCGAGGAGCC	3	no match			
2481	AACAGGAGAAGGUGGCUUUGAGA	3	1	no annotation		
2482	AACUUGAGACCCAACAACGCAACCGA	3	no match			
2483	UUCCCCAUGAACGAGGAAUUCCCAGU	3	no match			
2484	UACGACGAGAAUGAGAUCAGCGCGA	3	0	transposon	LTR	frogger
2485	GGGAAACCAAAGCUUUUGGGCUCCGGGG	3	0	ncRNA		rRNA
2486	AAGACAAUCUGGGAGGUGCCCG	3	no match			
2487	UGAGACUACAAACGGAUUGCGUACC	3	1	no annotation		

2488	UCGACCUGUAUUCAGAAAUGUAAA	3	0	repeat		SAR_DM
2489	AGCGAGUGGACUGGGCCAGGCAUGC	3	0	transposon	LTR	ninja-Dsim-like
2490	AUUGACGUUAUGCUACUCUCAGAG	3	0	transposon	LINE	Doc3
2491	AUACAUGCCGACGGGCGCUGACCCCC	3	no match			
2492	AUAUCGACGGCGGUGUUUGGCACCU	3	no match			
2493	GAGCGCACGUGUUAGGACCCGAAC	3	no match			
2494	GAUCGAAGAUGAUCAGAUACCGUCGU	3	no match			
2495	CCGAACUUGAGACCCCAACCGCUCCUC	3	1	no annotation		
2496	UAAACCGUCAUUCCAGCGCAAGUUGCC	3	0	transposon	LTR	ninja-Dsim-like
2497	AUCCUCAA	3	0	transposon	LTR	297
2498	CGGUAUGUACUAUCCGGCUGAGCCU	3	no match			
2499	AGCGCCGCGGAGCCUCGGUUGGCCUCGGAU	3	no match			
2500	CAUGACAGUUCACGAUUGAUUCCCCA	3	0	transposon	LTR	opus
2501	GAAGAGAUAACUGAGGCGCUAAAG	3	0	transposon	Low_complexity	F-element
2502	CGCAUCUUAAAGUAAAUGGCCUACGC	3	0	transposon	LINE	Cr1a
2503	CUGACAAUGACAGGGCAAUUUAUA	3	0	transposon	LTR	297
2504	GAACUGGCAACAUGUCGAGCUCCCAGA	3	no match			
2505	UGAUCUCGGAAGCUAAGCAGGGUCGGG	3	no match			
2506	AUACCUAUACAUUACCGCUAAAGU	3	0	ncRNA		rRNA
2507	GCAACUAGCGACGCUGCUAUACCC	3	0	coding gene		CG12717-RA|CG12717
2508	GCCCGAAAACACGAGAGCCGAGCGCUGGU	3	0	transposon	LTR	opus
2509	GGCUUUGAGAGGCGCCGUAGUUAC	3	0	no annotation		
2510	AGAGGCGACAUUCAUAACCGAGCGA	3	0	transposon	LTR	GATE
2511	ACACUGGAUCCUCAA	3	1	coding gene		CG9886-RA|CG9886
2512	ACCGAUCCAGACGAGGGACGAUGAC	3	0	coding gene		CG2750-RA|CG2750
2513	AGCCGCGUGCCCACCGACUACGUGCAGG	3	1	repeat		trf
2514	CCCGAUAUUAAUCUAUCCUGAUUUU	3	0	transposon	LTR	Idefix
2515	GACACAGAAAAGAAGACUCCGAAGGC	3	0	transposon	LINE	Rt1b
2516	UAGUUUACCGGAACGGUCUGGAAUGGCA	3	no match			
2517	UCCGCAUGUGAAUGUUGGCCCCAGA	3	0	no annotation		
2518	UGUUUCAUCGCUAGACGGCUCGGGCGG	3	1	AT-chX		AT-chX-B
2519	UCGGGAUAAGGAUUGGCUCUAAGGGUC	3	no match			
2520	CAGAACUUGAAGUCGCCGAGGCAG	3	0	transposon	LINE	F
2521	UACGUAAACAACCCCAGGCUGACUCCGG	3	0	transposon	LTR	Max
2522	CGAACGCUUAAUGAACACGAGGUCA	3	0	transposon	LTR	McClintock
2523	GCAUUCGAGACCCAACGGCACGUG	3	0	transposon	LINE	INE-1
2524	UCGAGGAUGAACCCGGGGUGGAGCGCGA	3	1	coding gene		CG33715-RE|Msp-300
2525	AUCCUCAACUGUGAGGGGGAUCCUCAA	3	no match			
2526	ACACUGAACGCGCCCGAUCUCGUC	3	no match			
2527	CCAACUAUAACAAGCUGAUAUCCGA	3	0	transposon	LINE	G2
2528	CACGCAGUUACCGACCGUUGUGUCAUU	3	0	transposon	LINE	Quasimodo
2529	GAAACUCCAUAAAAAUCAUUGCCGC	3	0	transposon	LTR	accord
2530	ACUGAAGUGGAGAAGGGUUUCGU	3	0	ncRNA		rRNA
2531	CACAGAGUACGGGGCUUACUAA	3	0	transposon	LINE	baggins
2532	GCCGAACUUGAGACCGCCCCUGGACGAA	3	1	Su(Ste)		Su(Ste)
2533	UCCCCGCGGAAAGACGGAAAGACCCAUG	3	no match			
2534	UCGACAGUUCCUCGGGCUUGUUCUAC	3	0	Su(Ste)		Su(Ste)
2535	ACGAGCGCACGCGUUAGGACCCGAAA	3	no match			
2536	AUAAACCUGGAUAUAGAUGAGCCGA	3	2	transposon	LTR	flea
2537	CCAUCCUAAAACUGAUAGACAGAUGG	3	0	Su(Ste)		Su(Ste)
2538	UGGCCACACUGUCUCCUCCUGAGACU	3	no match			
2539	UUAAUGCCGGAAGCGAUCCCACUGCUGA	3	0	coding gene		CG41580-RA|CG41580
2540	ACGACCAAGAGAGCUCUGAUGACGA	3	0	transposon	LINE	G6
2541	ACGAGUGUUAUUGUGGGCCGGUAC	3	0	ncRNA		rRNA
2542	GACUGUAAAUCUUGAGAUCGGGC	3	no match			
2543	ACAACAAACAACACAUACCCGCUG	3	0	transposon	LTR	GATE
2544	GGGCAACUGUGAGGGGAUCCUCAA	3	no match			
2545	CGGCCAUAGUGAUCCGGUGGUCCCG	3	no match			
2546	ACAUACUCACACACUCCCACACACACAC	3	0	repeat		(TG)n
2547	UGUCCUGGGCUGCACGCGCGCUACAAU	3	0	ncRNA		rRNA
2548	UGACUCAAGUUCAAAUAGGUUUGC	3	0	transposon	LINE	Doc3
2549	UAGAUGUCCUGGGCUGCACGCGCGCUA	3	0	ncRNA		rRNA
2550	CUUAAACGAGUGUUAUUGUGGGCCGG	3	0	ncRNA		rRNA
2551	CCGAUUAUCAAACCUUUUUUGAAGC	3	0	transposon	LTR	Quasimodo
2552	GGUUAAUUCCGAUAACGAACGAGAC	3	0	ncRNA		rRNA
2553	AACGAGAUUCCCACUGUCCCUAU	3	1	ncRNA		rRNA
2554	ACUCGAAUCAACAAAUUGCGCCGA	3	0	coding gene		CG41545-RA|CG41545
2555	UUCAAGGUUACCCCAGGAUGCUGUGCC	3	0	repeat		trf
2556	AAUAAUACAAACAGAAUUCCCGAAUGA	3	0	transposon	LINE	Cr1a
2557	GACAAAACAACAUCCCGGGAAAUGAGC	3	0	transposon	LINE	baggins
2558	UGCCGAGAGCUGAUGACGAGCUUUCUU	3	no match			
2559	UCCGAAAUGAACGUCUUCAUUGGACC	3	0	transposon	LTR	gypsy8
2560	UGGCUCCAUUUCCGGUGACCGUGUG	3	0	coding gene		CG13340-RA|CG13340
2561	UUAAUGCCGGAAGCGAUCCACUGC	3	no match			
2562	UGAGACCGCCCCUGGACGACCCACG	3	0	Su(Ste)		Su(Ste)
2563	AUCCUGAACGCGCCCGAUCUCGUC	3	no match			
2564	CGGCUCAUCUCAUCCUGAGUCGC	3	no match			
2565	ACCGUCACCAUACCCGACACUUACCCC	3	0	transposon	LTR	opus
2566	UGUUUCAUCGUUAGACGGCUCGGCU	3	2	AT-chX		AT-chX-B
2567	GGUUCGAUCCCGUUCACCUCCACC	3	no match			
2568	CCGGAUAUUACAGGCGAUUUAUUCC	3	0	transposon	LTR	HMS-Beagle
2569	CUGGGAAACGGGGCGCGGCUGGA	3	no match			
2570	GUAACCGAGAAACGUGUUGACAUCUGU	3	0	transposon	LTR	invader2

2571	ACCGACGACAGAUCUUUCAAAUGUC	3	0	ncRNA		rRNA
2572	AAGCCACUGUGAUUGAUACAACAGACAA	3	no match			
2573	AGGUGGAAAUAAAGAAAGUACA	3	no match			
2574	AAAACUGAUAGACAGAUGGACGGAG	3	0	Su(Ste)		Su(Ste)
2575	UGGGAACUCUAAUGGGACUGCCGG	3	no match			
2576	GAAAAUGGUCGAGUGGAUAUACAGAG	3	1	transposon	LTR	McClintock
2577	CCACUCCCGACCCGGGGAGGUAGUGACGAA	3	no match			
2578	AGACGUGAUACUCCUCUACCCAGU	3	0	transposon	LTR	HMS-Beagle
2579	CUGCGGGAUGAACCGAACGCCGGGUU	3	no match			
2580	GGCGGCGUGUCCGAGCGGCCGAAGGU	3	no match			
2581	CAGACCCGAUGCUGCGCUCCUACUCUAG	3	0	coding gene		CG7212-RA|cdm
2582	CCGAACUUGAGACCGCCCCUGGACGACCC	3	0	Su(Ste)		Su(Ste)
2583	GCACACAGAGAAGAGCAGAUGAGAGCA	3	no match			
2584	GACCUCAGAUCAGACGUGGCGACCC	3	no match			
2585	UGAACGCGCCCGAUCUCGUCUGAUCU	3	no match			
2586	AGAUCAGACGUGGCGACCCGCUGAA	3	no match			
2587	AACCCUAGAAGCAAUUGAUGAUGUGA	3	0	transposon	LTR	GATE
2588	GACAAUCCCAAACGCAGACAUCGAGC	3	0	transposon	Low_complexity	F-element
2589	CGCAAACCCAAAAUUACAAUGCACG	3	1	no annotation		
2590	CAGUGUUGGCUGAAGUACUCCAGG	3	1	Su(Ste)		Su(Ste)
2591	UCCUUGGAAUGUGUUGGCAAGUAGCA	3	1	transposon	LTR	ninja-Dsim-like
2592	AAAAGAGAGUAAUUUUGGCCGUCAC	3	0	transposon	DNA	FB
2593	UCCGAAGUGACAGACCGAAUCGCAA	3	0	transposon	LTR	Quasimodo
2594	CACGCGGGCGGCCCGGGUUCGUUUCCC	3	1	ncRNA		tRNA
2595	AGUAUCUAGGACUACCAUGGUUGCA	3	0	ncRNA		rRNA
2596	AUGGCUCUCUCAGUCGCUUCCCGGGU	3	1	ncRNA		rRNA
2597	GCCCGAAAAAACGACUUGGUUGACACU	3	0	transposon	LTR	gypsy
2598	AGCGUUGGAUUGUUCACCCAUGC	3	0	ncRNA		rRNA
2599	GCGCGGGUAAACGGCGGGAGUAA	3	1	ncRNA		rRNA
2600	ACGUCGGCUACCCACCCGACCCGUC	3	no match			
2601	GCAACGGGUAACGGGGAAUCAGGGUU	3	0	ncRNA		rRNA
2602	AUGUACAAUGCUGCCGAGUUUGCCAA	3	0	transposon	LINE	G2
2603	UGCUCAUUCCGAACAUACACCGACGA	3	1	transposon	LINE	Doc
2604	UUUGUCAAUAAACAAGAUUCCCGAA	3	0	transposon	LTR	micropia
2605	UGACGUGCAUCAUUCCGUUGUGCGAAA	3	0	coding gene		CG13340-RA|CG13340
2606	UGCUCUCGAAUGUAUGCCCGAUAUUUAU	3	no match			
2607	ACUGAUCACAUUGCCUUCGUGUUA	3	0	transposon	LINE	Cr1a
2608	UAAGAAGAUGUCCGGACAGCUGGGC	3	0	Su(Ste)		Su(Ste)
2609	AAAACCUGUAGAUGAUCCUGAGUCGC	3	1	transposon	LINE	Helena
2610	GCGAAUGGCAUCCGGAUGAGAAAU	3	0	transposon	LTR	flea
2611	UGGGCUCCGGGGGAAGUAUGGUUGCA	3	0	ncRNA		rRNA
2612	ACCGAGCCUGGUGAUAGCUGGUUGUC	3	no match			
2613	CGAGGCCCUGUAAUUGGAAUGAGUAC	3	0	ncRNA		rRNA
2614	UGAUCCAAUACCAUCCCCGUCAAGGUGCC	3	no match			
2615	CGUAACAAGGUUUCCGUAGGUGAACC	3	0	ncRNA		rRNA
2616	UGGCUCCCUCAGUCGCUUCCCGGGAAA	3	1	ncRNA		rRNA
2617	AAUUACAUUAGACGAGACGGCCCUACA	3	0	transposon	LTR	opus
2618	UCAAGCAGGCGUUCGACAGAGUGU	3	0	transposon	LINE	G2
2619	UGCUCUCGAAUGUAUGCCCGAU	3	0	transposon	LINE	R1-element
2620	AGGGACGGUCGGGGGCAUCAGU	3	2	no annotation		
2621	AUGGUUUCAACGGGUAACGGGGAAUCA	3	1	ncRNA		rRNA
2622	CCGUUACUCAGCUAGUAUAAUUGC	3	0	transposon	LINE	INE-1
2623	UACGAACAGAAUACGAGUUGAUGG	3	0	no annotation		
2624	GCCGAGCUGUGACCAUCGAGCUAG	3	0	transposon	LINE	R1-element
2625	GAAGACGAAAUCCAAACUGACAAUGGGA	3	0	transposon	LINE	Porto1
2626	GCGGGUGUUGACGCGAUGUGAUUU	3	2	ncRNA		rRNA
2627	UAGUCCCGCAAAUCUUAGUCGAGUAAA	3	no match			
2628	GACAAACUUACCAUCGAGAUACCACC	3	0	transposon	LTR	accord
2629	UGUUUCAUCGUUAGACGGCUCGGGG	3	1	AT-chX		AT-chX-B
2630	AAAAUUACAAAGACGACGGCCCAC	3	0	transposon	LTR	springer
2631	GUUCGAGAAAAAUUGACGAAUUGCUGA	3	0	transposon	LTR	invader2
2632	GGGCUGGGUCGGUCGGGCUGGGGCGC	3	no match			
2633	CAGACAUCUACUUCCCCCCGCUCC	3	0	transposon	LTR	invader1
2634	GCAGCAACACCCGAACGUGCGAGU	3	0	coding gene		CG17736-RA|schuy
2635	GCAACCACGACUCCUCGGUAAACUCGGU	3	0	coding gene		CG6773-RA|sec13
2636	CUCAGCGAUGUAUGGGGCAAGUCAAC	3	0	coding gene		CG42398-RA|SteXh:CG42398
2637	CACAUCAACAGAAAGCGUAUCCAGGCC	3	0	transposon	LINE	G2
2638	UGCUCUCGAAUGUAUGCCCGACAUAC	3	1	transposon	LINE	R1-element
2639	UAACAAGAUGCAUACGAUGUGGCUC	3	0	transposon	DNA	1360
2640	GCACACAGAGGAGAGCAUACGCCGGU	3	no match			
2641	ACCCUGAACGCGCCCGAUCUCU	3	no match			
2642	AUCGAGGAGCUCAUUCGGAGUCAGGAC	3	0	coding gene		CG14039-RA|qtc
2643	GACACAGAGGGAGCAGAUGAGAGCA	3	no match			
2644	UAACAAGAUGCAUACGAUGUGGCUCC	3	0	transposon	DNA	1360
2645	CACCAAGAUCAUCAACCCGUGUUA	3	0	transposon	LINE	Cr1a
2646	CCGAACUCGACGCGGUUGUCAUGAA	3	0	no annotation		
2647	UACACACUCAUCAACCAAAGGACGUAG	3	0	no annotation		
2648	CAAGCUAAAACGUGGAUUCCCGCUGGA	3	0	transposon	LINE	R1
2649	CGUGAAGCGAACCCGGUGAACUGAA	3	no match			
2650	AGCUGCCAUACUGCGAUCCACUGAAG	3	0	ncRNA		rRNA
2651	AGAGCAUAAACCCAUACCACGACGAA	3	no match			
2652	GAUUCCGCAGUGCGUUGAUAUGGC	3	0	Su(Ste)		Su(Ste)
2653	GCAUUCGAGACCCAACGACACACAUG	3	0	transposon	LINE	Rt1b

2654	UGUUCCAUCGUUAGACGGCUCGGG	3	1	AT-chX		AT-chX-B
2655	AGCUUCAUCCAAGAGAACUGUGCCC	3	0	transposon	LINE	R1-element
2656	UACAAGCUGAGCUAGACCACUGUGU	3	0	transposon	LINE	RR49783
2657	UUGAAAAUCCGGGGGAGAGGGUGU	3	no match			
2658	UGCCUCAACAUUUGGUGACCUCGACGU	3	0	no annotation		
2659	GGAGAAGAUAGCCACCUCGGAGU	3	0	coding gene		CG7832-RA|l(3)L1231
2660	AUUCCGUGGGUGGUGGUGCAUGGCCGU	3	no match			
2661	UAGGACUCCCGGGGGAAACAUGGUC	3	0	transposon	Low_complexity	F-element
2662	UAAUGUCCCUUAAAUCAGCACUCG	3	0	no annotation		
2663	GCCGGGGCGAGGAGGAGGACUAGCUG	3	0	Su(Ste)		Su(Ste)
2664	CAUGUCAUAACAACUAAACACGAAGACC	3	0	transposon	LTR	Quasimodo
2665	GGCUUUGGAGGCGCCGUAGUUACCACC	3	no match			
2666	AGCGUCCGACGAAUACCCUGGGGGG	3	no match			
2667	CGGCAGGGUUCGUGACUGGGGUG	3	no match			
2668	AAAGGAAUUGACGGAAGGGCACC	3	0	ncRNA		rRNA
2669	GACCGAGCCAGAUGUACCAGUUCCU	3	0	coding gene		CG30488-RA|CG30488
2670	UAGGACAAUGAUGCGACCCGUCUCUA	3	0	transposon	LTR	ninja-Dsim-like
2671	AUCCUCAACUGUGAGGGGAUCAUCAA	3	no match			
2672	AGAUGUAACGGGGCUCAAGUGA	3	no match			
2673	CAAACCGAGAGGAGACUUUCACC	3	1	coding gene		CG12717-RA|CG12717
2674	AUAGCGACAACCAGAGUUCCCAGU	3	0	transposon	LTR	DM88
2675	GCCCCGUACGUGAAAUGACCUGUG	3	no match			
2676	CGCGCAUGAAUGGAUUAACGAGAUU	3	0	ncRNA		rRNA
2677	GUAGUGACGAAAAAUAACAAUACAG	3	0	ncRNA		rRNA
2678	UAUACCGUUACCUUCACCACCUGGCGA	3	0	transposon	LINE	Doc4
2679	ACCAAGAUGAUCAACUCGUGUUAU	3	0	transposon	LINE	Cr1a
2680	AAAGGAUUGACGGAAGGGCACCACC	3	2	ncRNA		rRNA
2681	AUGCUCUCGAAUGUAUGCCCGAUAUA	3	0	transposon	LINE	R1-element
2682	GCAUUCCUGCGUAGUACGAGAGGAAC	3	0	ncRNA		rRNA
2683	GUAGCAAAAAUAACUCCGAAUUAUGCC	3	1	transposon	LINE	Cr1a
2684	UCCGCAGUGCGUUGAUAUGGCCAGC	3	0	Su(Ste)		Su(Ste)
2685	CCGAACUUGUGACCGCCCCUGGACGACC	3	1	Su(Ste)		Su(Ste)
2686	GAUGGUCCGAAAGCACCGACGCCU	3	no match			
2687	GGCUAAGGAGGGAUGAACCGGGCCAGG	3	0	ncRNA		7SLRNA-RA
2688	UGAGACUUUAUUACACCUUGGCCUA	3	0	transposon	LTR	gypsy8
2689	CAUCAAGCCUGAGAGGUGAUGCGUAGCC	3	no match			
2690	GCCCCUGAUGAAGAAGAGAUGCCCA	3	0	transposon	LTR	ninja-Dsim-like
2691	ACCGAAGAAAACAAAGCCUCUGCCAAC	3	1	transposon	LINE	F
2692	UGCUCUCGAAUGUAUGCCCGAUAUACGGA	3	0	transposon	LINE	R1-element
2693	UGAUCCAAUAAAAACUGUACACAGACG	3	0	transposon	LTR	invader6
2694	CACUGCAAUACUAAGUCGAUCCAG	3	0	transposon	LINE	Doc3
2695	GACUGAGAAGAUAUUGUACCCGUGA	3	0	coding gene		pseudogene
2696	AUCGAUAUUAUCCCAGCACUACGAAGCG	3	0	no annotation		
2697	AGCUCGAUCGUGUGACUCCGGGUAG	3	0	transposon	LINE	Doc2
2698	GCGCGCGAUGGUUAGCUCAGACCGG	3	no match			
2699	GCACACUACAUGAGUCAGUCACUUGA	3	0	transposon	LINE	Cr1a
2700	AAGGGAACGGGCUUGGCGGAAUC	3	no match			
2701	CGGGACAUCAUGCUGGCUGAUGACCAGUA	3	1	transposon	LTR	GATE
2702	UGGCUUUCUCAGUCGCUUCCCGGGA	3	1	ncRNA		rRNA
2703	UUCGGCCCGAACCCUUCUGGUGUGU	3	0	transposon	LINE	baggins
2704	UGCGCGAAGAAGGUGAACGAGAAGGC	3	1	transposon	LINE	G6
2705	CGAGGCCCUGUAAUUGGAAUGAGUA	3	0	ncRNA		rRNA
2706	GUGGCUAUGAGAGGCGCCGUAGUUUU	3	no match			
2707	GGCGGGUGUUGACGCGAUGUGAUUUC	3	no match			
2708	UUCAAGGUUACCCCCAGGAUGCUGU	3	no match			
2709	ACAAAUAGGACAAUUAAUUCGAGA	3	no match			
2710	AUGGGUAUUUACUCGACGCGUUCCGGA	3	0	coding gene		CG11598-RB|CG11598
2711	UCACACAGAGGAGAGCAGAUGAGAGCA	3	no match			
2712	UGGGACAUGUGCUUUUAUUAGGCUAAA	3	0	ncRNA		rRNA
2713	AGGGACAUUAUUGUGGAAAUGUGACG	3	no match			
2714	UAAUACAUGCCGACGGGCGCUGACCCCC	3	no match			
2715	CCACUUGUCGUGCUGUCGCUGGUCUGA	3	1	coding gene		CG32351-RA|CG32351
2716	GAUCCUCAACUGUGAGGGGGAUCCUCAA	3	no match			
2717	AAGACAAUUAUCCGAUACCGCUUAU	3	0	transposon	LTR	flea
2718	UAGACAGAAAAUAAGGGGACACGCCG	3	0	transposon	LTR	opus
2719	GUUUCGCAAAAGUUGACCGAACUUGA	3	0	ncRNA		rRNA
2720	UCGGAACUUUUGUAACUUUUUCGA	3	0	transposon	DNA	FB
2721	UCAGAUAAAACGUGUUAGUAAAU	3	no match			
2722	GAGUCCGCUAGCUCAUUUCCCGGGAUGU	3	1	transposon	LINE	baggins
2723	UAAACCAUUCGUAGACGACCUGCU	3	no match			
2724	CUAUCCGUUAGGCAGCUGUGACAG	3	no match			
2725	CAAGCCGAGAGGUGACCCAGUUGAGAG	3	no match			
2726	UCACACAGAGGAGGGCAGAUGAGAGCA	3	no match			
2727	UCGCCGAAUCCCGGGGCCGAGGGAGCGA	3	no match			
2728	UGCUGAGCAACGAUUAGACCCCGA	3	0	transposon	LTR	micropia
2729	GAAGUGGAGAAGGGUUCCAUGUG	3	2	ncRNA		rRNA
2730	AGAGGGACGGCCGGGGGCAUUCG	3	no match			
2731	UGGAUAGGGGACCUCAACGGCUA	3	no match			
2732	CUGUAGCCCGAUGACCAGGUAAAUCCGGU	3	no match			
2733	UGCUCAUUCCGAACAUACACCGACACU	3	2	transposon	LINE	Doc
2734	GGAGAAGCCGGCGGGAGCCCCGGGGAG	3	no match			
2735	UCGGCCGAUCGAAAGGGAGUCGGGUU	3	no match			
2736	CUCUCUCAGUCGCUUCCCGGGAA	3	0	ncRNA		rRNA

2737	UUCAGGGGGAUCUUUGAGUGAUGA	3	0	transposon	LINE	baggins
2738	AAAACCUGAAGAUGGUCCUGAGUCGC	3	1	transposon	LINE	Helena
2739	CCCGAAAGCAGACGAGGUGACGUACC	3	1	transposon	Low_complexity	F-element
2740	AACCAAAGCUUUUGGGCUCCGGGGG	3	0	ncRNA		rRNA
2741	CAUGAGAGGUUUGGCCAUAUAAGCA	3	0	transposon	LTR	copia
2742	AUUGCUUGUACCUGAGUCGUUGCG	3	0	transposon	LINE	Circe
2743	AAAGGAGUCUAACACGUGCGCGAGUC	3	no match			
2744	ACCGUCCCAGGUAGACCGUUAUACCAAU	3	0	transposon	LTR	G4
2745	AAGAGUAAUAUGAGCGCCCAGAGU	3	1	transposon	LTR	opus
2746	GGUCGUUCUUCCCGGAGGAGAUCUCGUCC	3	no match			
2747	AAAUCAGCCGACCAAUAGCAUUAA	3	no match			
2748	AUCAAGAACGAAAGUUAGAGGUUCGAA	3	0	ncRNA		rRNA
2749	GCCCUAAACGAAUCGACAUGUCAAC	3	0	transposon	LINE	INE-1
2750	CACGCAGUUACCGACCGUUGUGUCU	3	0	transposon	LINE	Rt1a
2751	AUGGAACAGAAACCCGCGCCAGAG	3	0	transposon	LTR	ninja-Dsim-like
2752	GCUAAACCAUUCGUAGACGACCUGCU	3	no match			
2753	CCUGUAAGCGGAUCCCAUAGGAUG	3	no match			
2754	UGUUCAUCGUUAGACGGCUCGGG	3	2	AT-chX		AT-chX-B
2755	AUCUGGAGGUCCUGUGUUCGAUC	3	no match			
2756	GUGAGGUCUCCGGACGUGAUCAC	3	0	ncRNA		rRNA
2757	UACGAAGUUUCUGAAAGAUGUUGGC	3	0	transposon	DNA	Tc1-2
2758	CUUCGGGAUAAGGAUUGGCUCU	3	0	ncRNA		rRNA
2759	GGGAUCCUCAACUGUGAGGGGAUCCUCAA	3	no match			
2760	ACAUCAAUCGCUGAAAAGGCACCGGGAU	3	1	transposon	LTR	accord
2761	UGCUCUCGAAUGUAUGCCCGAUAUCA	3	2	transposon	LINE	R1-element
2762	ACUCGCCGAAUCCCGGGCCGAGGGAG	3	no match			
2763	CGAGAUAGUACGAAUCGGUUACAA	3	1	transposon	LINE	I
2764	UCUCAUCGUCGUAGAACAAGCCCUA	3	1	Su(Ste)		Su(Ste)
2765	AGAACUGGCGAUGCGGGAUGAACCG	3	no match			
2766	ACCAUAAACGAUGCCAGCUAGC	3	0	ncRNA		rRNA
2767	CAACCUCGAACCUGCUAGCAACAA	3	0	transposon	LINE	G2
2768	UGUUUCAUCGUUACGGCUCGGGCGG	3	no match			
2769	CAGGGUUCGAUUCCGGAGAGGG	3	0	ncRNA		rRNA
2770	CGGCUCAUCUCAUCCUGGGGCUGUAG	3	no match			
2771	GCAUUCGAGACCCAACGACACACA	3	0	transposon	LINE	Rt1b
2772	CAAUUCUGCAACUCGAGCGACGCC	3	0	transposon	LTR	Quasimodo
2773	ACCCAGAACGCGCCCGAUCUCGU	3	no match			
2774	ACCUGAUUGAUAGCGUAGACGACGACCC	3	0	transposon	LINE	Rt1a
2775	ACCGAAAGUAUAUGAGAAGAACACC	3	no match			
2776	GAUGUGCCUAGGACGGAAAUAUGACCA	3	0	coding gene		CG17991-RA|CG17991
2777	CUUCGGGAUAAGGAUUGGCUCUGA	3	0	ncRNA		rRNA
2778	GAGGGCGUGAAACCGUUAAGAGGU	3	no match			
2779	GGGUUCCGGGGGGAGUAUGGUUGCA	3	2	ncRNA		rRNA
2780	UCUGGAGUAAUAAGGCGGGCAGUGAA	3	0	transposon	LTR	diver2
2781	UUCCGGAGAGGGAGCCUGAGAAA	3	0	ncRNA		rRNA
2782	AAGGAGUCUAACACGUGCGCGAGUA	3	no match			
2783	GUGCCCGUAGCUCAGCUGGAUAGAGCAA	3	no match			
2784	CGCAAACCCAAAAUAACAAUGCAG	3	0	repeat		trf
2785	ACUCAUAUCCGAGGCCCUGUAAUU	3	0	ncRNA		rRNA
2786	CUGAAGUUGAGUGGUGAAUCCGUGG	3	0	coding gene		CG41545-RA|CG41545
2787	UAUUCGGAAACGUUUUCUCCAGUA	3	0	transposon	LTR	frogger
2788	GGUGCCAAAAUUAUCGAGCUGCCCUAC	3	0	coding gene		CG8137-RA|Spn2
2789	AACUCGCCGAAUCCCGGGGCCGAGGGA	3	no match			
2790	GACAAAGAGAAGCGGUGUGUGAUCGAU	3	1	transposon	LTR	297
2791	AUUCCGAUAACGAACGAGACUCAAAUA	3	0	ncRNA		rRNA
2792	UCCAACUGUGAGGGGAUCUCAA	3	no match			
2793	GGCGAGAUGGGUGACAGCCCCGUACGGGAC	3	no match			
2794	AGCCGCAGACGCAGACCUACGAGGUGGAC	3	0	coding gene		CG7349-RA|CG7349
2795	UAGCUUCAUCCAAGCGAACUGUGCCC	3	0	transposon	LINE	R1-element
2796	GUUGCAAUUAAACAGAUUCCGCAGU	3	0	Su(Ste)		Su(Ste)
2797	AAUCAUGCCGACGGGCGCUGACCCC	3	no match			
2798	UCUCAUCAUCGUAGAACAAGCCCGAU	3	0	Su(Ste)		Su(Ste)
2799	AAGAGGAGAAGGUGGCUUUGAGAG	3	0	no annotation		
2800	AAGCAGCGUCGGGCGCGGUUAGUAC	3	0	ncRNA		5SrRNA
2801	AGCUCGAUCAUGUGACUCCGGGU	3	0	transposon	LINE	Doc2
2802	AUGCCUGAACGCCUCUAAGGUCGUAUCC	3	0	ncRNA		rRNA
2803	GUGGCAAGAGCGAUUUGAAGGAGUA	3	0	transposon	LTR	micropia
2804	UCCUCUUUAGAAUAACUCGAUAUU	3	0	no annotation		
2805	AGGGACAUUAUUGUGGAAAAUGUGAC	3	0	no annotation		
2806	UGUUUCAUCGUUAGACGGCUCG	3	0	AT-chX		AT-chX-B
2807	AAAACCUAGAAGAUGAUCCUGAGUC	3	no match			
2808	CACGAUGAUCGAUGACGAACUACGAAGG	3	0	transposon	LTR	ninja-Dsim-like
2809	ACCCAAAAGCAGACGAGGUAACGUACC	3	2	transposon	Low_complexity	F-element
2810	ACCGCAACGACCUCCCAGCCCAGC	3	0	transposon	LINE	Doc
2811	CGAGAAACUUAUCGACUAAUCGAC	3	0	transposon	Unknown	RR50960
2812	CUUGGAUUGACUCUAGACCGGAAGC	3	1	transposon	LINE	G
2813	AUCUGUGGGGUUCCAAGCAGCUGAA	3	0	coding gene		CG4931-RA|Sra-1
2814	UUGAUCUUGACACCAAACCGAGCAGU	3	0	transposon	LTR	accord
2815	UGGAGUAUAAGGGUGAGUCCAAGA	3	0	coding gene		CG18743-RA|Hsp70Ab
2816	CGGCAGGGUUCGUGACUGGGGUGAA	3	no match			
2817	AUUCCGAUAACGAACGAGACUCA	3	0	ncRNA		rRNA
2818	GUGUUAUUGUGGGCCGGUACUAUUACU	3	0	ncRNA		rRNA
2819	AUUCCGAUAACGAACGAGACUCAAA	3	0	ncRNA		rRNA

2820	UGCUCUCGAACGCGUGCCCGAUAU	3	0	transposon	LINE	R1-element
2821	ACGCCGACGAGGAGGACAAGAACGA	3	0	coding gene		CG32351-RA|CG32351
2822	GUAACAACGAAGAUGAUUCUGCCC	3	1	transposon	LTR	invader4
2823	ACCACCCUGAACGCGCCCGAUCUCGA	3	no match			
2824	AUCCUCGAUAGUAUAGUGGUUAGUAUCCC	3	0	ncRNA		tRNA
2825	AUGGCUCUCUCAGUCGCUUCCCAA	3	2	ncRNA		rRNA
2826	GGUGAGAUAAUCCCGCAAUAUGGCA	3	0	no annotation		
2827	CUAACUGGAACGUACCUUGAGCAU	3	0	ncRNA		rRNA
2828	GGAGAACUUUGGAGUGAAGACCGCGC	3	0	transposon	LTR	ninja-Dsim-like
2829	UAUUCCGAUAAACGCCCAUUACACUUGC	3	0	transposon	LTR	gypsy3
2830	CUACCGAUUGAAUUAUUUAGUGAGGU	3	0	ncRNA		rRNA
2831	AGCUCGCGUUGAAUUCGUCCCUGC	3	no match			
2832	UGGGCCUGGAGUACUUCAGCCAGA	3	0	Su(Ste)		Su(Ste)
2833	GAUAAUUCCAGACCAACCAAAGA	3	0	transposon	LTR	Idefix
2834	UCCGAUUUCAAGUACCAGGCGAAAC	3	2	repeat		RSP
2835	AUGUCCACCAAAGGCAUGCCCGA	3	0	transposon	LTR	opus
2836	UCUAGGCUGAGCGGAGAACCAGAGUUGA	3	0	transposon	Low_complexity	F-element
2837	ACGAGUAAUAUGAGCGCCCAGAGA	3	1	transposon	LTR	opus
2838	UCUGGGUCGGGGUUUCGUACGUA	3	no match			
2839	CCCUAAGACGCCUGCAGAGAACGCA	3	0	transposon	LINE	G2
2840	CAACCUAUCCAUCAACCCAGACGA	3	0	coding gene		CG18743-RA|Hsp70Ab
2841	UGAAAGUAGUUGGUGGGCACGUGGC	3	0	Su(Ste)		Su(Ste)
2842	ACUGAUGAUGUGUUGUUGCCAUG	3	no match			
2843	UGGACAAAAAAACGAAUGACACAG	3	0	transposon	LINE	accord2
2844	AUUCCGAAGACAAGUAUACCGUGG	3	0	transposon	LTR	micropia
2845	CUUCGGGAUAAGGAUUGGCUCUGAAG	3	0	ncRNA		rRNA
2846	CCUGACAUUACAACAAAUAUUUUUUGA	3	0	Su(Ste)		Su(Ste)
2847	UCCAACGAGAGAGCGGAGAGCCAGCG	3	0	transposon	Unknown	RR50960
2848	UCCGUGGGUGGUGGUGCAUGGCCGUU	3	2	ncRNA		rRNA
2849	CAGGACCAUAUCCGAAGACGCGGC	3	0	transposon	LINE	Rt1b
2850	GAGGCAAUCACCGAGAGCAUCAACGG	3	0	transposon	LINE	F
2851	GACAAAUCUACGAAAAUACGACGUC	3	0	transposon	LTR	diver2
2852	CAGUUCAUUCUUGAGGCCUUCGAG	3	0	transposon	LINE	G2
2853	AAGUUAACGAUCACAAAGGCAUCACGC	3	0	no annotation		
2854	UCACGUCCAGUGUCUGGUUGAAGUACU	3	0	no annotation		
2855	AGCAUACGAGGAAGCAUACGAAGAUAU	3	2	transposon	LTR	invader3
2856	CUGCCCUAUCAACUUUUGAUGGUAG	3	0	ncRNA		rRNA
2857	UGUUUCAUCGUUAGACGACUCGGGCG	3	1	AT-chX		AT-chX-B
2858	AUCAGCUGAACAACACUUUUCUCCGA	3	0	transposon	LTR	gypsy12
2859	CAACAUUGAAUACCGAAAAUAUAA	3	0	transposon	LTR	gypsy12
2860	ACGAGUAAUAUGAGCGCCCAGAGUG	3	0	transposon	LTR	opus
2861	UGGGAACUCUAAUGGGACUGCCGGUG	3	no match			
2862	GAUCGUUCCUUCGAUGACUGAUU	3	0	transposon	LTR	flea
2863	GCUAAUCCCACUGCCAUGCUGCUCUGC	3	0	coding gene		CG6439-RA|CG6439
2864	UGUUUCAUCGUUGGACGGCUCGGGC	3	1	AT-chX		AT-chX-B
2865	AGGUAGCCAAAUGCCUCGUCAUCU	3	0	ncRNA		rRNA
2866	ACCAGAGAUCUAGAGGGCCCAAGG	3	1	no annotation		
2867	CAGGACUUUGCAUACCGGUUCUCGGAA	3	0	transposon	Simple_repeat	I
2868	UCUCAUCGUCGUAGAACAAGCCCGAAA	3	1	Su(Ste)		Su(Ste)
2869	CCCAGAUGCCUCAGGUUGCACCCA	3	0	transposon	LINE	R1
2870	UUGUAACUAGAGCGAUGUAGAUACAC	3	0	no annotation		
2871	AAGCCAGUUACCGAAGCUGCGGA	3	no match			
2872	CGCCCGUCGCUACUACCGAUUGAA	3	0	ncRNA		rRNA
2873	AAGGAAUUGACGGAAGGGCACCAC	3	0	ncRNA		rRNA
2874	AUUACCCAGGUGCGUGUUCUGUAAGAA	3	0	coding gene		CG6694-RA|CG6694
2875	ACUACUGACUGACCGAUAGUGAACC	3	no match			
2876	CGAGAGAAGACUUUCACCCGAUGAG	3	0	coding gene		CG12717-RA|CG12717
2877	AGAAGACUCACCCGACGAGAAUCAAA	3	0	no annotation		
2878	AGAGAUCAAAGAGUGGAUUCGAGG	3	0	transposon	LINE	Rt1a
2879	UACAUCGAAAAGCGAGUUGAUGUGU	3	1	transposon	LTR	Max
2880	AUUGGUUGAACUCUAGAUAACA	3	0	ncRNA		rRNA
2881	UUGGAAGAAACACGAUAUGAACUAAA	3	0	transposon	LINE	Porto1
2882	AAUCGAAUGAGACCUUGGAGGACU	3	0	ncRNA		rRNA
2883	GAAGCAUCGGGUUAGUAAGUAUACGGGC	3	no match			
2884	UGUGCUAGAAGACGAUUUUCGGA	3	0	transposon	LINE	Cr1a
2885	AUCUCAUCGUCGUAGAACAAGCCCGA	3	1	Su(Ste)		Su(Ste)
2886	GAUGUCCUGGGCUGCACGCGCGCUACA	3	0	ncRNA		rRNA
2887	AAUCUCGCGCCGGGCCGUACCCAU	3	no match			
2888	GCAGAACUCCGAUGGCGAUAGCAAC	3	0	no annotation		
2889	CAGACUGACAAAACACAUCCUUUAGGC	3	0	transposon	LINE	BS
2890	GAUUCCGUGGGUGGUGGUGCGUGGCCG	3	no match			
2891	GAUGACUAUAUCAUUCUCGGGAUUAGA	3	0	transposon	LTR	gypsy8
2892	CUAAGUUGUAUUGACAAAUUUAGCAA	3	0	transposon	LTR	opus
2893	UCCCGCAAUAUGGCAAUCGCUCGA	3	0	coding gene		CG40609-RA|CG40609
2894	CAAGACCAGAACCAAUGGACGUGGA	3	0	transposon	LINE	HMS-Beagle2
2895	CGAGAAUCCGAGCAUUUUGUCACA	3	0	transposon	LTR	HMS-Beagle
2896	UCUGAAGAUAAGUGGAGUUGGAAAGC	3	0	transposon	LTR	ninja-Dsim-like
2897	ACCCGAUCGGACCUCACGCAGCUGCAGA	3	0	transposon	LTR	invader4
2898	UGUUUCAUCGUUAGGCGGCUCGGGC	3	1	AT-chX		AT-chX-B
2899	GGGUGUUGACGCGAUGUGAUUA	3	no match			
2900	CAGUGCCAAGGCUAAAGAUGAUUCGU	3	0	coding gene		CG2658-RA|CG2658
2901	UUUGUAAGGACCUGAGAACUUGUACGC	3	0	transposon	LINE	Doc4
2902	GGUGGAGUGAUUUGUCUGCUUA	3	2	ncRNA		rRNA

2903	UACUUGUUCCCCGGAUAGUUUAGUUACG	3	0	ncRNA		rRNA
2904	UCCGAAGACAACGAGGAACAGAGC	3	0	transposon	LTR	1731
2905	UUUGAACAUCGAGAAUACUGCACAG	3	0	transposon	LINE	Doc
2906	CCCGAAAGCAGAUGAUGUAGCGUACC	3	0	transposon	LINE	F
2907	CCGUAGGAUGGCUGUGUUGCAACCC	3	0	transposon	LTR	GATE
2908	UGCUCAUUCCGAACAUACACCGACA	3	0	transposon	LINE	Doc
2909	CACACAACAAAACUAUCACCCAGC	3	0	transposon	Low_complexity	F-element
2910	CGAGAUGUUCAACUACACGGAUUCGC	3	0	transposon	LTR	3S18
2911	UACGUAAACAACCCCAGGCUGACUCCGU	3	1	transposon	LTR	Max
2912	UUCAACAAGGAUAAUAUGCCCGCUCA	3	0	coding gene		CG40601-RA|CG40601
2913	GAGGCCCUGUAAUUGGAAUGAGUA	3	0	ncRNA		rRNA
2914	UAGACAGAGAUUUAACCAGAAGCCACAA	3	0	transposon	LTR	gypsy4
2915	GGGAAUACCGGGUGCUGUAGGCUUU	3	no match			
2916	CGAGUAAUAUGAGCGCCCAGAGUG	3	0	transposon	LTR	opus
2917	AGAGAGCGAAAACUUAAAUACCGAGA	3	0	transposon	LTR	Dm88
2918	UCCGAUCUCCUGAGUCGGACUCCGAGC	3	0	transposon	LINE	R1
2919	CCGCUUCAUCUCGGCACCGGACCCA	3	0	coding gene		CG31538-RA|CG31538
2920	AAGACAAAGAUCGCAAACCUGCUGCA	3	0	transposon	LTR	gypsy6
2921	AACUGGGAAGGUGGGACUUGAGUUU	3	0	transposon	LINE	G2
2922	AAUACAAACAGAAUUCCCGAAUGAGAG	3	0	transposon	LINE	Cr1a
2923	AGACCGAUAUACCUUCGACGACCAGACU	3	0	coding gene		CG5089-RA|CG5089
2924	GAGAUGACAAUGUAGUGAACGCCA	3	0	transposon	LTR	invader4
2925	UAUGAAGAUGGACAACGGGGAUGUU	3	no match			
2926	GACCUCAGAUCAGACGUGGCGAC	3	no match			
2927	CACGACCCCCAGCUACGUGGCCUUCACCG	3	no match			
2928	GACGACGAGCGAGAUCCGGAUGUC	3	0	transposon	LTR	ninja-Dsim-like
2929	AGGAACGGUCGGGGGCAUCAGU	3	no match			
2930	ACCCUGAACUCGCCCGAUCUCGU	3	no match			
2931	CAAGAGUCAAGCCGAUUAUAGUUUAGG	3	0	transposon	LTR	diver2
2932	UUAUAAGGAUGAUGAUACCGAUUA	3	0	no annotation		
2933	ACUGAAUAAAACGAAAUGGAUGACA	3	0	transposon	LTR	invader4
2934	CUUGAGCGAGGCAUGUCUGUUAU	3	0	transposon	LTR	Dm88
2935	GAGCGCACGUGUUAGCACCCGAAAG	3	no match			
2936	CUUUAGAUGAGAUUCCGCUGGCGC	3	0	transposon	LTR	invader1
2937	CACUCAGUUACCGACCGUUGUGUCUG	3	1	transposon	LINE	Rt1a
2938	CCAGUCCGAGUGUGUUUCUGAUCCUACC	3	0	coding gene		CG30372-RB|CG30372
2939	GGCCCAGAAGCGAAGACCGGCGAAUUC	3	0	coding gene		CG15525-RB|CG15525
2940	UGCUCUCGAAUGUAUGCCCGAUAUUUU	3	no match			
2941	UGAGACUACAAACGUAUUGCGGACC	3	1	no annotation		
2942	CCCAAGAAAAAUCCCAGACCUGAUCGA	3	0	transposon	LINE	Doc
2943	CUGGCUGCAACGAGGUGACCGAGGCGG	3	0	coding gene		CG32085-RA|CG32085
2944	ACCCUGAACAAGCCCGA	3	1	coding gene		CG3615-RA|Atg9
2945	ACGAGUAAUAUGAGCGCCCAGAGUU	3	1	transposon	LTR	opus
2946	ACUGAUGAUGUGUUGUUGCCAUGGU	3	no match			
2947	AUGGUGUAAUGGUUAGCACUUUGG	3	0	ncRNA		tRNA
2948	CAGAUACCCGUUACUCAGCUAGUA	3	0	transposon	LINE	INE-1
2949	CGCGGAAAGACGGAAAGACCCCAUGA	3	no match			
2950	UUGAACGUGUCCUGAAAGUAGUUGG	3	0	Su(Ste)		Su(Ste)
2951	UCUGAAGAUAAGUGGAGUUGGAAA	3	0	transposon	LTR	ninja-Dsim-like
2952	CAUCAUUGUUUUCCCGAUGCAUCCGA	3	0	transposon	LINE	R1
2953	UGGAUACAAAACGAGAGAAAGCGGC	3	0	transposon	LTR	GATE
2954	CCCCAUGAACGAGGAAUUCCCAGU	3	no match			
2955	AGAACCAUCACUGGGGCACCGUGGUACA	3	0	transposon	LINE	F
2956	CAGUUCAUCAAAGCGCCUUACUG	3	0	transposon	LTR	flea
2957	AUUGACAUCAUUCAGCGAGCACAA	3	0	transposon	Low_complexity	F-element
2958	GAGCACCGUUUCCCGUCCGAUC	3	no match			
2959	ACCCUGUGGACCCUGUGGACCUUGUUG	3	0	coding gene		CG17945-RA|Mst84Dc
2960	ACCCUACUGAUGGUGUGUUGUUG	3	no match			
2961	CGACUCGUUUUUAAAUUUCGCAA	3	0	transposon	LINE	Cr1a
2962	CAGACUCGGAACGCCUCAUUGGUGA	3	0	coding gene		CG31359-RA|Hsp70Bb
2963	GCUGACGAUAUUGCCCUGUUGUAC	3	0	transposon	LINE	BS
2964	UGGGCCCGAACCCUUCUGGUGUGUUA	3	1	transposon	LINE	baggins
2965	CAAGUUCUCGGCUGAUCAGUACCC	3	0	transposon	LTR	Quasimodo
2966	UGGAUCCUCAACUGUGAGGGGAUCCUCAA	3	no match			
2967	GCUGGGUCGGUCGGGCUGGGGCGC	3	no match			
2968	UUGACUCUAGUCUGGCACGGUGAAGA	3	no match			
2969	AUGCCUUGGCGAUUACAGGCGACGAAGGAC	3	no match			
2970	UCCGAGGCCCUGUAAUUGGAAUGAGUU	3	1	ncRNA		rRNA
2971	CAGCACACAGAGGAGAGCAGAUGAGAGCA	3	no match			
2972	UAGUACAAUAGAUCCGACUGGGCCA	3	0	transposon	LINE	baggins
2973	GCCGAACGUAGUCUCUCGAGAGCGAU	3	1	transposon	LTR	invader1
2974	GUUCGUGGAGUGAUUUGUCUGGUU	3	0	ncRNA		rRNA
2975	AGGGACAUUAUUGUCGAAAAUGUGACA	3	2	no annotation		
2976	CUGGUUAAUUCCGAUAACGAACGAGACU	3	0	ncRNA		rRNA
2977	CUCGCCGAAUCCCGGGGCCGAGGG	3	no match			
2978	CUGGAACGAACUCAUCUGGCCCGCGUA	3	no match			
2979	AAACCUAGAAGAUGAUCCUGAGUCGC	3	no match			
2980	UGUUCUCGAAUGUAUGCCCGAUAU	3	1	transposon	LINE	R1-element
2981	CCCGCCAUAACACUUCUCGUCCCUGCGA	3	0	transposon	LTR	invader4
2982	CCGAACAUCGAAGAUUGGCCUACCCU	3	0	transposon	LTR	297
2983	GUGGAAGUCAAAGGAGUUGAAAUCGCC	3	0	transposon	LTR	Quasimodo
2984	CCCAAUUCAAGAAGUCGGUGUACC	3	1	transposon	LINE	R1-element
2985	UUUGUAAGGACCUGAGAACUUGUACG	3	0	transposon	LINE	Doc4

2986	GUUAGAGGUUCGAAGGCGAUCAGA	3	0	ncRNA		rRNA
2987	UACCGACGACUAAUUAUGGACAAU	3	0	no annotation		
2988	ACCUGAUUGAUAGCGUAGACGACG	3	0	transposon	LINE	Rt1a
2989	GCAUCCGUUACUGUCGAGUCAA	3	0	transposon	LTR	gypsy12
2990	CGGACUCUAAGACGGUGCUGAGAUG	3	0	transposon	LTR	ninja-Dsim-like
2991	CACCAAGCGUUGGAUUGUUCGCC	3	1	ncRNA		rRNA
2992	ACCCCCGCCACGCAGUUUUAUCCGGUAA	3	no match			
2993	AAAACCUGAAGAUCGAUCCCUGUAGUCGCC	3	no match			
2994	CCAGCGAAUACCGAGGGUUUUGGC	3	0	transposon	LTR	GATE
2995	UGUUUCAUUGUUAGACGGCUCGGGC	3	1	AT-chX		AT-chX-B
2996	CCCGAAAGCAGACGAGGUAGCGUAC	3	0	transposon	LINE	F
2997	GACCGAGUAGUGUAGUGGGUGACCAU	3	no match			
2998	ACAACGGCGGGAGUAACUAUGACUCU	3	1	ncRNA		rRNA
2999	AUCCAUGAGCAGGUUGAAGCCGGUGU	3	no match			
3000	GUGAUCGAUACGCCCCCUCAGAGU	3	0	transposon	LTR	Max
3001	GGACUGUGAGGGGAUCCUCAA	3	no match			
3002	GAUAGGCAGAUGUGCCCGAGCGUGG	3	0	transposon	LINE	Cr1a
3003	UUAUGGCUCUCUCAGUCGCUUCCCGG	3	0	ncRNA		rRNA
3004	AGGUGGGGAGUUUGACUGGGGCGGUAC	3	1	ncRNA		rRNA
3005	GAGUCCAAGAGAUUUGCCCCCGAUCC	3	no match			
3006	CCGAACUUGAGACCGCCCCUGGACGAAU	3	1	coding gene		CG40820-RA|CG40820
3007	CAAAACUGCAAGUUUAGAGACGGGUC	3	0	transposon	LTR	ninja-Dsim-like
3008	UGCCAAACGAACCCAUAGCGCAGA	3	0	transposon	LINE	INE-1
3009	CGCUGAGAGCUGAGAGCCCGCAGA	3	0	ncRNA		alphagamma-element
3010	UGCUCUCGGAUGUAUGCCCGAUAUA	3	1	transposon	LINE	R1-element
3011	ACUCGCCGAAUCCCGGGGGCCGAGGGGAGC	3	no match			
3012	AAGAGGCGUGAAACCGUUAAGAGGU	3	no match			
3013	GCAUAAUCAAAACAUCCGAUGACGAAG	3	0	transposon	LTR	McClintock
3014	AGGGAUGAUAAGGAGCCGAAAAACGCCU	3	0	transposon	LTR	springer
3015	GCUGACAAAACUAAGAGUGCCCAAGA	3	0	transposon	LTR	springer
3016	AUCAAUCGCUGAUAAGGCACCGGGAGA	3	2	transposon	LINE	accord2
3017	GGCGAGCGUGAGAUGGCUAACGA	3	0	coding gene		CG8542-RA|Hsc70-5
3018	UGCUCUCGAAUGUAUGCCCGAUUA	3	2	transposon	LINE	R1-element
3019	AACGAUUCUAUUUUUCCGUGCCCAGC	3	0	transposon	DNA	FB
3020	GCGGGUGUAGCUCAGUUGGUUAGAGCGCC	3	no match			
3021	GAAACCUAGAACGACAUAAAUGGCAC	3	0	transposon	LTR	HMS-Beagle
3022	UAAACCUGAAGAGAGCCCUGAAU	3	0	transposon	LINE	Rt1b
3023	AGAUCCUCAACUGUGAGGGGAUCCUCAA	3	no match			
3024	GCCGAACGUAGUCUCUCGAGAGCGAAU	3	2	transposon	LTR	invader1
3025	AACGCAGGUGUCCUAAGGCGAGCU	3	no match			
3026	GCGUAAAACAAUUCAAGAUCUGGCCGG	3	0	transposon	LTR	blood
3027	GACUCGUUGACUGCGCACAGUUGCAUC	3	1	transposon	LTR	micropia
3028	ACGACGAGCUCCCCGAUUUGUGAUUCC	3	0	coding gene		CG34269-RA|CG34269
3029	ACGGAAUGGAAUCGAGCGCUGAG	3	0	transposon	LINE	X
3030	CAAGUUCUCGGCUGAUCAGUACCCG	3	1	transposon	LTR	Quasimodo
3031	GUUCCUCGGGAUCAAAGGCAACGAGU	3	1	Su(Ste)		Su(Ste)
3032	UCUCAUCGUCGUAGAACAAGCCC	3	0	Su(Ste)		Su(Ste)
3033	UGACGUAAUUCGGUUGACCUGUUCGA	3	1	transposon	LINE	Doc
3034	ACGGAAGACCCGCUCCCCGGCGC	3	no match			
3035	GACAGAAACGAGGCCACCGCCGUCCC	3	no match			
3036	UACGAGAACAUCAUCAUCCUCGGC	3	no match			
3037	UCAGAGCAUACAGUAGGCCUAAGU	3	0	transposon	LTR	mdg3
3038	CGGGAUAAGGAUUGGCUCUGAAGA	3	0	ncRNA		rRNA
3039	AGGUGGAAAUGAAGAAAGUACA	3	no match			
3040	ACUUCAUAACCGAGCGACUGUUCAAUC	3	2	transposon	LTR	GATE
3041	AUCAAGGCCUUCUGGUGCAACCGAA	3	0	Su(Ste)		Su(Ste)
3042	CGAGGUCUAAUCCAGCUCAGCAGC	3	0	transposon	Low_complexity	F-element
3043	ACCAGAGAUCUAGAGGGCCCAAGAU	3	1	no annotation		
3044	UAGGCAUGGAACUAGACCUGUUGAG	3	0	transposon	LINE	INE-1
3045	GAUCAUGGAAGUCGAAAUCCGCUAAGGA	3	0	ncRNA		rRNA
3046	AGGAUCUUCAACUGUGAGGGGAUCCUCAA	3	no match			
3047	AUCCCACGGAACACCCGAGCAAUG	3	0	transposon	LINE	G2
3048	AUUUCCUGGGCUGCACGCGCGCUAC	3	1	ncRNA		rRNA
3049	AUCGCAAAGAAACUGUUUCGAAGGACC	3	0	coding gene		CG12717-RA|CG12717
3050	UGUGCUAGAAGACGAUUUUCGGGCCG	3	0	transposon	LINE	Cr1a
3051	GGCGGCGGGGGUGCCGGGCAGCUUCCGGGA	3	no match			
3052	GCGGACAGACAGCAAUUUUGGCC	3	0	transposon	DNA	1360
3053	CGUUGACCAAUUUCCGAACAUACGCCAC	3	0	transposon	LTR	DM88
3054	CAAGAAAUGACGAAAGCAAAGACAA	3	0	transposon	DNA	1360
3055	CGGGUAACGGGGAAUCAGGGUUCUAU	3	1	ncRNA		rRNA
3056	ACAAAAGAUAAUACGGCCAAUGUGG	3	0	transposon	LINE	RR49783
3057	AGAAUAAGAAGACGUCGACGACGAA	3	0	transposon	LTR	HMS-Beagle
3058	AUAUCCGUGAACUACAUCCCAACGA	3	0	no annotation		
3059	UGAACCAGAAAACAAGCCUCCUAGAAA	3	0	transposon	Low_complexity	F-element
3060	CAAGCAAGCAAAUUCAUUUUUUAGA	3	1	transposon	DNA	1360
3061	CAAGAUAUGACCAAAGUAAAGACAC	3	0	transposon	DNA	1360
3062	CAACAUGUCGAGCCCAACAACAACAA	3	no match			
3063	GUUCGUCCUGAGUGACCGCGUACCUUU	3	no match			
3064	CAGUCUAUCAAUCCGAAUUGCAUCA	3	0	transposon	DNA	Ivk
3065	AUCUAGGACUACCAUGGUUGCAACGG	3	0	ncRNA		rRNA
3066	AUCCUCAACUUGAGGGGAUCCUCAA	3	no match			
3067	CUCCUGAUAUGACGCGUGACAUGCCA	3	0	transposon	LINE	G2
3068	CAAGAUAUGACCAAAGCAAAGACUA	3	1	transposon	DNA	1360

3069	AGAUGCAACUGAAGCGCGAUCUCGA	3	0	coding gene		CG6223-RA|betaCop
3070	CAAGAUGAAGAAGAACCGCAUUCA	3	1	coding gene		CG4750-RA|loopin-1
3071	AUAGAAGAAAUUGCCGUUACAA	3	0	transposon	LTR	G4
3072	CAAGAGUUCCCGUGGUAGGCAGUAGU	3	0	ncRNA		7SLRNA-RA
3073	AUCGAUAUUAUCCCAGCACUACGAA	3	0	no annotation		
3074	CCAGACGUGAUACUCCACUACCCA	3	0	transposon	LTR	HMS-Beagle2265
3075	UACGUAACUAUACACUUCAUAAAGACC	3	0	transposon	LINE	Helena
3076	AAGAUAAAGGAGACGCACCGUUGCA	3	1	transposon	LTR	flea
3077	CCCGUAGAAAACCUUUGGAGUGC	3	0	transposon	LTR	HMS-Beagle
3078	UCCCCAAAACAUAACAGACCCUCCUGAA	3	0	transposon	DNA	Tc1-2
3079	AAUAACUGUAAUUAAUAACUGGCCCCC	3	0	transposon	LINE	HMS-Beagle2
3080	GACCAAUGUUGGCGACGAGAAGG	3	0	transposon	LTR	GATE
3081	GACGAACUACCGAGCCCGAGCCUU	3	0	transposon	LTR	Max
3082	ACUCGCCGAAUCCCGGGGGCCGAGGGA	3	no match			
3083	GACUGAGCGGGGACGAGGAGGAUG	3	0	coding gene		pseudogene
3084	AGAGAAUUGAUUUUGCGUUUGAAUGA	3	0	transposon	DNA	invader2
3085	ACUUCGGGAUAAGGAUUGGCUC	3	0	ncRNA		rRNA
3086	AGAAACUUCAUCUCAGUCGGAUUAGCCC	3	0	coding gene		CG41521-RA|CG41521
3087	CUGGCCAUACCCCUCUGCCUCCCGAAG	3	no match			
3088	GAAACCGAAAAAUCACAAUUUGCC	3	0	transposon	LTR	McClintock
3089	ACUUACAAUUGUGGUUAGUACUAUACC	3	0	ncRNA		rRNA
3090	UGCACCGAAAAUAUUUGCGAGGAGAU	3	0	repeat		trf
3091	UCGCCGAAUCCCGGGGCCGAGGGAGU	3	no match			
3092	CCGCCAGCUAAGGUCCCUAAAAUUGGCU	3	no match			
3093	CAAACCGAUAUGCUAUCUUGGCUGAC	3	1	transposon	LINE	Doc
3094	AACGAGAAUAUACACAGAGACUUCCAA	3	0	transposon	LINE	Doc3
3095	AGCAAACUCUUUCGUCAUGCCGAGG	3	0	transposon	LINE	R1-element
3096	CAGACCUAUACAGAUUGGAACAUU	3	0	transposon	LTR	McClintock
3097	UGGUAUCCAGGACUUUGCAUACCGG	3	0	transposon	Simple_repeat	I
3098	ACGAGUAAUAUGAUCGCCCAGAGU	3	1	transposon	LTR	opus
3099	GCAACCCAAGGGACAAGGCAGCAC	3	0	transposon	LINE	G2
3100	UGUCCUUUGUGGGCUCCGACCAGG	3	0	coding gene		CG17896-RA|CG17896
3101	CACUAUCAGACAGCAACAUGACCC	3	0	transposon	LTR	McClintock
3102	CCAAAAAACAACUAACACCUACCGACU	3	0	transposon	LINE	TART-A
3103	CAAGAACGAAAGUUAGAGGUUCGAAG	3	0	ncRNA		rRNA
3104	UGGUAAAGAAAACCAUGUAGCAGAUGC	3	0	transposon	LTR	Quasimodo
3105	ACACCGCCCGUCGCUACUACCGAU	3	0	ncRNA		rRNA
3106	AAAUACCAUAGCAAGCUUCACACGC	3	0	transposon	Low_complexity	F-element
3107	GACAUCAUUUCGAGACCCAUUUUA	3	0	transposon	LTR	gypsy12
3108	GCAACGGGUAACGGGGAAUCAAGGUU	3	1	ncRNA		rRNA
3109	GGAUCCUCAACUGUGAGGGGUCCUCAA	3	no match			
3110	GAUGGGGGCCGCUUGGGAACACCGC	3	no match			
3111	CCCGGAAUGUUCUCAGGAAGAUGACUCA	3	0	ncRNA		snoRNA
3112	UGGACAGAAACCACCCGACGGGUAGC	3	0	transposon	LINE	INE-1
3113	AAAAGGAAUUGACGGAAGGGCACCACC	3	1	ncRNA		rRNA
3114	GAUCUCCUGAGUCGGACUCCGAG	3	0	transposon	LINE	R1
3115	ACCGAAUUAAUCGCGAGGCUGCAU	3	1	transposon	LTR	mdg3
3116	UAGAACAAGCCCGAGGAACUGUCGAA	3	0	Su(Ste)		Su(Ste)
3117	CUUCAUCCAAGCGAACUGUGCCCG	3	0	transposon	LINE	R1-element
3118	GGGUUCCGGGGGGGAGUAUGGUUGC	3	no match			
3119	GACCAAAUCAACACAUCCGACCAGC	3	0	no annotation		
3120	UCCCACGGAACACCCGAGCAAUGCCACC	3	0	transposon	LINE	G2
3121	AUACCAAGAUGAUUCAGUGCGCCG	3	0	coding gene		CG7642-RA|ry
3122	UAGUGAGGUCUCCGGACGUGAUCACUG	3	0	ncRNA		rRNA
3123	AGGCGUAGUCGAUGGAUGACGGGUU	3	no match			
3124	UUAACCAGAACUGAGUCCAGAACAGA	3	1	transposon	LTR	gypsy3
3125	UGAUCCUGAAGCCGGUGAUCGAUAGU	3	0	repeat		trf
3126	CCAGCUCGGGGAGGUAGUGACGAAAA	3	0	ncRNA		rRNA
3127	CGUAAAACAAUUCAAGAUCUGGCCGG	3	0	transposon	LTR	blood
3128	UCGGCCCGAACCCUUCUGGUGUGUUGG	3	0	transposon	LINE	baggins
3129	GAAGAUAUCAGGGGCCUUCGCAG	3	0	transposon	LTR	HMS-Beagle
3130	UUUGCAAAAACGAGAGCGAUAGGC	3	2	transposon	LTR	micropia
3131	UCGCGGUGAAUGCGUUCCCGGGUCUUGUA	3	no match			
3132	AACCGAUAUGGUUGGACCACAAUCUU	3	0	ncRNA		7SLRNA-RA
3133	GCCGAACGUAGUCUCUCGAGAGCGAGA	3	0	transposon	LTR	invader1
3134	UGUCCCAAGGCCAGCUCAGUGCGGAC	3	0	ncRNA		rRNA
3135	UGGCUCUCUCAGUCGCUUCCCGGGAAA	3	0	ncRNA		rRNA
3136	AGCAACUGAACCCAAGAAAAUCGCC	3	0	transposon	LINE	Doc
3137	GGAUCCUCAACUGUGAGGGGUCUCAA	3	no match			
3138	AUGGAACAGAAACCCGCGCCAGAGG	3	0	transposon	LTR	ninja-Dsim-like
3139	CUUCGGGAUAAGGAUUGGCUCUGAAGA	3	0	ncRNA		rRNA
3140	UCUCAUCGUCGUAGAACAAGCCCGAUAU	3	no match			
3141	CCGUACUUUACUUUUGCAACAAAAC	3	0	transposon	DNA	1360
3142	ACCAGGUGGCGCUGAACCCGCAGA	3	no match			
3143	GGACUCAUAUCCGAGGCCCUGUAAUUGGA	3	0	ncRNA		rRNA
3144	GCGUAAAACAAUUCAAGAUCUGGCCU	3	1	transposon	LTR	blood
3145	ACGAGUAAUUGAGCGCCCAGAGU	3	no match			
3146	AAUUCCGUAAACGAAGCCCAAAGUUU	3	1	transposon	LTR	Rt1c
3147	AACGUAGAGAGCAACAUGCCUGGAGUC	3	0	coding gene		CG5338-RB|RpS19b
3148	UGCUCUCGAAUGCUUGCCCGAUA	3	0	transposon	LINE	R1-element
3149	CCGGCGACGGUGUGGGUCCCGAGCUGGUCU	3	0	coding gene		CG6439-RA|CG6439
3150	GGCACGAAACGGUAUUGGCACCAU	3	0	transposon	LTR	Idefix
3151	AGCCUGGGCAACAUAGCGAGACCCCGUC	3	no match			

3152	CAAGCAUUCGACAAAGUCUGGCUCGAC	3	0	transposon	Low_complexity	F-element
3153	UGGAACGAAAUUGGCCUGAUUAGC	3	0	transposon	LTR	GATE
3154	CAAAGGCAUGCCCGAUGUCCGCGAG	3	0	transposon	LTR	opus
3155	GUGGUGGUGCAUGGCCGUUCUUAGUUC	2	0	ncRNA		rRNA
3156	GCGGAAAGAGAGCGAUUUUGGCC	2	0	transposon	DNA	1360
3157	UGCUCAUUGAACUCCCAAAGUGUGC	2	0	transposon	LINE	Doc
3158	UCCUCAACCAGCACGACAAGGCCCAA	2	1	transposon	LINE	G2
3159	CAAGUUCUCGGCUGAUCAGUACCCA	2	0	transposon	LTR	Quasimodo
3160	GCCAGAGAACCCUCUAAUUGGGCCCGU	2	1	transposon	LINE	baggins
3161	CCGACAAGGAGACGGGUGACGAGAAGUC	2	0	coding gene		CG8583-RA|sec63
3162	UUCGUAACAAAAUUCCCCAUCGACC	2	0	repeat		SAR_DM
3163	AAAACAGACACGAACGAAAUUUUGA	2	0	transposon	LTR	Idefix
3164	ACCCGAAAUAACCGAAACGGAAGACC	2	0	transposon	LTR	accord
3165	AACUGAAUAAACGAAAUGGAUGACGC	2	1	transposon	LTR	invader4
3166	ACUCGCCGAAUCCCCGGGGCCGAGGGAGC	2	no match			
3167	AAACCAAAGCUUUGGGCUCCGGGGGA	2	no match			
3168	AGAACCAUCACUGGGGCACCGUGGUAC	2	0	transposon	Low_complexity	F-element
3169	GCAUUCGAGACCCAACGGCACGUGAUC	2	2	transposon	LINE	INE-1
3170	GGACCCUGUGGACCCUGUGGACCUUGUU	2	0	coding gene		CG17945-RA|Mst84Dc
3171	CCAGACGUAAUACUCCACUACCCGGAA	2	1	transposon	LTR	HMS-Beagle
3172	CAUCCUAGAAUCGACCGACCAGCUGUGC	2	0	transposon	LTR	invader3
3173	ACCCCUGAACGCGCCCGAUCUCGU	2	no match			
3174	AAAGCUUAUCCUUUGCUUGAUGAUUCGA	2	1	ncRNA		rRNA
3175	GCAUUCGAGACUCAAUGGCACGUCAU	2	1	transposon	LINE	Rt1b
3176	AAUGGGCGUGGGUUUUGUGUUGAGA	2	0	transposon	LINE	Circe
3177	UCACCAAGCGUUGGAUUGUUCACCCA	2	0	ncRNA		rRNA
3178	UACUUCGUUACUACCCAUUUGGAA	2	0	no annotation		
3179	GGCUUUGAGAGGCGCCGUAGUUAA	2	1	no annotation		
3180	GUCUGGUUAAUUCCGAUAACGAACGAGG	2	1	ncRNA		rRNA
3181	AGCUUCAUCCAAGCGAACUGUGUCCG	2	1	transposon	LINE	R1
3182	UAUGGCUCUCUCAUUCGCUUCCCGGGA	2	1	ncRNA		rRNA
3183	AGAGAUGGAUUGGUGCCUUCGGGA	2	no match			
3184	GCGAAUGAUAAGUUUCAUGCCACCGG	2	0	no annotation		
3185	UUCUUGACAACCCGAUCGGACCUCA	2	0	transposon	LTR	invader4
3186	UAUGGCUCUCUCUGUCGCUUCCCGGGA	2	1	ncRNA		rRNA
3187	UCUACACCAAAGUAAGCCGUGCC	2	0	coding gene		CG31359-RA|Hsp70Bb
3188	GUGCUAGAAGACGAUUUUCGGGC	2	0	transposon	LINE	Cr1a
3189	ACCAGAGAUCUAGAGGGCCCAAGAA	2	0	no annotation		
3190	AUCAACUGUGAGGGGAUCCUCAA	2	no match			
3191	UACACAGAGGAGAGCAGAUGAGAGCA	2	no match			
3192	GGAAUCGACGGCGCCCAGCGAUACAAC	2	1	coding gene		CG32656-RA|CG32656
3193	AAGAGUUCCCGUGGUAGGCAGUAGU	2	0	ncRNA		7SLRNA-RA
3194	UCCGGGAAGAACGACCGGCUCUCGCC	2	no match			
3195	UGAGAUAUGCUAGCAGACUGGAGAACC	2	0	transposon	LINE	G2
3196	CAGACAAGAAGGUGGCAUCGGUG	2	0	transposon	DNA	H
3197	AGGUCAUCGGUUCGAUCCCGUUUA	2	no match			
3198	GAGGGCGUGAAACCGUUAAGAGG	2	no match			
3199	CGUACGCUGACGAUAUUGCCCUGUU	2	0	transposon	LINE	BS
3200	AGAGGCGUGAAACCGUUAAGAGGU	2	no match			
3201	CGGAGAGGGAUCCUGAGAAACGGC	2	1	ncRNA		rRNA
3202	UGAGCGAACAAACCCAACAUAGUGCA	2	0	transposon	LTR	McClintock
3203	CGGACUCUAAGACGGUGCUGAGAU	2	0	transposon	LTR	ninja-Dsim-like
3204	AGGAAUAAUGGAAUAGGACCGCGGUUC	2	no match			
3205	UAUUUACAAAACCGUUGCCUGCUGC	2	0	transposon	LINE	Cr1a
3206	UUCGAUUUCAAGUACCAGGCGAACA	2	0	repeat		RSP
3207	CGGAUCAGUUCGCCCAAGUCGCCGCC	2	0	coding gene		CG5151-RA|CG5151
3208	GCCGGGCCCGUAUAACGUUAAUGA	2	1	ncRNA		rRNA
3209	ACGAGUUCGAUCGCGAGCGGCAGGCCC	2	0	coding gene		CG10958-RA|CG10958
3210	UGCUCUCGAAUGCUUGCCCGAUAUU	2	1	transposon	LINE	R1-element
3211	AAGAGGACGUGAAACCGUUAAGAGGU	2	no match			
3212	CUGGGAACACCGCGUGUUGUUGGCCU	2	1	ncRNA		5SrRNA
3213	GACGGAGCUGCUGAUUCAUCCGGACA	2	1	coding gene		CG6750-RA|CG6750
3214	CGGAGGUGUCCCAAGGCCAGCUCA	2	0	ncRNA		rRNA
3215	ACAGCUAUGUACCCGACGGCAAGAC	2	0	transposon	LINE	G2
3216	AUACAAACAGAAUUCCCGAAUGAGAG	2	0	transposon	LINE	Cr1a
3217	ACCAACUACAACACAUCGAGCGACCG	2	0	transposon	LINE	Doc3
3218	AAGAGACAAAGCAGGCUGCACGAAA	2	0	transposon	LTR	micropia
3219	UGUUUCGUCGUUAGACGGCUCGG	2	1	AT-chX		AT-chX-B
3220	ACGGAGGUGUCCCAAGGCCAGCUCAG	2	0	ncRNA		rRNA
3221	GAAAGAAAAGGGAGGUCGAGGUUU	2	0	coding gene		CG42481-RA|CG42481
3222	CAGCCUGAGAGGUGAUGCGUAGCCGA	2	no match			
3223	GACGCGAACACGAAGCUCGAGGUC	2	0	transposon	LTR	GATE
3224	GUAACCUGGAACUUAACCCAAACAA	2	0	transposon	LTR	297
3225	CGUAGAACAAGCCCGAGGAACUGUCGC	2	1	Su(Ste)		Su(Ste)
3226	AGCAGAGACCGUACCCGAAACCG	2	no match			
3227	AUAUCCGUGAACUACAUCCCAACGACCG	2	0	no annotation		
3228	GCUUGGGAACACCGCGUGUUGUUGGCCU	2	0	ncRNA		5SrRNA
3229	UAGGCGAAACGAACCGGGUACCUC	2	0	transposon	Simple_repeat	I
3230	AGGUGGGGAGUUUGGCUGGGGCGGC	2	no match			
3231	ACUGAGAAGAUAUUGUACCCGUGAA	2	0	coding gene		pseudogene
3232	AGUGGGAGACGGUGGUGGGUGAUGA	2	no match			
3233	CAAGCGGUGGAGCAUGUGGUUUAAUU	2	no match			
3234	AAAGAGAGCAAUUUUGGCCGUCAC	2	0	transposon	LINE	Cr1a

3235	AUGGCUCUCUCAGUCGCUUUCCCGGGA	2	no match			
3236	UAGUGAGGUCUCCGGACGUGAUCACU	2	0	ncRNA		rRNA
3237	GCGAAGAUGUAACGGGGCUCAAGACGU	2	no match			
3238	GAAGAUGAAAGCAGAACCGAGUCUGA	2	0	transposon	LINE	Rt1a
3239	AAAAACCGAAUGCGAUCCGGAUU	2	0	coding gene		CG4714-RA|CG4714
3240	ACCACUGAAACAGCAGAAGCCGCA	2	0	transposon	LINE	X
3241	GUUCGAUCCCGGAUGGAACCACCA	2	2	ncRNA		tRNA
3242	CGUAAACAAUUCAAGAUCUGGCCGGC	2	no match			
3243	UGCCCGAGGAUUCCGACACCUACUUGCA	2	0	coding gene		CG7269-RA|Hel25E
3244	GAAUCCCGGGGCCGAGGGAGCGAGACCCGUC	2	no match			
3245	UGAGGUCUCCGGACGUGAUCACUGUGA	2	0	ncRNA		rRNA
3246	ACUGGGAGAAGGUGGGCUUUGUG	2	no match			
3247	GUGAUGUCAACGUUUGAUCCCCUG	2	0	transposon	LTR	ninja-Dsim-like
3248	UGCCGAACGUAGUCUCUCGAGAGCG	2	0	transposon	LTR	invader1
3249	AAGGACACUAGCUUCUUAAAUGGACAA	2	0	ncRNA		rRNA
3250	AAGCGCUGGUAAACGGCGGGAGU	2	2	ncRNA		rRNA
3251	GAGUGCCUAGUGGGCCACUUUUGGU	2	no match			
3252	AACACAAAGACGGGGUGACCACGGGA	2	0	coding gene		CG5546-RA|MED19
3253	AGAACGAUGACGAUCCCACGGCUGGAU	2	1	transposon	LTR	ninja-Dsim-like
3254	ACGAGGUAACCACGAGCAGCGAGGAGC	2	0	coding gene		CG11666-RA|CG11666
3255	CGCGUCAGAAAACGUUACGCCGACGAG	2	1	repeat		trf
3256	GACGAGGAGGUCGAUGAGAUGAUCCGGG	2	0	coding gene		CG8472-RA|Cam
3257	CCACCCUGAACGCGCCCGAUCUCG	2	no match			
3258	ACAUUCAUAACCGAGCGACUGUUUAA	2	1	transposon	LTR	GATE
3259	AGAUCAACUGUGAGGGGAUCCUCAA	2	no match			
3260	AGGACAAUUAUCCGAUACCGCUUAUC	2	1	transposon	LTR	flea
3261	CAACCGACUACAAAGAUGCCAUACG	2	0	transposon	Unknown	RR50960
3262	ACCCCGAACGCGCCCGAUCUAGUC	2	no match			
3263	AGCUCGAUCAUGUGACUCCGGGUAG	2	0	transposon	LINE	Doc2
3264	UGUACUACCCCUUCAUGACUCAAAGAU	2	0	transposon	LINE	Fw2
3265	GUGGGUGGUGGUGCAUGGCCGUUC	2	1	ncRNA		rRNA
3266	UCCGGGAACAUGCCACCUCCGUUGUUGU	2	0	coding gene		CG1763-RA|nod
3267	GCCAAUGAAAGUCACCCCUAAACCCGA	2	0	transposon	LTR	rover
3268	UUAGCCGAUCUAGAUGCCCAGCAUU	2	2	transposon	LINE	Doc2
3269	CCGAACGUACCACGGUGCCCCAGUGA	2	0	transposon	Low_complexity	F-element
3270	UAAUAGGGAACGUGAGCUGGGUUU	2	no match			
3271	AUGGCUCUCUCAGUCGCUUCCCUGGA	2	1	ncRNA		rRNA
3272	CGGAUAUACAUUUUGCAAGUGACG	2	0	transposon	LINE	TART
3273	UGCGGUGAAUACGUUCCCGGGGCUUGUA	2	no match			
3274	GACCAUGAGAUUACACUGGAGGAUGUAC	2	0	transposon	LINE	Cr1a
3275	AAAGCGCGAAGAAGAUGACUGGCC	2	no match			
3276	GGUUCCGGGGGGAGUAUGGUUG	2	2	ncRNA		rRNA
3277	UACUUGUUCCCCCGGAUAGUUUAGUUAC	2	no match			
3278	AAGUCCCGCAACGAGCGCAACCCUCGC	2	no match			
3279	AGAGAGCGAUUUUGGCCGUCACUAA	2	0	transposon	DNA	1360
3280	GAGAAGAACGAUCCAGACAAGCCCA	2	0	coding gene		CG12366-RA|O-fut1
3281	UUAGAUCGUUAACAGUUACUUGGAU	2	0	ncRNA		rRNA
3282	UCAACUGUGAGUGGAUCCUCAA	2	no match			
3283	UCUAAUGAGACUGCCGGUGACAA	2	no match			
3284	UUCGUCGAUAAACAGAUUUCUGAGUC	2	0	no annotation		
3285	CCAGUCAGUAGAGCACUACCGCCGG	2	0	repeat		trf
3286	AGAUCAGACGUGGCGGCCCCGGGU	2	no match			
3287	AAAACCCGAGAUGAUCCUGAGUC	2	no match			
3288	UCGAGAUCGCAGGAAUAAGGUUGG	2	no match			
3289	UACGUCAACAUAUCAGGGCAGCCCG	2	0	transposon	LINE	Doc4
3290	AGGGCGAGAGCCGGUCGUUCUUCCCGGAGG	2	no match			
3291	CCCAAGAAAAUCGCCUGGCUACGACC	2	0	transposon	LINE	Doc
3292	GUGUACCCUGCACCCCAGCCUACAUCCA	2	0	coding gene		CG17567-RA|CG17567
3293	AAACUCUGGACCAGAUGAUAGAGGGC	2	no match			
3294	UCGCAUCGUCGUAGAACAAGCCCGAGG	2	0	Su(Ste)		Su(Ste)
3295	ACCCAGUUCAUUCUUGAGGCCUUCGAC	2	1	transposon	LINE	G2
3296	GCAACUCAAACUCGAAACGGCAUAA	2	2	transposon	LTR	ninja-Dsim-like
3297	UAUGGCUCUCUUAGUCGCUUCCCGGGA	2	1	ncRNA		rRNA
3298	GCUGAAUCAACAUAAACAUCGGGC	2	0	no annotation		
3299	GAGACCGAAUGAUUUAACCCGGA	2	0	transposon	LTR	Rt1c
3300	CAACAAACAACACAUACCCGCUGG	2	0	transposon	LTR	GATE
3301	AGAGCAUAAAGACGACGAGAGUGA	2	1	transposon	LINE	Doc3
3302	GGAGUUAAAAAAUCACGUCAUCCCUAG	2	0	Su(Ste)		Su(Ste)
3303	GCAGCGACGAGAGCGAUGGGCCUG	2	0	transposon	LTR	GATE
3304	AACCAAAGCUUUUGGGCUCCGGGGGAA	2	0	ncRNA		rRNA
3305	UAUCUUCAACAAGGAUAAUAUGCCCGC	2	0	coding gene		CG40601-RA|CG40601
3306	AGUCGGUCCUGAGAGAUGGGCGAGCG	2	no match			
3307	CGACUCGAAAUAGUAACUACGUAAU	2	0	transposon	LTR	17.6
3308	UUUACUGAGACAUAGUGACAUCGGCCA	2	1	no annotation		
3309	AUCACUGAGAUGCCCCGGCGCGGA	2	0	coding gene		CG14355-RA|CG14355
3310	GCAGUCGUGGCCGAGCGGUUAAGG	2	0	ncRNA		tRNA
3311	AUAUCCGUACUACAUCCCAACGACC	2	no match			
3312	UCUCCGGACGUGAUCACUGUGAC	2	0	ncRNA		rRNA
3313	UCCUUAACAAGAACGACGUUGUCC	2	0	transposon	LTR	diver2
3314	UACAAGCAAGACCACCACACCCGACU	2	1	transposon	LTR	gypsy12
3315	CAAGAUAUGACCAGAGCAAAGCAAU	2	1	Su(Ste)		Su(Ste)
3316	UCCUGAUGGAAGACCGAAACAGUUCUGA	2	1	ncRNA		rRNA
3317	AGGGGUGAUAACGAGUCUGCUUG	2	no match			

3318	ACAGAGAGAAGACUCACCCGACGUC	2	2	no annotation		
3319	CUAUCCUGAAUUCACGCCCGCUGU	2	0	transposon	LTR	3S18
3320	ACUCGCCGAAUCCCGGGUCCGAGGGAG	2	no match			
3321	UGAGUAACGGAUAUACGAUAUAGAU	2	0	no annotation		
3322	CGCAAACCCAAAAUAACAAUGCAGG	2	0	repeat		trf
3323	CCCCUGACACAAGCACCCAGACCAU	2	1	transposon	LTR	McClintock
3324	CCGAACUUGAGACCCAACAACGCAAC	2	no match			
3325	AGAGACAAUUGACGGUACCGCAC	2	0	transposon	LTR	Dm88
3326	GAGUUCGAAUCUCUUAGCAACCACC	2	no match			
3327	CACCCUGAACGCGCCCGAUCCCCUC	2	no match			
3328	CCCCUGGACAACCCACUAAGGCCUUGA	2	no match			
3329	ACCGACGAUGAAAUGGCUGAGCUG	2	0	no annotation		
3330	UAACCAUGAGACCGAUAGUAAACA	2	0	ncRNA		rRNA
3331	AGCGAAAUGACUCUGUCCCGAUUC	2	0	transposon	LINE	R1-element
3332	ACUCGCCGAAUCCCGGGGGCCGAGGG	2	no match			
3333	GUCCCGGGACAUGGCGUGGUGGGUAGUU	2	no match			
3334	ACCCUGAACGCGCCCGAUCUCGUU	2	no match			
3335	GUGGCUUUGAGAGGCGCCGUAGUUG	2	1	no annotation		
3336	CGAGAUAGUACGAAUCGAUUACGAAA	2	0	transposon	LINE	I
3337	UCUGGCGAAGAUGACUACGAGGA	2	0	coding gene		CG9351-RB|flfl
3338	GUAGGGAGUUUGACUGGGGCGGUACA	2	0	ncRNA		rRNA
3339	GCAACCGAUAACGCCCACCGAGGUCC	2	1	transposon	LTR	ninja-Dsim-like
3340	CAAUCCAGAAUGUUCUCCUUGAGG	2	0	transposon	LINE	G2
3341	AGACCAAGUAUCCAAGUUUCUCCGA	2	0	coding gene		CG4691-RA|CG4691
3342	GUAUCACGAGAAGGAACAUACCAAGGA	2	0	coding gene		CG34074-RA|mt:CoIII
3343	AACACCGAGAAGUGACGACGAGUGUCU	2	no match			
3344	GUCCAGAGGAGGCGAGAAUUUUACA	2	0	AT-chX		AT-chX-B
3345	UCCCAGCUCGGGGAGGUAGUGACGAAA	2	0	ncRNA		rRNA
3346	CAUUAAUGAAGUGUUGGCCCAAU	2	0	transposon	LTR	HMS-Beagle
3347	UUGUUCCCCGGAUAGUUUAGUUA	2	0	ncRNA		rRNA
3348	AUCGAGGAAUAAAGUUUACAAGU	2	0	transposon	DNA	transib2
3349	AAAAAGGUCGUAUCCGUGCUGGACUGCA	2	no match			
3350	AGAACUGGCGCUGCGGGAUGAACCGAAC	2	2	ncRNA		rRNA
3351	UUGACUAUUAUUUGGAGAACCACAGCA	2	0	transposon	LTR	opus
3352	AUUGAGCGAACCCGGGGAACUGAAAC	2	no match			
3353	GAUGGAAGACCGAAACAGUUCCGA	2	1	ncRNA		rRNA
3354	AAGACCGUAAACGAAAAACUAAGAAUC	2	0	transposon	LTR	Idefix
3355	CACUAUAACAACGGACCAGGGAAU	2	1	transposon	LINE	Doc4
3356	CCGAACGUACCACGGUGCCCCAGUGAUG	2	0	transposon	Low_complexity	F-element
3357	UGUUAUCUAAGGCGACAUGGGUGA	2	1	transposon	LINE	Cr1a
3358	AACCGAUUAACGUUGAAAAGUUACCGGA	2	no match			
3359	AGUGCGCAGGUUGUGGGAAAAAGUACC	2	no match			
3360	ACAAACAGUACACCUACUAGCCCA	2	0	transposon	LINE	baggins
3361	CCGCUGAGCCACUUUCAUGCUUGGGAUU	2	1	ncRNA		rRNA
3362	UGCUCAUUCCGAACAUACACCGA	2	0	transposon	LINE	Doc
3363	CCGCCCGUCGCUACUACCGAUUGAAUUA	2	0	ncRNA		rRNA
3364	UGAGACUACAAACGGAUUGCGGACCC	2	1	no annotation		
3365	UAAACGAGUGUUAUUGUGGGCCGGUAC	2	0	ncRNA		rRNA
3366	AAGUCUGGUGCCAGCAGCCGCGGUAAU	2	0	ncRNA		rRNA
3367	UUUGCAAAAACGAGAGCGAUAGAUC	2	1	transposon	LTR	micropia
3368	UGGGUGCUAACGGGGCCGAUUUC	2	0	transposon	LTR	Rt1c
3369	GGUAGAGAGUGAUGCCGAGUAUAA	2	0	repeat		trf
3370	GGAUCAACUGUGAGGGGAUCUCAA	2	no match			
3371	GGAUCCUCAACUGUGGGGGAAUCCUCAA	2	no match			
3372	ACCCAUUACAUCCUAUACGGUAAUUU	2	0	transposon	LTR	GATE
3373	ACGAGUAUAUGAGCGCCCAGAGU	2	no match			
3374	UCUCAUCGUCGUAGACAAGCCCGA	2	no match			
3375	ACUUCGGGAUAAGUAUUGGCUCU	2	1	ncRNA		rRNA
3376	CCCCGCGGAAAGACGGAAAGACCCCAUG	2	no match			
3377	ACGACGCGAGAGGUGAAAUUCU	2	0	ncRNA		rRNA
3378	CAGCAAACAAACAUCCUUGACGGAAUC	2	0	transposon	LTR	invader6
3379	ACCGAUUAAUCGCGAGGCUGCAAG	2	no match			
3380	UCCAGGGGCGGUCUCAAGUUCGGC	2	0	Su(Ste)		Su(Ste)
3381	CGUGGACGGCGUAAGUCCUUGUCA	2	1	coding gene		CG9284-RA|CG9284
3382	ACCUUCUCGAGCUAGCGGUUGA	2	0	transposon	LTR	3S18
3383	CCUCAAAUCAUUCACUGAGGGUAGA	2	0	transposon	LINE	G
3384	CAGCCAAGAGUUCCCGUGGUAGGC	2	0	ncRNA		7SLRNA-RA
3385	UGUUACAAAAAAAAAGCUUCGUCUCGA	2	0	transposon	LTR	gypsy3
3386	ACCGAUUCGGCGACAGCAGCGAGGGU	2	0	coding gene		CG14472-RA|poe
3387	UAUCCAUCAGUCCCAGCCCAAGUA	2	0	transposon	LTR	McClintock
3388	AAACCAAGCUUUUGGGCUCCGGGGGA	2	2	ncRNA		rRNA
3389	CACGAAUUUACUACUGCUUUCAUCAGG	2	0	transposon	LTR	GATE
3390	AACUCGCCGAAUCCCGGGGCCGAGGGAG	2	no match			
3391	UGAGACUACAAACGGGAUUGCGGACC	2	no match			
3392	UCCGGGAACAUGCCACCUCCGUUG	2	0	coding gene		CG1763-RA|nod
3393	CGAGAUGUUCAACUACACGUAUUCGCC	2	1	transposon	LTR	3S18
3394	AUGUCUAUGAGAUGACCGUAAUGAG	2	0	transposon	LTR	ninja-Dsim-like
3395	GACCAUAUCCGAAGACGCGGCUCUAA	2	1	transposon	LINE	Rt1b
3396	UCGCCGAAUCCCGGGGCCGAGGGAGCGAGA	2	no match			
3397	GGCUCCGGGGGAAGUAUGGUUGCA	2	0	ncRNA		rRNA
3398	GCUGCAGGACGGUAUAAUUCGACC	2	0	transposon	LTR	opus
3399	AUCACCCUGAACGCGCCCGAUCUCGU	2	no match			
3400	CUUGGGAACACCGCGUGUUGUUGGCCU	2	0	ncRNA		5SrRNA

3401	UGAGUAAAUUGGACGAUGAGGCGU	2	1	transposon	LTR	Rt1c
3402	UACCACUCAUCAACCAAAGGACGUA	2	no match			
3403	CACGCAAAAAGGCUCUGAGAACACGCAA	2	0	transposon	LINE	G6
3404	AGGGAUGAGAUGCCAGCUGAGCAU	2	0	transposon	LINE	baggins
3405	AUGACCAAGAACUGUGGGCACGAUA	2	0	transposon	LTR	RR50204
3406	UCUCAUCGUCGUAGAACAAGCCGGA	2	1	Su(Ste)		Su(Ste)
3407	CUGCGAUCCACUGAAGCUUAUCCUU	2	0	ncRNA		rRNA
3408	CAACGGGUAACGGGGAAUCAGGG	2	0	ncRNA		rRNA
3409	UUCCUCGGGAUCAAGGGCAACGAGU	2	0	coding gene		CG40635-RA|CG40635
3410	UACGAACCCGAACGAAACCAUUUUGA	2	0	transposon	LINE	accord2
3411	GACAAUGUGAACAAGACGAGCGCCGG	2	0	transposon	LINE	Doc
3412	CACACAGAGGAAGCAGAUGAGAGCA	2	no match			
3413	AUAACGUGUCCUGAAAGUAGUUGGUGG	2	2	Su(Ste)		Su(Ste)
3414	AGAGAUGAUCAGCAGUUCCCCUG	2	no match			
3415	ACACCGACGAGGGCGAGUUCGACUGGAC	2	0	coding gene		CG6963-RA|gish
3416	ACCCUGAACGCGCCCGAU	2	2	no annotation		
3417	CGGGACAUGGUGUCGGACCGACUUG	2	0	transposon	Simple_repeat	I
3418	GACCCUGAUCCGUGAACGUGGAAUGGA	2	0	coding gene		CG4750-RA|loopin-1
3419	UCCAGACGUAAUACUCCACUACCCGG	2	0	transposon	LTR	HMS-Beagle
3420	CAAUAUGUAAGAACACCAUGUACGGGU	2	0	transposon	LTR	Stalker2
3421	AGGAAAUGAUAGAACAAGGGAUUGU	2	1	transposon	LTR	Idefix
3422	UCUCCCCGUGAGGCACCUGUAUU	2	no match			
3423	UUCAACGGGUAACGGGGAAUAAGGGUUC	2	2	ncRNA		rRNA
3424	GAAACCAACCGGGAUUGCCUUAGUAAC	2	no match			
3425	GCUGAGAAAAAACGAAGUAUAACGA	2	no match			
3426	UGGUGCCGGCGGUGAUGGGGGUUACU	2	0	coding gene		CG3506-RA|vas
3427	GCAGGACGGUGAUCAUGGAAGU	2	0	ncRNA		rRNA
3428	GUGAGGUCUCCGGACGUGAUCACUGUGA	2	0	ncRNA		rRNA
3429	CAAAGCAUAAAGGACCCUAUGUCAAA	2	1	transposon	Unknown	RR50960
3430	GCACGUGACACGAGAGACAAAUGGC	2	0	transposon	LINE	Rt1b
3431	CCGAACUUGAGAUCGCCCCUGGACGUACC	2	2	coding gene		CG42398-RA|SteXh:CG42398
3432	ACGAGCGCACGUGUUAGGACCCGA	2	no match			
3433	CACCAAGAUAAUCAACCCGUGUUA	2	0	transposon	LINE	Cr1a
3434	AGAUUCUUGAGAAUUUGGACGCCGUU	2	0	transposon	LINE	Cr1a
3435	ACCCUGAACGCGCCCGAUCUCG	2	no match			
3436	UACGAAUUCAACGAGGUAUGGCA	2	0	repeat		SAR_DM
3437	UAAACCAUUAAUGCUGUGCCCAAA	2	1	transposon	LTR	HMS-Beagle
3438	ACUACUGGACCAGAUGAUAGAGGGC	2	0	transposon	Unknown	RR50960
3439	CCUCUAACUGGAACGUACCUUGAGC	2	0	ncRNA		rRNA
3440	CGAGAGCAUAUAAUGCCCCGUAGGUUG	2	0	transposon	LINE	R1-element
3441	UCAGCACCAUGGACGAGAUCUCCUCCGG	2	no match			
3442	CCAGAACGUGUAUUGCUACCAACG	2	0	transposon	DNA	H
3443	GAGAUUCCCACUACGGCCGGGGGCAU	2	no match			
3444	CACCAAGAUCAUCAAACCCGUGUUA	2	0	transposon	LINE	Cr1a
3445	ACCCUGAACGCGCCCGAUCCCGUC	2	no match			
3446	UCUGCAAGAAGCCCAAACGUGAACCA	2	0	transposon	Unknown	RR50960
3447	AGAACGAUGACGAUCCCACGGCUGG	2	0	transposon	LTR	ninja-Dsim-like
3448	AUGUCGAGCUAAAGGUGACUGGGU	2	0	coding gene		CG7008-RA|Tudor-SN
3449	AAAGGACAUCAACGAGACUCCGUCA	2	no match			
3450	AACCUAAGGACGAGAGAGCUUACAGAG	2	1	transposon	LINE	G
3451	CACUUGAUCGAGUGUCUGCAGGGCAAA	2	1	coding gene		CG3180-RA|RpII140
3452	CUCAGCGAUCUAUGGGGCAAGUCAA	2	1	coding gene		CG42398-RA|SteXh:CG42398
3453	AGGAAGGCAGCAGGCGCGUAAAUUAC	2	0	ncRNA		rRNA
3454	GCGAAACAGAGAGAAUGCCACCCGAA	2	0	transposon	Simple_repeat	I
3455	CCACUUUCAUGCUUGGGAUUGUGAA	2	0	ncRNA		rRNA
3456	UGAUUUGGAUGACAGAAGAAGGUGUA	2	1	transposon	LINE	Doc2
3457	AAGGUAACGAUGGAGUUUGACGGC	2	1	transposon	LTR	flea
3458	UAAUAAUAAUAAAUAAAAGGUGACCGU	2	1	AT-chX		AT-chX-A
3459	GGAAAACCUGAAGAUGAUCCUGAGUC	2	0	transposon	LINE	Helena
3460	ACGGUACCCGGAGAAGAAGCCCCG	2	no match			
3461	CGUAAAACAAUUCAAGAUCUGGCC	2	0	transposon	LTR	blood
3462	AAAAAACCUGAAGAUGAUCCUGAGUC	2	2	transposon	LINE	Helena
3463	UCCCCUGGACCUGUACUGACGCUCAUGC	2	no match			
3464	GGGGACGUAGCUCAGUGGUAGAG	2	0	ncRNA		tRNA
3465	UAUGUAGGGGCGGUGCAUUCGAGA	2	0	transposon	LTR	Stalker4
3466	UACGAUUUCGAGUACAAAGCGAACA	2	0	repeat		RSP
3467	ACUCGCCGAAUCCCGGGGCCGAGGGAUC	2	no match			
3468	ACCAAAGCUUUUGGGCUCCGGGGG	2	0	ncRNA		rRNA
3469	UCCGAAGUGACAGACCGAAUCGCA	2	0	transposon	LTR	Quasimodo
3470	UGGUGCAUGGCCGUUCUUAGUUCGUGGA	2	0	ncRNA		rRNA
3471	GGCCCAGCGUCGUUAUGCCAACCAGC	2	0	coding gene		CG6276-RA|CG6276
3472	AGCUUCAUCCAAGCGAACUGUGCCCGAU	2	1	transposon	LINE	R1-element
3473	ACAUAACGGAACCGCUAGCAGAUCGC	2	0	transposon	LTR	opus
3474	GCGACGUCGCUUUUUGAUCCUUCGAU	2	0	ncRNA		rRNA
3475	GGUGAGCCCGUUCGGACGGUUC	2	no match			
3476	CAGUCCGAAACACACCCCCCGAUUAA	2	0	no annotation		
3477	GAUACUCGAAGAUACCACGGUGAA	2	0	transposon	LTR	ninja-Dsim-like
3478	UUGUGUCCCGGGACCCACCAAAUUGUGA	2	0	transposon	LINE	baggins
3479	AAGGUGCAGGUGGACUACAAGGG	2	no match			
3480	AGGGUUGACAGGACCAAAGACACUAGA	2	0	Su(Ste)		Su(Ste)
3481	GCCGGAGCGAGGAGGAGGACUAGC	2	0	Su(Ste)		Su(Ste)
3482	AGAGCAUAAAGACGACGAAAGUGACU	2	1	transposon	LINE	Doc3
3483	AAGCACCGUUUCCCGUCCGAUCAA	2	no match			

3484	UGCUUAGUAUCCACUUGCGAUUCCGGC	2	0	coding gene		pseudogene
3485	GGUCAACGAGGCCGAGAAGUACCGCAA	2	0	coding gene		CG4264-RA|Hsc70-4
3486	AAGUGUGUAAGACAGAUUGAUAGC	2	0	ncRNA		rRNA
3487	CGUUUCAUGAGGAAUGGGCCCAGUG	2	0	transposon	LINE	G2
3488	AUCACGAUCGAGAACGACAAGGGA	2	1	coding gene		CG18743-RA|Hsp70Ab
3489	AAGAAAUGGAAAUAGGAGUAAUCUG	2	0	transposon	LTR	rooA
3490	UCGCCGAAUCCCGGGGCCGAGGUAGC	2	no match			
3491	CCCAACUGUGAGGGGAUCCUCAA	2	no match			
3492	AGAUGUCCGGGCUGCACGCGCGCUAC	2	no match			
3493	AGAUACUCAAGACUGGACACGAU	2	0	transposon	LINE	Circe
3494	UGUGCUAGAAGACGAUUUUCGGGC	2	0	transposon	LINE	Cr1a
3495	GAGUACUUCAGCCAGACACUGGAUG	2	1	Su(Ste)		Su(Ste)
3496	CGAGCGCAACGUGCUCAUCUUCGAC	2	2	coding gene		CG4264-RA|Hsc70-4
3497	CAGGUAGGGAGUUUGACUGGGGC	2	0	ncRNA		rRNA
3498	UUGGGAACACCGCGUGUUGUUGGCCUC	2	0	ncRNA		5SrRNA
3499	GACGCCCACCCAUUGGAAGCAAACGGU	2	0	coding gene		CG6563-RA|Art3
3500	UGUUUUCAUCGUUAGACGGCUCGGGCG	2	2	no annotation		
3501	CAAACCAAUCCCAACCAAACCCAU	2	1	transposon	LINE	G2
3502	UCCGGAUGUGCUGACCCCUGCGA	2	no match			
3503	CUACGUCCGUUGGAUUAUGCCUGA	2	0	ncRNA		rRNA
3504	CAGUCCAAGAAGCUCUGCAGUUUCACCU	2	1	transposon	LTR	GATE
3505	UUCUUGCGAGGGGAUUUGUAUGGGUG	2	0	transposon	LINE	Doc
3506	GUCUUAGGGAUUGGCUAUGGUCUAGAA	2	0	transposon	LINE	G2
3507	AGUGACGCGCAUGAAUGGAUGAAC	2	1	ncRNA		rRNA
3508	UUCGAGAGAGCCUGGAGCCACCACU	2	0	transposon	DNA	1360
3509	CAGUCCGAAACACACCCCCGAUUAA	2	no match			
3510	CUCGCGAUAAUUUGGUCAUUGGACAA	2	0	transposon	LTR	opus
3511	AGCUCCGAAGCACUAAGGCCCACC	2	1	coding gene		CG18455-RA|Optix
3512	UGUUUCAUCGUUAGACGGCUCGGGCGA	2	1	AT-chX		AT-chX-B
3513	ACUCGCCGAAUCCCAGGGCCGAGGG	2	no match			
3514	GCACCCAGAGGAGAGCAGAUGAGAGC	2	no match			
3515	ACCGAAGAAAACAAAGCCUCCGCCAA	2	0	transposon	Low_complexity	F-element
3516	AAGCUUCAUCCAAGCGAACUGUGCCCGA	2	1	transposon	LINE	R1-element
3517	CGCAUAAUCAAAGCAUCCGAUGACGA	2	0	transposon	LTR	McClintock
3518	GAAACCAAAGCUUUUGGGCUCCGGG	2	0	ncRNA		rRNA
3519	UUCGUGGAGUGAUUUGUCUGGUUAAUU	2	0	ncRNA		rRNA
3520	GUCGGUCCUGAGAGAUGGGCUAGCGC	2	no match			
3521	CCAGACGUAAUACUCCACUACCCCGG	2	2	transposon	LTR	HMS-Beagle
3522	AAGGAUCCUCAACUGUGAGGCAA	2	no match			
3523	UAGGCAAACUAGGUGGGUCCCGGGGCA	2	1	transposon	LINE	baggins
3524	UGUUUCAUCGUUAAACGGCUCGGGC	2	1	AT-chX		AT-chX-B
3525	AACGCGAAGAUGAUAAACCCCAUGUGG	2	0	coding gene		CG6612-RA|Adk3
3526	CAAGAGUUCCCGUGGUAGGCAGUAG	2	0	ncRNA		7SLRNA-RA
3527	UACGACGAGAACGAGAUCAGCGCGU	2	1	transposon	LTR	frogger
3528	CCACCCUGAACGCGCCCGAUCGCGU	2	no match			
3529	CCCGAAAGCAGACGAGGUAACGUACU	2	1	transposon	Low_complexity	F-element
3530	ACGAACCGAAAAUCCUUUACUGGA	2	1	transposon	LTR	DM88
3531	AAUAGUAAGAAGUUCAACCGGUGA	2	0	transposon	LTR	invader2
3532	CCGAGGCCCUGUAAUUGGAAUGAGUACA	2	0	ncRNA		rRNA
3533	AACCAUAUCGGACGACGCAGCGCAC	2	0	transposon	LINE	Rt1a
3534	UGGCUUUGAGAGGCGCCGUAGUU	2	0	no annotation		
3535	CGGCGCGAGCCCUGGGAGGAGUU	2	no match			
3536	CAAGAUAUGACCAAAGUCAAAGACAC	2	1	transposon	LINE	F
3537	ACCAGAAAGCACCGUUUCCCGUCCGAUC	2	no match			
3538	AAAACCUUCGUAGACGACCUGCUU	2	no match			
3539	CAAGAUAUGACCAAAACAAUGCAACG	2	0	no annotation		
3540	GACAUAAACAUCCGCAGCCGCAUU	2	0	transposon	LTR	Quasimodo
3541	CUUGUACCGACGACAGAUCUUUCAA	2	0	ncRNA		rRNA
3542	UAUCCUGAGAAUCGAUUCGCAAGAAGA	2	0	transposon	LINE	Cr1a
3543	AGAUGAUUUAAUUCUAGCGUCCCAGU	2	1	transposon	LTR	1731
3544	GAUGCCGAAGUGGAGAAGGGUUC	2	no match			
3545	GAAGCUGAACGCGCCCGAUCUCGU	2	no match			
3546	CUGGAGUACAUCAGCCAUACACUGGA	2	2	Su(Ste)		Su(Ste)
3547	UGAUCGAGGCCCAGCCCGUGGACGGUGU	2	no match			
3548	UAAGAGAAUAUGCAAAGGAAGUUGGCCC	2	0	transposon	LINE	Cr1a
3549	UUAGUGCCAACGACUUUACCCGUAAAC	2	1	transposon	LINE	F
3550	GAUUACGUCCCUGCCCUUUGUACACA	2	0	ncRNA		rRNA
3551	UACGAAUGAUGUUGGCUCCAAGAG	2	0	transposon	LINE	Cr1a
3552	UGCUUGGACUACAUAUGGUUGA	2	0	ncRNA		rRNA
3553	CUUCGGGAUAAGGAUUGGCUCUG	2	0	ncRNA		rRNA
3554	CCCAAACUUAGCACCAUUGCGUAAGGAA	2	1	transposon	LINE	Circe
3555	AGCAACCGAUUUUCGAUGUGGAGACUA	2	0	coding gene		CG8517-RA|CG8517
3556	AGCUCUCGAAUGUAUGCCCGAUA	2	1	transposon	LINE	R1-element
3557	GACGCGCAUGAAUGGAUGAACGAGAU	2	1	ncRNA		rRNA
3558	GCUUUGAGAGGCGCCGUAGUUACCAC	2	0	no annotation		
3559	CAAAAGGGCAAAUGCUGACUUGA	2	0	ncRNA		rRNA
3560	UUCGGGAUAAGGAUUGGCUCUGAAGA	2	0	ncRNA		rRNA
3561	GCGAUAACGAACGAGACCUUAACCUACU	2	no match			
3562	UCUCAUCGUCGUAGAACAAGCCGA	2	2	Su(Ste)		Su(Ste)
3563	GAAGGGGGGCUCGAAGAUGCGGGGAA	2	1	transposon	LTR	opus
3564	GAGGUCUCCGGACGUGGUCACUGUGU	2	2	ncRNA		rRNA
3565	UGCCCCUGCCGAACCCAUGUCCCUU	2	no match			
3566	AGAGAUAUGACCACGCGGGAGGU	2	0	transposon	LTR	rooA

3567	GUGGCAUGUUUCCGGAUUCCCGACAGC	2	1	ncRNA		alphagamma-element
3568	UUCAAGAACAAACGAUACUGUUCGGC	2	1	transposon	LINE	G2
3569	GCGGCAACAAUCCUGUGAUCCGGAGA	2	no match			
3570	UCUCAUCGUCGUAGAAUAAGCCCGA	2	1	Su(Ste)		Su(Ste)
3571	AGGGCGAAACCUCUGAGAGUUGUGGC	2	0	coding gene		CG7816-RA|CG7816
3572	ACGGGAAAACUUACCAGGUCCGAAA	2	1	ncRNA		rRNA
3573	UUUGCAAAAACGAGAGCGAUAGAGUC	2	2	transposon	LTR	micropia
3574	UUUCCCGAAAUCCGAGUGAAUAUGG	2	0	no annotation		
3575	AUGAGAGACAUAUGCAGAGCGGGU	2	0	transposon	LINE	G2
3576	UGUCCUGAAAGUAGUUGGCAGAGGAAC	2	no match			
3577	GCGAUCCACUGAAGCUUAUCCUUUGC	2	0	ncRNA		rRNA
3578	GAGAGAUAUGGUAGAUGGGCUAGAA	2	0	ncRNA		rRNA
3579	UUCAAACUGAAACCGAGGAUAGCCA	2	0	transposon	LINE	Doc2
3580	AUCUAGGACUACCAUGGUUGCAACGGGU	2	0	ncRNA		rRNA
3581	UACUCGAAGACACCACGGUGAACG	2	1	transposon	LTR	ninja-Dsim-like
3582	GUCGGGGAGGCGGCAAUAUC	2	no match			
3583	CAGAGGAAUCAGGACCCGCAUUCUGA	2	0	transposon	LTR	297
3584	GCGAUACUUGAGGGGCACAGCUGA	2	0	transposon	LTR	DM88
3585	AUGGAACACAUAGGGGCUUCGAGGAAA	2	1	transposon	LTR	HMS-Beagle
3586	CGUUCAAGGACCACGCGAGCCACGCCAU	2	1	transposon	LINE	Rt1b
3587	CCCUGAACGCGCCCGACCUCGUC	2	no match			
3588	UAGACAGAAAAUAAGGGGACACGCC	2	0	transposon	LTR	opus
3589	AAUUCAAGGACAUUGCCAGGUAGGG	2	0	ncRNA		rRNA
3590	AACUGAAUGUGGAGACGUCGGCGCGAG	2	no match			
3591	UGUGAGACGAGUAAUAUGAGCGCCCAGAG	2	no match			
3592	AAAGCCUGAAGAUGAUCCUGAGUCGC	2	1	transposon	LINE	Helena
3593	GAGUCAAACAUCCAAGUACCAGAGCA	2	0	transposon	LTR	frogger
3594	GAUGUCGUAUCACCCAGGACGUCCGAA	2	1	transposon	LINE	INE-1
3595	CAUACCAGAAGUUAGAUGUGAAGAGCC	2	0	transposon	LTR	gypsy12
3596	ACGCCCGACGCGCACUGUUAUCCAGUUC	2	0	coding gene		CG4750-RA|loopin-1
3597	AAAGCAUCGCGAAGGCCCGCGGCGGGU	2	no match			
3598	UAUGGCUCUCUCAGUCGCUUCCUGGGA	2	1	ncRNA		rRNA
3599	UUGACCUAUAUUCAGCAGAUUAUCCCG	2	0	transposon	LINE	G2
3600	UAGUGGCUCAGAAAAGUAAAUAUUCG	2	0	no annotation		
3601	GACGAAAUAAACAAGCACUCGAGC	2	0	transposon	Simple_repeat	I
3602	AGAUUCUUGAGAAUUUGGACGCCGU	2	0	transposon	LINE	Cr1a
3603	CAAGCUAGACUAUGUAUCCCCGGGG	2	1	transposon	LINE	Doc
3604	ACAAACAGUACACCUACUAGCCCAUGUU	2	0	transposon	LINE	baggins
3605	CCUCAUUCAAAGCCAGCUGGAUGCC	2	1	transposon	LINE	G
3606	CCCGAAGAACAAUUUUACCUACAAGACC	2	0	transposon	LINE	G6
3607	CGAAUCAACAAAUUGCGCCGAUAU	2	0	coding gene		CG41545-RA|CG41545
3608	AAACGGGGUAGGUCCCUCCCUGGGUGGCA	2	no match			
3609	ACGACUCGGGUAUUACCUUUUCGU	2	no match			
3610	ACUUCAUAACCGAGCGACUGUUCAAU	2	2	transposon	LTR	GATE
3611	AAAUAGAUCCUGAAACCGCGUGCUUAC	2	no match			
3612	UAACUCGAAACUUUUGCCCCCCGAAU	2	no match			
3613	AAAAUCUGAAGAUGAUCCUGAGUCGC	2	1	transposon	LINE	Helena
3614	GCAACAGAGGAGAGCAGAUGAGAGCA	2	no match			
3615	CAAUACUGUAUUCGCUGGACCUU	2	0	transposon	LTR	Burdock
3616	ACAACCCUAAAAGAUUGGCUACGUG	2	1	transposon	LTR	mdg3
3617	ACCACCCUGAACGCGCCCGAUCUCGUUC	2	no match			
3618	GAGGAGGAUGAUGAUGAUGAAGAG	2	0	repeat		(CGA)n
3619	UUGGAUUGCCUCUAGACCGGAAGCUC	2	2	transposon	LINE	G
3620	UCUGACGAGAACUCGAGCCGCGUAGCCU	2	0	no annotation		
3621	GCGAACGUUGCGAUACCAGUCCUU	2	0	transposon	LINE	INE-1
3622	CGGGACAUGGUGUCGGACCGACUUGGGA	2	0	transposon	Simple_repeat	I
3623	CCCCUGGACAACCCACGUCCCUAUGCUA	2	no match			
3624	UACAAUGAUCUGAUCGGGUUUUUCU	2	0	transposon	LINE	Cr1a
3625	AGUCCAAUCCGACGAUACCAGGCUUU	2	no match			
3626	UGCACUAAUAUAAGCCCUGACGUCCGU	2	1	transposon	LTR	G4
3627	UGCUAGAAGACGAUUUUCGUGCCGA	2	1	transposon	LINE	Cr1a
3628	AACCAAGCUAAAGUACAGAUCGAA	2	0	transposon	Unknown	RR50960
3629	AGCGAAUGAUUAGAGGCCUUA	2	0	ncRNA		rRNA
3630	ACCCAUUACAUCCUAUACGGUCAUU	2	0	transposon	LTR	GATE
3631	UGCUCUCGAAUGUAUGCCCGAUAUG	2	1	transposon	LINE	R1-element
3632	GUCCGAGGCCCACUUCGAUGAGUACGA	2	0	coding gene		CG6770-RA|CG6770
3633	ACCCUGAACAAUACACUACACCGCAG	2	0	transposon	LTR	mdg3
3634	UAUCGAGAAACUUAUCGACUGAUCGAC	2	0	transposon	LTR	mdg3
3635	UGCUCUCGAAUGUAUGCCCGAUAUAAA	2	2	transposon	LINE	R1-element
3636	GGUUCCUUAGAUCGUUAACAGUUA	2	0	ncRNA		rRNA
3637	UGAUCCUGAAGCCGGUGUUCGACAU	2	1	Su(Ste)		Su(Ste)
3638	CACCAAGAUCAUCAACCCGUGUUAU	2	0	transposon	LINE	Cr1a
3639	UGAUUCAUCCAGGGAAUGUCAGCGCCG	2	0	transposon	LTR	ninja-Dsim-like
3640	CCGAACUUGAGACCGCCCCUGGAA	2	1	Su(Ste)		Su(Ste)
3641	AAUGACUGUAAAUUGAUAUACCG	2	0	transposon	LTR	G4
3642	UUGUAUGUUAGUUUCAGUUUCAAA	2	0	transposon	LTR	HMS-Beagle
3643	CGCAAGCUUAUUUCCCGGGAUGUUG	2	2	transposon	LINE	Fw2
3644	CUCGGGUCAAUAGGACUGGGCGGCGG	2	no match			
3645	UUGUCGGCUAGCUCAUUUCCCGGGAU	2	1	transposon	LINE	Fw2
3646	GCUUUCAACACCGCCAGAUGGCCCGU	2	0	transposon	LINE	F
3647	AUCCCACGGAACACCCGAGCAAUGCCACC	2	0	transposon	LINE	G2
3648	AUGUCAUAAAAUAGAUAACCUGAUUG	2	1	transposon	LINE	Cr1a
3649	GAGGCAAUCACCGAGAGCAUCAACGC	2	1	transposon	LINE	F

3650	CACGAUGAUCGAUGACGAGCUACGAA	2	0	transposon	LTR	ninja-Dsim-like
3651	UUCCGAGGAACCAACUGAUCCAGCU	2	1	Su(Ste)		Su(Ste)
3652	UUACCAGGUCCGAACAUAAGUGUGUA	2	0	ncRNA		rRNA
3653	UGCGAGGAGGCCCGUAUUACUGUCACA	2	0	transposon	LTR	ninja-Dsim-like
3654	CCCUCAAACACAGCAGCCGAUUCCGACC	2	0	transposon	LINE	G2
3655	GAGAGUGGACGGGAGCGGCGGGGGCGGC	2	no match			
3656	GACCGGCAACGACGUUCCACAUUA	2	no match			
3657	CAAGCCAAUCUAGACGUACGGAA	2	0	transposon	LINE	G2
3658	CAGGUACGGACCAAUGGCACAAU	2	0	repeat		LSU-rRNA_Dme
3659	AGCUCAACAACGUGCGAGAGGUGCAGCC	2	0	transposon	LTR	ninja-Dsim-like
3660	CAACGGGAUACCUCAUACUCGAGU	2	0	transposon	LTR	micropia
3661	CAGACAGAGACGGAGCUUUCUCCAGUAU	2	1	transposon	LTR	297
3662	AGGUCUGUGAUGCCCUUAGACGUUC	2	2	ncRNA		rRNA
3663	AACGCGCCCGAUCUCGUCUGAUC	2	no match			
3664	GAAUCAGUUAACUUCCCGACAUCAGA	2	0	transposon	LTR	invader6
3665	AGAUACUGAAACGAGUUGGCAUUCUU	2	0	transposon	DNA	RR51476
3666	ACUCGCCGAAUCCCCGGGGCCGAGGGAG	2	no match			
3667	GUGGAGCCGGGGCGUGGAAUGCGAGUGUC	2	no match			
3668	CUACGAGUGGCUGGACAAAAUGGUCG	2	0	coding gene		CG2915-RA|CG2915
3669	UGCAACUCGAGCGACGCCGUCUCUGC	2	0	transposon	LTR	Quasimodo
3670	GUACAAUAGAUCCGACUGGGUAA	2	1	transposon	LINE	baggins
3671	CCCGAUGAGAAGAAUGGCCCGUAAAAGC	2	2	transposon	LINE	RR49783
3672	AACCCUACUGAUGAUGUGUUGUUGCC	2	no match			
3673	UAGAGAUGCCCGCUGUACUGUCGAAA	2	0	transposon	LTR	diver2
3674	CGCUCUCGAAUGUAUGCCCGAUA	2	1	transposon	LINE	R1-element
3675	AUAGGGACGGUCGGGGGCAUCAGUA	2	no match			
3676	CCGGAGUGGACUGCCGUUAUCAGCCC	2	0	ncRNA		7SLRNA-RA
3677	CAAGAUAUGACCAGAGCAAAGCAACGU	2	0	Su(Ste)		Su(Ste)
3678	CCCGCGGAAAGACGGAAAGACCCCAUGA	2	no match			
3679	UACGUCCGUUGGAUUAUGCCUGAA	2	0	ncRNA		rRNA
3680	CCACAGAGUACGGGGCUUACUAAAGUC	2	0	transposon	LINE	baggins
3681	CGAGAGGUGAAAUUCUUGGACCGUCG	2	0	ncRNA		rRNA
3682	ACCGGAAACAUUAGUCGAGGCGUA	2	0	transposon	LINE	Rt1b
3683	AUCCUCAACUGUGAGGGAUCCUAA	2	no match			
3684	GAUCCUCAAUGUGAGGGGAUCCUCAA	2	no match			
3685	AGCGUGUGCCUACCCUACGCCGGC	2	no match			
3686	AAAGACGACAAAUGGACUAAGCCUAAA	2	0	transposon	LTR	rooA
3687	CCAAACUUAACGUAAACACAGCUUCCGC	2	0	transposon	LTR	HMS-Beagle
3688	UACAUCAGUAAAUUACGGACCCCG	2	0	transposon	LINE	McClintock
3689	ACCGAAGAGACGCCGGCCGCUAC	2	0	transposon	LTR	G4
3690	UACAAUUACGUCACGAGCAACACAGAU	2	0	transposon	LTR	Dm88
3691	GCACACAGAGGAGAGAGAUGAGAGCA	2	no match			
3692	UAAUAAAACAACGACCCGCUACCGGAG	2	0	transposon	LTR	opus
3693	AUCCGCAGUAGGUCUCCAAGGUGAA	2	2	ncRNA		rRNA
3694	GCGAUCAGAUACCGCCCUAGUUCU	2	0	ncRNA		rRNA
3695	GCCGAAUCCCGGGGCCGAGGGAGCGA	2	no match			
3696	ACCCUUGAUAGACGGUUGACUUUCG	2	1	transposon	LINE	Rt1b
3697	AGCCGAACGUAGUCUCUCGAGAGCGA	2	1	transposon	LTR	invader1
3698	UGCUUCAUCGUUAGACGGCUCGGG	2	1	AT-chX		AT-chX-B
3699	CCCGAAAAAAUCCCCCGGCUGCGAC	2	0	transposon	Low_complexity	F-element
3700	CACACAAGGUGGUCAUGAUUACGG	2	0	coding gene		CG6230-RA|CG6230
3701	UCGUCUGUGAAGGGAUCCUCAA	2	no match			
3702	GCGUUCGUUGACGUUGAUGGCUGC	2	0	transposon	LTR	HMS-Beagle
3703	CUGGUUGAUCCUGCCAGUAGUUAUAUG	2	0	ncRNA		rRNA
3704	CGGGAACACCGCGUGUUGUUGGCCU	2	1	ncRNA		5SrRNA
3705	CCCGAAAAAUCCCCCGGCUGCGAC	2	1	transposon	LINE	F
3706	GCUCCGAAAACAAGGCCCGUCGCUAA	2	0	transposon	LINE	G6
3707	ACCACCCUGAACGCGCCCCGAUCUCGU	2	no match			
3708	AGAGCGUCCGACGAAUACCCUGGGGG	2	no match			
3709	AGCAACUGAACCAAGAAAAUCACCUGC	2	0	transposon	LINE	Doc
3710	AAAGCCGAGCACUACUAAUUGCCC	2	no match			
3711	UUUUCAUCGUUAGACGGCUCGGGC	2	1	AT-chX		AT-chX-B
3712	UCCUGGGUCACACGGAGGCCGACG	2	0	coding gene		CG3356-RA|CG3356
3713	CUGAUGAAAAAUAUGUUGCCGUUU	2	0	transposon	LINE	TART-A
3714	CCGAACUUCAUACAGAUGCCUCUAA	2	0	transposon	LTR	flea
3715	UGGGCCCGAACCCUUCUGGUGUUUA	2	2	transposon	LINE	baggins
3716	ACCAACGGGAUACCUCACACUCGAGUAA	2	1	transposon	LTR	micropia
3717	GAUGUGUUGUUGCCAUGGCAAUCCUGC	2	no match			
3718	GCGUAUAACAACUACUUCUCGGUGA	2	0	transposon	LTR	accord
3719	AGUAACUUCGGGAUAAGGAUUGGCUCU	2	1	ncRNA		rRNA
3720	GACUGGCGGAUAACCCGGUUCAAGUU	2	0	coding gene		CG9970-RA|CG9970
3721	GGCAACUGUGGGGGGAUCCUCAA	2	no match			
3722	UUCCGAGGAACCCAUUGAUCCAGCUGC	2	0	no annotation		
3723	ACGACGCGAGAGGUGAAAUUCUUGGACC	2	0	ncRNA		rRNA
3724	ACAGAGAGAAGACUAACCCGACGAGA	2	1	no annotation		
3725	UCCUGUAGUACGAGGUCAUGCCUAGAAA	2	0	transposon	LTR	opus
3726	GCCACCGAGAUAUGAGGACGACGA	2	0	transposon	LTR	ninja-Dsim-like
3727	UAGGACUUAAAGAUGAUGAACCCAUU	2	1	transposon	DNA	NOF
3728	AUGCCCGAUGUCCGCGAGAUUGC	2	0	transposon	LTR	opus
3729	GAGACCCGUCGCCGCGCUCUCCCCCUC	2	no match			
3730	UACAUAUGGUUGAGGGU	2	0	ncRNA		rRNA
3731	ACAUCACAUAGAAGGAAGAUCCAGCGA	2	0	transposon	LINE	G2
3732	GGAUCGUGCUGCGGAUAUUACUGCU	2	0	coding gene		CG17945-RA|Mst84Dc

3733	AUCCCAACUGUGAGGGGAUCCUCAA	2	no match			
3734	GAAUGAGUUAAACGAGAAUGUAGAGCA	2	0	transposon	LTR	micropia
3735	AUGGAACACAUAGGGGCUUCGAGU	2	1	transposon	LTR	HMS-Beagle
3736	AUCGUCUAGUGGUUAGGACCCCA	2	0	ncRNA		tRNA
3737	CAGGAUUUUAAUUUUCAACCCGAACA	2	0	transposon	LTR	flea
3738	GAAGGCCCGCGGCGGGUGUUGACGCGAU	2	no match			
3739	GCACACAGAGGAGAGCACGAUGAGAGCA	2	no match			
3740	UAGAUGUCCUGGGCUGCACACGCGCUAC	2	1	ncRNA		rRNA
3741	GAAACCAUCAGACGCCUCCAGAGGAAA	2	0	transposon	LINE	G
3742	GCCGGUGAUCGAUAGUUCCUCUGGC	2	0	repeat		trf
3743	GUCAUGCGAGGCGAACCGGGUGAACUG	2	no match			
3744	UCGGAAAUUAAACGAGAAGACUGAC	2	0	transposon	LTR	INE-1
3745	UCUCAUCGCCGUAGAACAAGCCCGA	2	1	Su(Ste)		Su(Ste)
3746	CACCAAGAUAAUCAACCCGUGUU	2	0	transposon	LINE	Cr1a
3747	UUCGUUCCUCGGGAUCAAGGGCAA	2	0	coding gene		CG40635-RA|CG40635
3748	AGCGCCGCGGAGCCUCGGUUGGCCU	2	no match			
3749	UCUGAAACAAGAAGAGGACGACGAUC	2	0	transposon	Low_complexity	F-element
3750	AACGACACGCUUUGUUUUACCCA	2	0	transposon	DNA	RR51476
3751	AAGGGCUGGGUCGGUCGGGCUGGGGCG	2	no match			
3752	UAUCCGUGAACUACAUCCCAACGACC	2	0	no annotation		
3753	UGGGGACACCGCGUGUUGUUGGCCU	2	1	ncRNA		5SrRNA
3754	UACCACCCUGAACGCGCCCGAUCUCGUC	2	no match			
3755	AAUGAUAUGCACCGAAACUAUAUGAG	2	0	Su(Ste)		Su(Ste)
3756	AAGGGCCUAGAGGAUCUCUGUUGC	2	0	coding gene		CG12700-RA|skpD
3757	GACUUAUAUCCGAGGCCCUGUAAU	2	1	ncRNA		rRNA
3758	UUCACUAUUAAUGGAGACCGCUAUGU	2	0	transposon	DNA	Tc1-2
3759	GCGAUCAGAUACCGCCCUAGUU	2	0	ncRNA		rRNA
3760	CGAGAUCAAACCGCCGAGCUGCAACAA	2	0	repeat		(TTG)n
3761	GCAAUCAUCCAAGGCUGAGAUCCG	2	0	transposon	LTR	mdg3
3762	GUGAGGCAGGAACCGUACCCCAAAC	2	no match			
3763	ACUCGCCGAAUCCCGGGGCCGAGUGA	2	no match			
3764	UCUGCUGGCCUUCGAGUUGUACUA	2	0	coding gene		CG12202-RA|Nat1
3765	CAAUCUCAGAUAAAAACCCGAACGUAG	2	0	transposon	LTR	gypsy4
3766	GGCAAAUUAGAUCCGUAACUUCGG	2	0	ncRNA		rRNA
3767	GCGUAUUACCGGUGGAGUUCUUAUA	2	0	ncRNA		rRNA
3768	UCCCCUGCGUAGGCGCUGAGCUCUUA	2	0	transposon	LINE	R1-element
3769	AUUGGCGAGUGACCGAUAGCGAAC	2	no match			
3770	UCACCAAGCGUUGGAUUGUUCACCC	2	0	ncRNA		rRNA
3771	UAACCGAGAAACGUGUUGACAUCUG	2	0	transposon	LTR	invader2
3772	UCGCCGAAUCCCGGGGCCGAGGGAGUC	2	no match			
3773	UGAGACUACGAACGGAUUGCGGAC	2	1	no annotation		
3774	UGCACCGACAAAGGAGUAUUGCCAGA	2	0	transposon	LTR	gypsy8
3775	GCCAAUUACAUUAGACGAGACGGCCCU	2	0	transposon	LTR	opus
3776	GACCUCAGAUCAGACGUGGCGACCCG	2	no match			
3777	UAACGACCCAUUUCUGUUACUUUG	2	0	repeat		trf
3778	UUCCGAGGAACCAAUUGAUCUAGCUGC	2	1	Su(Ste)		Su(Ste)
3779	UCUAACAAAAAUUCGAGAGAGCCUGGA	2	0	transposon	LINE	Cr1a
3780	ACCGAGAAAGCAGUAGCGCUGGGUU	2	no match			
3781	CGCCGAAAGACCAAGGGUUCCUGCG	2	no match			
3782	UUACGAGCUCAGCGAGGUAUGACAU	2	0	repeat		SAR_DM
3783	UUCGAGAGCAUAUAAUGCCCCGUA	2	0	transposon	LINE	R1-element
3784	CCUGCAGAGAACGCACCCGCUUGUA	2	1	transposon	LINE	TART-A
3785	GUCAAUAAUAAUAAAUAAAAGGUGACC	2	1	no annotation		
3786	AGCUUCAUCCAAGCGAACUGUGUCCCGA	2	no match			
3787	AGAAGCACCGUUUCCCGUCCGAUC	2	no match			
3788	UACGAGGAAGCAGUGGAGGAUUGCU	2	0	transposon	LTR	ninja-Dsim-like
3789	AAUAGUGUGUUAGAGAGAGAUGAC	2	0	transposon	LTR	Max
3790	ACACAUACAAAUAACAAUACUGCCAUGC	2	0	ncRNA		ncRNA
3791	GGUUCCGGGGGAGUAUGGUUG	2	no match			
3792	AUCCUCAACUGUGAGGGGAUCCUCAG	2	no match			
3793	UUGGGCGGCGAGGACUUUGACAACCG	2	0	coding gene		CG18743-RA|Hsp70Ab
3794	AGCCAUGAGAAGAGGAAGAUGAUCGAA	2	0	transposon	LTR	Dm88
3795	CGAGAGUGGACGGGAGCGGCGGGGGCG	2	no match			
3796	UGGUUAAUUCCGAUAACGAACGAGAC	2	0	ncRNA		rRNA
3797	UAAGAUCGGACUCGAGCCUUAGCGA	2	0	transposon	LINE	Rt1b
3798	GCUGAACGUAGUCUCUCGAGAGCG	2	1	transposon	LTR	invader1
3799	AGAGCAGAGAACCCUCUAAUUCGGCC	2	0	transposon	LINE	baggins
3800	AUUCGAUUCAUGGUGUGACUACA	2	0	transposon	LTR	springer
3801	UCAACUGUGAGGGGAUCCUCUAA	2	no match			
3802	ACAAAAGGGUAAAAGUCCCCUUGAU	2	no match			
3803	UGCGAGAGGUCCCGGGUUCAUUUCCC	2	1	ncRNA		tRNA
3804	UGAGAUAUGCUAGCAGACUGGAGU	2	1	transposon	LINE	G2
3805	UGGAAAACCUGAAGAUGAUCCUGAGUC	2	0	transposon	LINE	Helena
3806	CAUGCAAGGGAACGUGAGCUGGG	2	0	ncRNA		rRNA
3807	CCGCUGGCGGUGUGAUGAGCGUGUU	2	0	coding gene		CG4264-RA|Hsc70-4
3808	UGACUCUCGAAUGUAUGCCCGAUAU	2	no match			
3809	UAUAACGACCGGUAGACCAGCACUAA	2	0	transposon	LTR	McClintock
3810	AUCUCACGGAACACCCGAGCAAUGCC	2	1	transposon	LINE	G2
3811	UUCCGAGGAACCAACUGAUCCAGCUGCU	2	1	Su(Ste)		Su(Ste)
3812	AGGCGGGGCUAGUGACUGGAGUGA	2	no match			
3813	ACAACAAACAACACAUACCCGCUGGCCU	2	1	transposon	LTR	GATE
3814	GCUUAUCCUUUGCUUGAUGAUUCGA	2	0	ncRNA		rRNA
3815	GCCGAACGUAGUCUCUCGGGAGCG	2	1	transposon	LTR	invader1

3816	AAGAACGAAAGUCGGAGGUUCGAA	2	2	ncRNA		rRNA
3817	GAGUCCAAGAGAUUUGCUCCCGAGGAGA	2	0	coding gene		CG31359-RA|Hsp70Bb
3818	CAAGAGAUGUUGUUGUCGAUGAAA	2	0	transposon	LINE	Cr1a
3819	UGCUCUCGAAUGUACGCCCGAUA	2	1	transposon	LINE	R1-element
3820	GCGCUAGUCCCAAGACCCAAAUCAGCC	2	0	coding gene		CG3330-RA|CG3330
3821	GGAAAAGAAACCAACCGGGAUUGCCUU	2	no match			
3822	GCUCCUGAAGCAGGACUCGCAGAUCCUC	2	1	coding gene		CG9074-RA|Mst57Da
3823	AACCAAGCUAGAGUACAGAUCGAA	2	0	transposon	LINE	jockey
3824	ACCCUGAACGCGCCCGAUCUCGUCUGAU	2	no match			
3825	UCAACCAGCACGACAAGGCCCAUACGA	2	0	transposon	LINE	G2
3826	ACCAAGAUCAUAAGCCCGUGAUAU	2	0	transposon	LINE	Cr1a
3827	CAGAACGUCAUCGAACCCUAGAGGC	2	0	transposon	LTR	GATE
3828	AACAGAGAGAAGACUCACCCGACGAG	2	0	no annotation		
3829	AGCCAGGAAAGAGCAGAGAACCCUC	2	0	transposon	LINE	baggins
3830	CAAGCCAAAGAGGGAGCCCGGAUC	2	0	transposon	LTR	flea
3831	ACCAAGGCGCUUGAGAGAACUCGGGU	2	no match			
3832	AAGACUACUGCCGCCUAUGCUCCC	2	0	transposon	LINE	R1-element
3833	AAGAAUACUAUUUGGGUUAAAC	2	0	ncRNA		rRNA
3834	GGAACCUCAACUGUGAGGGGAUCCUCAA	2	no match			
3835	UGCUCUCGAAUGUGAGGGGAUCCUCAA	2	no match			
3836	AUUUGACAGAGUGUGGCUCGAUGGA	2	0	transposon	LINE	Doc
3837	ACCGUGGAGACCGGGGUUCGACUCC	2	no match			
3838	GCAAUCGUUUUUAGACGAGUAUUCCGA	2	0	transposon	LTR	Rt1c
3839	AUUCCGAUAACGAACGAGACUCAA	2	0	ncRNA		rRNA
3840	UCUCAUCGUCGUAGAACAAGCCCGGG	2	1	Su(Ste)		Su(Ste)
3841	GGCUCUCUCAGUCGCUUCCCGGG	2	0	ncRNA		rRNA
3842	AACAUCGAGAAUACUGCACAGCC	2	0	transposon	LINE	Doc
3843	UCACAUCGUCGUAGAACAAGCCCGA	2	1	Su(Ste)		Su(Ste)
3844	AAGACAACUAUCCGAUACCGCUUAUC	2	0	transposon	LTR	diver2
3845	ACCUGGCAGACUGGAAUGGACCGC	2	0	transposon	Low_complexity	F-element
3846	GGCGGGUGUUGACGCGAUGUGAUUC	2	no match			
3847	GCGAGAGGUCCCGGGUUCAUUUCCC	2	1	ncRNA		tRNA
3848	GCCAAUUACAUUAGACGAGACGGCCU	2	1	transposon	LTR	opus
3849	CUUCGGGAUAAGGAUUGGCUCUA	2	1	ncRNA		rRNA
3850	GACAGACGGACAUGACUAGAUCGAC	2	0	transposon	LINE	INE-1
3851	UUCGCAAAAAGCACGGAACGAUCGAA	2	1	transposon	LINE	Doc2
3852	ACAUUCAUAACCGAGCCACUGUUCAAUC	2	1	transposon	LTR	GATE
3853	ACCGAAGAGACGCCGGCCGCUACGU	2	0	transposon	LTR	G4
3854	AGAUGUCCGGGGCUGCACGCGCGCUA	2	1	ncRNA		rRNA
3855	UACUCGAAGACACCACGGUGAAAGU	2	0	transposon	LTR	ninja-Dsim-like
3856	GUCCCGCUGGUCUAGGGGUUAUGAUCCU	2	no match			
3857	CAGAUCAACAAAGUACGAUGGGAACCA	2	no match			
3858	UUCUGCCCGAACCCUUCUGGUGUGU	2	1	transposon	LINE	baggins
3859	UACGAAAUAAGCUGUCUGAGUUAA	2	0	coding gene		CG40084-RA|CG40084
3860	GUAAAGGCUUACCAAGGCGACGAU	2	no match			
3861	ACCGAAUUAAUCGCGAGGCCGCAAGA	2	1	transposon	LTR	mdg3
3862	CAAGAGUCAAGCCGAUUAUAGUUUAG	2	0	transposon	LTR	diver2
3863	AAAAUCUCGCGCCGGGCCGUACCCAU	2	no match			
3864	AGAGGAGAAGGUGGCUUUGAGAGG	2	0	no annotation		
3865	UGCUCUCGAAUGUAGUGUCCCGAUA	2	no match			
3866	UGCUACUCGAAUGUAUGCCCGAUAU	2	no match			
3867	GUCGAGAAGAAGAUGAAGCGCAA	2	0	transposon	LTR	ninja-Dsim-like
3868	UACUUGUUCCCGGAUAGUUUAGUUAC	2	no match			
3869	UUGGGCCCGAACCCUUCUGUGGUG	2	0	transposon	LINE	baggins
3870	UGUUAUAAAAUCCUACGAUCGGGAGA	2	0	transposon	Unknown	RR50960
3871	CGAGCGUGAUGACUCCUGGGGCAU	2	0	coding gene		CG10859-RA|CG10859
3872	GAUACUCGAAGACACCACUGAUGCUGC	2	0	transposon	LTR	ninja-Dsim-like
3873	ACGACCAUGAGAGUGCAGCUGAGGU	2	1	transposon	LTR	1731
3874	AGCCAAGAGGAUACUACGAUGGUGCC	2	no match			
3875	UCCUUCGAGCGGGAUCACCUGGAU	2	1	transposon	LTR	GATE
3876	UUUUUCAUCGUUAGACGGCUCGGGC	2	1	AT-chX		AT-chX-B
3877	GCGCCGCGGAGCCUCGGUUGGCCUCGGUA	2	no match			
3878	UACUCGAUCAAAAGGCCCUGGCAU	2	1	no annotation		
3879	AGCGGGGCCUCACGAUCCUUCUGACCUU	2	no match			
3880	ACUCGCCGAAUCCCGGGGCUGAGGGAGC	2	no match			
3881	GAUCGUCGAAUUUGGGUAUAGGGGC	2	no match			
3882	CCUACGAGAUGGAAGCUUAUAUCGC	2	0	no annotation		
3883	ACCCUACGGAUGAUGUGUUGUUGC	2	no match			
3884	UACUCUCGAAUGUAUGCCCGAUAU	2	1	transposon	LINE	R1-element
3885	AGCUUCAUCCAAUCGAACUGUGCCCGA	2	1	transposon	LINE	R1-element
3886	GUGUCAAAACCCGGUGGACUGCCGU	2	2	transposon	LINE	R1
3887	GGAUUGACAGAUUGAGAGCUCUUUCUU	2	no match			
3888	CUCCCACUAGAGAGUUGGCCCAGC	2	0	coding gene		CG10077-RA|CG10077
3889	ACGAAAUCGAUGUGUUACUGCUCA	2	0	transposon	LINE	X
3890	CUUGUUCCCCGGAUAGUUUAGUUACGUG	2	1	ncRNA		rRNA
3891	UAGUCUGGUCGCGGGUAGGAGCGGGG	2	0	transposon	LTR	invader1
3892	AAUUCCGUAAACGAAGCCCAAAGUUCA	2	0	transposon	LTR	Max
3893	AAUUCCGUAAACGAAGCCCAAAGUUCG	2	0	transposon	LTR	Rt1c
3894	UCGAACGAUAAACGGCGCUAACCA	2	0	transposon	LTR	GATE
3895	UCCUGGCAGAGACGGCGUCGCUCGAGU	2	0	transposon	LTR	Quasimodo
3896	UAUUGAAUAACGUACUCUUCGUCCCAGA	2	0	transposon	LTR	297
3897	UUCCGGAGAGGGAGCCUGAGAGACG	2	1	ncRNA		rRNA
3898	ACAAACAGAGUUCCCGAAUGAGAG	2	0	transposon	LINE	Cr1a

3899	ACCCUGAACGCGCCCGAUCUCGUUC	2	no match			
3900	UGCAUUUCCAAAAUUGACCUCAAAG	2	0	transposon	LINE	Circe
3901	ACCACUUUCAUGCUUGGGAUUGUGA	2	0	ncRNA		rRNA
3902	GGCGAGAUGGGUGACAGCCCCGUACGGGA	2	no match			
3903	ACGAGAGAGAUCCGGAUGUCUAUGA	2	0	transposon	LTR	ninja-Dsim-like
3904	UCUCAUAGUCGUAGAACAAGCCCGA	2	1	Su(Ste)		Su(Ste)
3905	GGAUCCUCAACUGGUGAGGGGAUCCUCAA	2	no match			
3906	UUAUUCGUUACGUAGAUUUAGAGUU	2	0	transposon	Unknown	RR50960
3907	CUGGUCGUUAAAGGGAGUUUUGGC	2	0	transposon	LINE	R1
3908	GCGACUCGGGGCCUGUCUAGGAACA	2	0	transposon	LTR	opus
3909	AUUCCCCAUGAACGAAAUUCCCAGU	2	no match			
3910	AUACCCUGGUAGUCCACGCCUUAAACG	2	no match			
3911	UGUUUCACCGUUAGACGGCUCGGGC	2	1	AT-chX		AT-chX-B
3912	CAGUCCGAAACACACCCCCCCGAUAAU	2	1	no annotation		
3913	AAAAACCUGAAGAUGAUCCUGAGUCGA	2	2	transposon	LINE	Helena
3914	ACCACCCUGAACGCGCCCCGAUCUCGUC	2	no match			
3915	CGUUUCAUCGUUAGACGGCUCGGG	2	1	AT-chX		AT-chX-B
3916	ACAUUCAUAACCGAGCGACUGUUCAAA	2	1	transposon	LTR	GATE
3917	GACCUCAGAUCAGACGUGGCGACCCGC	2	no match			
3918	GGCGAGCGAACCGGGACCAGGCCAGU	2	no match			
3919	AACGCGGUCACCGCGUUGGUCACCGGGUGGCCC	2	no match			
3920	UAUUGCACUUGAGACGGCCUGA	2	0	ncRNA		mir-312-RA
3921	UGAGGUCUCCGGACGUGAUCACUGUGAC	2	0	ncRNA		rRNA
3922	ACCACACCCGGGGACAAUGACCUUUU	2	0	coding gene		CG8498-RA|CG8498
3923	UGCUAAGGGAUCCAGUUGCGAUAAA	2	0	transposon	DNA	1360
3924	AAAUCCUCAACUGUGAGGGGAUCCUCAA	2	no match			
3925	ACCACCCUGAACGCGCCCGAUCUUGUC	2	no match			
3926	AAGAAACGAAAGUGCAGACCCAGG	2	0	transposon	LINE	Doc3
3927	AAAUCUGGAUACUAGGUAUCGACGCCGA	2	0	no annotation		
3928	ACCGGAUGAAUAACGAGAACCUACUC	2	0	transposon	LTR	gypsy
3929	GGGGAACUACUGCUCGAGAUAGAGG	2	0	transposon	LINE	Quasimodo
3930	UGGGAAUCCGAGGUGCCGUAGAU	2	no match			
3931	AACCACCCUACGCCAAGACUCGUU	2	0	transposon	LINE	G
3932	AGUACCUCGACCAGACGGUGGCCCCC	2	0	coding gene		CG11591-RA|CG11591
3933	GAUCUGAAUAAUUCUGCGAACGAGGA	2	0	coding gene		CG4916-RA|me31B
3934	GGGGAUGUAGCUCAGAUGGUAGAG	2	0	ncRNA		tRNA
3935	GGCACAAUACUUGUUCGAGCGAACA	2	0	ncRNA		rRNA
3936	UUACUGACGAAUUUAGUUACAAGCAGA	2	0	no annotation		
3937	AUCAAAAACCAUUCUGAUUUCAACGA	2	0	no annotation		
3938	AUGUGAACAGCAGUUGAACAUGGGUC	2	no match			
3939	AGUGACGCGCAUGAAUGGAUUAA	2	0	ncRNA		rRNA
3940	AAUGAUGUAGAUCGUGGUUCGUGG	2	0	no annotation		
3941	AAGAGGGACGGCCGGGGGCAAUCGU	2	no match			
3942	UUUCGAGAAGCGAACCCGGCGAACUGAA	2	no match			
3943	UGCGAGAGGCCCUGGGUUCAUUUCCCA	2	no match			
3944	GAUACUCGAAGACACCACGGUGAAAG	2	0	transposon	LTR	ninja-Dsim-like
3945	GCAGCCGGGCGUGGAAUGCGAGU	2	no match			
3946	AAGGACCGCAUCCAUGUACCGGAGG	2	0	transposon	LINE	R1-element
3947	CUUGAUAUGACCAGCUUGGACCGAA	2	0	Su(Ste)		Su(Ste)
3948	GGCGGGGAAGGUGGCUCGGGGGGC	2	no match			
3949	AUGUCUAUGAGAUGAACGUAAUGACG	2	0	transposon	LTR	ninja-Dsim-like
3950	GCUGAGAAAAAAACGAAGUAUAACGAAA	2	0	transposon	LINE	Doc
3951	UCCGGCGAUAAACUUCUUGUCCUUCG	2	0	transposon	LTR	G4
3952	UUCCUUAGAUCGUUAACAGUUACUUGG	2	0	ncRNA		rRNA
3953	UAUAUGCUCUCGAAUGUAUGCCCGAUA	2	0	transposon	LINE	R1-element
3954	AACUAAGACAGGGCAUUACGAGUAUA	2	0	transposon	LTR	McClintock
3955	UAGGCGGAUCGUUAAGUCAGAGGU	2	no match			
3956	GAGAGAACUCGGGUGAAGGAACUAGGC	2	no match			
3957	CAAGAUAUGACCAAAGCAAAGCAAC	2	0	Su(Ste)		Su(Ste)
3958	UUCGGUCCUGGGCGAAAUCACCA	2	no match			
3959	UGAUCUCGAAAGCUAAGCAGGGUC	2	no match			
3960	CCCGGAAGCGAUCCCACUGCUGAAG	2	1	AT-chX		AT-chX-A
3961	CUGACCUGUAUUCGGAUAAUGGA	2	0	transposon	LTR	3S18
3962	ACUGAGACACGGCCCAGACUCCUACGGGA	2	no match			
3963	UGCUGUUUACUUCGAGGACAAGGU	2	0	transposon	LTR	flea
3964	UCAAGGUAGUAACUAUGACGAUUUGA	2	0	transposon	LTR	flea
3965	GGACGCGUAGCUCAGCGGGAGAGCA	2	no match			
3966	GCAUUCGAGACCCAACGGCACAUCA	2	1	transposon	LINE	Rt1b
3967	CACCCUGAUUGCAGACGCAGACGAGC	2	0	coding gene		CG34222-RA|CG34222
3968	CGGUAAGACCCUGAUCGAUGCCCU	2	1	coding gene		CG8280-RA|Ef1alpha48D
3969	AAGCCGGUGAUCGAUAGUUCCUCU	2	0	repeat		trf
3970	CGGAGAUGGGGUCUUAUGGCUGGAA	2	no match			
3971	CUCUUCUUCAUUUCUGUUCAAGGGC	2	0	no annotation		
3972	GAAACCAAAGCUUUUGGGCUCCGGGG	2	0	ncRNA		rRNA
3973	GAGAGCGAGAAGAUGUCCGAGA	2	no match			
3974	UCGGCCCGAACCCUUCUGGUGUGUA	2	1	transposon	LINE	baggins
3975	CAAGCUAGACUAUGUAUCCCCGGGCAU	2	2	transposon	LINE	Doc
3976	UGGGAGGAGGAAGACGGUUACC	2	0	transposon	LTR	diver2
3977	AGAGACAAUUGACGGUACCGCACACC	2	0	transposon	LTR	Dm88
3978	CAACUCGGAUGUUAUCAUCAUGCA	2	0	transposon	LTR	flea
3979	CAUACCAACAGUAGACAAUGUUCCGA	2	0	transposon	LTR	accord
3980	UACUACAGGAAGUUCAUUCAGGACUA	2	0	transposon	LTR	opus
3981	UCCGAAGACAACGAGGAGCAGAG	2	0	transposon	LTR	1731

3982	GCAACGAGGCAGCACGGGGUAUC	2	0	transposon	LINE	BS
3983	GGAUCCUCAACUGUGAGGGCAUCCUCAA	2	no match			
3984	AACGAGUCCAAGCCAGGCAGCUAU	2	0	transposon	LINE	Rt1b
3985	UGACCGAACUUGAUUAUUUAGAGGAAG	2	0	ncRNA		rRNA
3986	GGGGAACUACUGCUCGAGAUAGAGGG	2	0	transposon	LINE	Quasimodo
3987	AACGCGAAGACAGAUUCCUGGGUGG	2	0	coding gene		CG8590-RA|Klp3A
3988	ACGGCCAUACCACCCUGAACGCGCC	2	no match			
3989	AAUAACUCCAGCAGAGACCGUACCCGA	2	no match			
3990	CUCAGAUCAGACGUGGCGACCCGC	2	no match			
3991	ACGGGAAAAACGCCGUGCGCAUUUUGAC	2	0	transposon	LINE	baggins
3992	AUGUUUCAUCGUUAGACGGCUCGG	2	1	AT-chX		AT-chX-B
3993	UCCUGAAGCCGGUGAUCGAUAGUU	2	0	repeat		trf
3994	CUGAAUGAUGGACUGAAAUGGUCUGG	2	no match			
3995	GCAAAGAGAACGCGACCAACUGAA	2	0	no annotation		
3996	CCGAACUUGAGACCCCAGCCGCUCG	2	1	no annotation		
3997	GGAUCCUCAACUGUGAGGUCAA	2	no match			
3998	CCAGCUUGGACCGAACACUUUGAACC	2	0	Su(Ste)		Su(Ste)
3999	CGCAAACCCAAAUCGACAAUGCACGA	2	0	repeat		trf
4000	AUCCUUCGGACAAAGGCACUGAGUUU	2	0	transposon	LTR	Quasimodo
4001	ACGGAAUGUAACCGAGCUCUGAGU	2	0	transposon	Simple_repeat	I
4002	GCGUUCGUUGACGUUGAUGGCUGCU	2	1	transposon	LTR	HMS-Beagle
4003	CCGCUUGGGAACACCGCGUGUUGUUGGC	2	0	ncRNA		5SrRNA
4004	CAAGAUAUGACCAAAACAAAGACAUA	2	2	transposon	DNA	1360
4005	AGCAGAGAACCCUCUAAUUCGGCCCGAA	2	0	transposon	LINE	baggins
4006	UCCGAAUCAAAUGCUACGUAACCAAGCCC	2	0	transposon	LTR	diver2
4007	UCGAAAUGCCGGGGCGAGGAGGAGGAC	2	1	Su(Ste)		Su(Ste)
4008	GGAGUUUGACUGGGGCGGUACACCUGU	2	2	ncRNA		rRNA
4009	AAAAAAGUGAAGAAGACGUCGACG	2	2	no annotation		
4010	UAAUGAAACAUGAUGCUUUCGCCGA	2	0	no annotation		
4011	AAGACAACAGUGCAGACGGCGCCCGC	2	1	transposon	LTR	diver2
4012	CCACCCAGAGAAGUACCAAAAAGC	2	0	transposon	LINE	G6
4013	AUGACAGACAUCCCACACAUCCGGA	2	0	transposon	LTR	HMS-Beagle
4014	UGCCGUGAUCGUCUAGUGGUUAGGAC	2	1	ncRNA		tRNA
4015	GAACUUGAGAUCGCCCCUGGACGA	2	0	coding gene		CG42398-RA|SteXh:CG42398
4016	ACUACUUUCAGGACACGUUCAACCAGA	2	0	Su(Ste)		Su(Ste)
4017	ACGAGCGCACGUGUUAGGACCCGAA	2	no match			
4018	GCCGAACUUGAGAUCGCCCCUGGACGACC	2	0	coding gene		CG42398-RA|SteXh:CG42398
4019	CAAAGCAUUGUGAUGGCCCUAGCGG	2	0	ncRNA		rRNA
4020	AUCCUCUACACCCUGUUUACAGCAGA	2	0	transposon	LINE	G2
4021	ACUGAUAACGAAUGGAUGACA	2	no match			
4022	ACGAGUACUAUGAGCGCCCAGAGUGA	2	1	transposon	LTR	opus
4023	UGGAGCUAACCGAUACUAAUUGUCCU	2	no match			
4024	GCCGAAUCCCGGGGCCGAGGGAGCGAG	2	no match			
4025	ACGGGAACCUCACCCGGCCCGGACACGG	2	no match			
4026	GACGACAUAAUUGUCUUUCGACAUC	2	no match			
4027	CCGAUAAUGAACCAAGCUUGACCUC	2	0	no annotation		
4028	AUGAGGCGCUCGGUGUCGGUGAAGGCC	2	no match			
4029	GACACCGGAACAUGUGGCUGCGAGU	2	1	transposon	LTR	diver2
4030	CGAAAGUAGGUCUUAGUGAUCCGGUG	2	no match			
4031	UGACAUCAUUCAGCGAGCACAAUCAA	2	0	transposon	Low_complexity	F-element
4032	AGAUUCAUCCAAGCGAACUGUGCCCGA	2	1	transposon	LINE	R1-element
4033	ACGGACCGAAAAUCCUUUACUGGAU	2	0	transposon	LTR	DM88
4034	ACCGAAAGCAGACGAGGUAACGUAC	2	1	transposon	Low_complexity	F-element
4035	GUUAAUAAUAAUAAAUAAAAGGUGACCG	2	0	no annotation		
4036	CUGUAAAUAAUUCAGCAGCACUGACC	2	0	transposon	LINE	Doc
4037	CUUUGGCUCAGUCACACAACAUUG	2	0	transposon	LTR	micropia
4038	UCGACUCGAAAUAGUAACUACGUAA	2	0	transposon	LTR	17.6
4039	UGAGACUACAAACGGAUUGUGGACC	2	1	no annotation		
4040	AAUCGCUCGAUGUCGUGAGGUGACCG	2	0	no annotation		
4041	UACGUCAACAUAUCAGGGCAGCCCGACC	2	0	transposon	LINE	Doc4
4042	AAAGAGGGACGGCCGGGGGGCAUUCGU	2	no match			
4043	UGACCAGUUGUCCCAUCAAAUCCGU	2	0	transposon	DNA	transib2
4044	CAGCAUCAACAAGUACAAUCCCAGA	2	0	coding gene		CG3891-RA|CG3891
4045	ACCCUGAACGCGCCCGAUCUCGA	2	no match			
4046	CACCCAUGCAAGGGAACGUGAGCUGG	2	0	ncRNA		rRNA
4047	ACCCAGUUCAUUCUUGAGGCCUUCGG	2	1	transposon	LINE	G2
4048	ACAACCUAAUCACGUUCGGUGCCCG	2	0	transposon	LINE	Rt1a
4049	ACUUGUUACAAAACAUAGAUCCAAGA	2	0	transposon	LINE	Cr1a
4050	GGCUAAUGACGGUACCGUAAGAAUA	2	no match			
4051	CAACUCUGUAUGCACACCCCGUUAGCUU	2	0	transposon	LINE	Circe
4052	GCAUUCGAGACUCAACGGCACGUCAC	2	0	transposon	LINE	Rt1b
4053	AGCGACGAAGACACCGGUUCAUGCUC	2	0	coding gene		CG5658-RA|Klp98A
4054	GCAUUCGAGACCCAACGGCACGUGACUA	2	2	transposon	LINE	INE-1
4055	AACCAUAAACGAUGCCAGCUAGC	2	0	ncRNA		rRNA
4056	UUCCCUGAACUCGGAGCUCGGCA	2	0	coding gene		CG6370-RA|CG6370
4057	AACUCUAAGCGGUGGAUCACUCGGCUC	2	0	ncRNA		rRNA
4058	AUACCUAGCGUUGAAGCCCCGAUAAA	2	0	transposon	LTR	HMS-Beagle
4059	ACCUGGAGAAGGAACUUGAUGUGCA	2	1	coding gene		CG13032-RA|CG13032
4060	GCGAAGGUGGCGGAAUUGGGCUUAGG	2	no match			
4061	UUCUCAUCGUCGUAGAACAAGCCCGAA	2	0	Su(Ste)		Su(Ste)
4062	UCCUGUUGAUCAUUUGCCCGUGGU	2	0	transposon	LINE	Cr1a
4063	UAUUACGAGCUCAACGAGGUACGGCAU	2	0	repeat		SAR_DM
4064	UACGAGAAUAUACCGCAAACCACA	2	0	transposon	LTR	gypsy12

4065	UACGACGAGAACGAGAUCAGCCCGA	2	1	transposon	LTR	frogger
4066	CCCCUGGACAACCCACUAAAUUAUAGA	2	no match			
4067	UUCAAGAGAAGCAGGCGGAAUUAACGA	2	0	transposon	LINE	X
4068	UUCGGAGACAAAGUUUACAUGCUAAGGA	2	0	ncRNA		snoRNA
4069	CAUCUUCUUAAUGCAGCUGCCGAACU	2	0	Su(Ste)		Su(Ste)
4070	AUGCACGACAGAGGAAGCAGAACA	2	0	repeat		HETRP_DM
4071	UGCCCGUAAAAACGACUUGGUUGACAC	2	0	transposon	LTR	gypsy
4072	ACUCGCCGAACCCCGGGGCCGAGGG	2	no match			
4073	GGGGGCAUUAGUAUUACGACGCGAGA	2	0	ncRNA		rRNA
4074	GGAUCGGUGGUUCAGUGGUAGAAUGCU	2	1	ncRNA		tRNA
4075	CAGUGCUGGCUGAAGUACUCCAGGCCC	2	0	Su(Ste)		Su(Ste)
4076	GUGACUCUAGAUAACCUCGGGCCGA	2	no match			
4077	UGCCGGAGCGAGGAGGAGGACUA	2	0	Su(Ste)		Su(Ste)
4078	UGAUGAAACACAUUACCAGAACCU	2	0	transposon	LTR	HMS-Beagle
4079	UCGAACGAUAAACGGUGCUAACCAGC	2	0	transposon	LTR	GATE
4080	AAUAAAACAACGACCCGCUACCGGA	2	0	transposon	LTR	opus
4081	UCCGAAGUGACCGACCGAAACAGUCGCGA	2	0	transposon	LTR	diver2
4082	UCACCAAGCGUUGGAUUGUUCACCCAU	2	0	ncRNA		rRNA
4083	CGGAUCCUCAACUGUGAGGGGAUCCUCAA	2	no match			
4084	CCGAAGAAAACAAAGCCUCCGCC	2	0	transposon	Low_complexity	F-element
4085	AUGGCUCUCUCAGUCACUUCCCGGGA	2	1	ncRNA		rRNA
4086	UACGAUUGAAAAAGAACUACUAGCU	2	0	transposon	LINE	McClintock
4087	UCCGAUAACGAACGAGACUCAAAU	2	0	ncRNA		rRNA
4088	CGUAACUUCGGGAUAAGGAUUGG	2	0	ncRNA		rRNA
4089	ACGAGUAAUAUGAGCGCCCAGAGC	2	1	transposon	LTR	opus
4090	CUGUUCUGAAUUUUACGUACCGCAGU	2	0	transposon	LINE	Fw2
4091	UGUUUCAUCGUUAGACGGCUCGU	2	1	AT-chX		AT-chX-B
4092	GCCUAUAAUUUUGCGACCUUAGUCAC	2	1	no annotation		
4093	UGCGCGAAGAAGGAGGACGAGAAA	2	1	transposon	LINE	G2
4094	AUGGCUCUCUCAGUCGCUUCCCGA	2	1	ncRNA		rRNA
4095	GACGUAGAAAAACUUGGAGACGAC	2	0	transposon	LTR	diver2
4096	ACCACCCUGAACGCGCCCGAUCUCGCC	2	no match			
4097	GCAUUCAUAACGAGGUUGCGCAUU	2	0	transposon	LINE	Doc4
4098	UCUCAUCGUCGUAGAACAAGCCCGAGC	2	1	Su(Ste)		Su(Ste)
4099	AUUCGUAGACGACCUGGGGCGGGGUUU	2	no match			
4100	AUGGUGACCACGGGUGACGGGGAAU	2	no match			
4101	AACAGGGCGUGAAACCGUUAAGAGGU	2	no match			
4102	GCAGAACUCCGAUGGCGAUAGCAACU	2	0	no annotation		
4103	UCCAUGAGGACAUAACGCUGAGCGAC	2	0	transposon	LINE	Quasimodo
4104	GGGGAUGAGCUGUGGAUAGGGGU	2	no match			
4105	GAUCACCUGAUGCCGCGCUAGUU	2	0	ncRNA		rRNA
4106	UCCGGGGCUUCUCAACUACAAGC	2	0	coding gene		CG3221-RB|dgt3
4107	UUAGUUUACCGGAACGGUCUGGAAUGGCC	2	no match			
4108	GGCCGUGAUCGUCUAGUGGUUAGGACC	2	1	ncRNA		tRNA
4109	UAAGAUCAACGAGAGAUGGCCAUUAA	2	0	transposon	LTR	opus
4110	CUUAAACGAGUGUUAUUGUGGGCCGGU	2	0	ncRNA		rRNA
4111	AAAUUCAUACCUGGAUACGCCCAA	2	0	transposon	LTR	flea
4112	AGAGAAAUGAUAGAAACCCAUGUA	2	0	transposon	LTR	Idefix
4113	CAAUGAGAUAGCAUUUGGCGAGAGU	2	0	transposon	LTR	diver2
4114	CUGCCAGUAGUUAUAUGCUUGUCUC	2	0	ncRNA		rRNA
4115	UCGGAAAUUAAACGAGAAGACAUUGC	2	0	transposon	LTR	INE-1
4116	CGCCCCCGGGGAGCCUGGCGCGC	2	no match			
4117	GCUAGUUCCGAAAAAAUCAUUACCGAA	2	0	transposon	LTR	Stalker2
4118	GAACCUGAAACCGUAUGCGUAC	2	no match			
4119	AAGAGCAAUGAACGAUAUUUUAAGA	2	0	no annotation		
4120	CUGGCCGGCAGGGACCUCACCGAUUAU	2	0	coding gene		CG5409-RA|Arp53D
4121	GUGUGGCGAGGAGUGCGGUUCUUUGU	2	no match			
4122	AAAACCUGAAAAUGAUCCUGAGUCGC	2	1	transposon	LINE	Helena
4123	CAAACCAAUCCCAACCAAACCCGUCAU	2	1	transposon	LINE	G2
4124	AUGCUCUCGAAUGUAUGCCCGAUAAUU	2	no match			
4125	ACGCUUUCACCGUGGAGACCGGGGUU	2	no match			
4126	CUGUUCACAUGAACUUGGAAUUCCCAGU	2	0	ncRNA		rRNA
4127	AGCUGCCAUACUGCGAUCCACUGAAGC	2	0	ncRNA		rRNA
4128	AUGGCUCUCUCAGUCGCUUCCGGGA	2	1	ncRNA		rRNA
4129	UUCCCGGAUCCUCGACCCAUCGAGUGC	2	0	coding gene		CG4691-RA|CG4691
4130	GCUCUCAUCACGAGGUUACACCCGCGU	2	0	no annotation		
4131	CAACAAGAUACCCGUACGGCUGU	2	0	transposon	Simple_repeat	I
4132	CUGUCUCGAAAACAAUCGCAGGG	2	0	transposon	LINE	F
4133	CAGAUCAACAAAGUACGAUGGGGAACCA	2	0	no annotation		
4134	GUAUACCUAUACAUUACCGCUAA	2	0	ncRNA		rRNA
4135	CUCAACUGUGAGGGAUCCUCAA	2	no match			
4136	GCGAGCGAACGCGGAUCAGGCCAGUGCCU	2	no match			
4137	CCCAACAAUAAAAAUACUGUGAGCAGG	2	0	no annotation		
4138	CUGUCGCGAAAUUGGUAUCCCCGAU	2	1	transposon	LTR	diver2
4139	GAUCCCAACUGUGAGGGGAUCCUCAA	2	no match			
4140	UAUGUUGGCCUGGAUACGCUUGU	2	1	transposon	LINE	G2
4141	AAGGAUACUCGAAGACAUCACGGUGA	2	1	transposon	LTR	ninja-Dsim-like
4142	CAAACCAAUCCCAACCAAGCCCGUCAA	2	1	transposon	LINE	G2
4143	AGAGAGACUACAAGAACCUGGAA	2	0	transposon	LTR	HMS-Beagle
4144	UGCUUUCAUCGCUGGCGAGGAUAUGGA	2	0	ncRNA		pseudogene
4145	GGCCGAAGUUCAAGCCCGGGUAUUCUG	2	0	repeat		trf
4146	AAAGCAGUCCUGAGGUGGGCGUGG	2	0	no annotation		
4147	AGCAGAGAACCCUCUAAUUCGGCCCGA	2	0	transposon	LINE	baggins

4148	AACGCAUUAAGCAAUCCGCCUGG	2	no match			
4149	AAGGCUAAAUACUACUGACUGACCGA	2	no match			
4150	CAGGAAGAGUGCCCGGCUGAACGGA	2	0	transposon	LTR	ninja-Dsim-like
4151	AGCAAAAGCAACAAUAAGUUACCGCGA	2	0	transposon	LTR	McClintock
4152	UCCAGACGCAGCCCGGCCAGGA	2	0	transposon	LTR	ninja-Dsim-like
4153	ACGCCUGUUAGUCCGAAUACAUACGA	2	0	no annotation		
4154	CGCAAUCCGAGGAAGUAGCUGUGCCGCA	2	0	no annotation		
4155	GGAUCCUCAACUGUGAGGGGUUCCUCAA	2	no match			
4156	UUCUCAGAAAACGAAUUCGACCACG	2	0	transposon	LTR	gypsy
4157	AGUGCAUUGACUUUAUUACAUAUUA	2	0	transposon	LTR	DM88
4158	ACAAGAAGUAUCCCAGCUACAACAGU	2	0	transposon	LTR	Quasimodo
4159	AUAAAGGGCAUCUAGGCGGCCGGGUA	2	no match			
4160	GAAUUCAUGAAGAAAGAAACCGAAU	2	0	transposon	LTR	Idefix
4161	CCUGAUGAAGAAGAGAUGCCCAGU	2	0	transposon	LTR	ninja-Dsim-like
4162	GCUUUGAGAGGCGCCGUAGUUACCACCC	2	0	no annotation		
4163	GAAACAGCAUCCCUACGAUUCCUCC	2	0	coding gene		CG1076-RA|CG1076
4164	ACAACUGUGAGGGGAUCCUCAA	2	no match			
4165	GUAACCGAGAAACGUGUUGACAU	2	0	transposon	LTR	invader2
4166	GCGUUCACCGCAUUGCUCCUGAUUAAA	2	0	coding gene		CG41580-RA|CG41580
4167	AAGACAAGUAUACCGUGGCACACCAUC	2	0	transposon	LTR	micropia
4168	AAGCCGGAGGAAGGUGGGGAUGACG	2	no match			
4169	AACUCGGAGGAAGGUGGGGAUGACGUC	2	no match			
4170	UGAGUCAUAAUGUAGUUCUUGGACGAG	2	1	transposon	LTR	flea
4171	GUCCAGAGGAGGCGAGAAUUUUACAU	2	0	AT-chX		AT-chX-B
4172	AAAACGCGAAUGGCAUCCGGAUGAGA	2	0	transposon	LTR	flea
4173	UGUUUCAUCGUUAGACGGCUCGUGC	2	1	AT-chX		AT-chX-B
4174	CAAGCUAGACUAUGUAUCCCCCGGGUAA	2	no match			
4175	UGUUUCAUCGUUAGACGGCUCGGCG	2	2	AT-chX		AT-chX-B
4176	GAUGUCUAUGAGAUGACCGUAAUGAGC	2	0	transposon	LTR	ninja-Dsim-like
4177	CACAUCAACAGAAAGCGUAUCCAGGCCA	2	0	transposon	LINE	G2
4178	ACCCUGAACGCGCCCCCUCGU	2	no match			
4179	GUCCCUUAAAUCAGCACUCGUAAA	2	0	no annotation		
4180	AACGGUACGAGGUAGUUGCUUUCCGAG	2	1	no annotation		
4181	GGGAGUGGGAGCGAAAUCGGGGAC	2	0	no annotation		
4182	CGUGUCCGGAAAAUGUUUACAGUA	2	0	coding gene		CG6817-RA|foi
4183	GGACUCUCCGCCUUCGGACCCGCC	2	0	coding gene		CG3640-RA|CG3640
4184	CGUGAGAGGAUCAGCAUGCCUGGCA	2	0	transposon	LTR	mdg1
4185	AGGUGGGGAGUUUGACUGGGGCGGU	2	1	ncRNA		rRNA
4186	GACAUCAUUUCGAGACCCAUUUUAAGC	2	0	transposon	LTR	gypsy12
4187	GUCUGGUUAAUUCCGAUAACGAACGG	2	1	ncRNA		rRNA
4188	UCUCAUCGUCGUAGAACGAGCCCGA	2	1	Su(Ste)		Su(Ste)
4189	GAGGUUCGAAGGCGAUCAGAUACCG	2	0	ncRNA		rRNA
4190	AAUCGAAUGAGACCUUGGAGGAC	2	0	ncRNA		rRNA
4191	AUGAGAUAUGCUAGCAGACUGGAGA	2	0	transposon	LINE	G2
4192	GCGUAGUCGAUGGGAAACGGGUUA	2	no match			
4193	AGAUACUUUAUGACGACACUGGAA	2	0	transposon	LTR	gypsy12
4194	CCGGUAAAGCGAAUGAUUAGAGGU	2	2	ncRNA		rRNA
4195	AGAUGUCCGGGCUGCACGCGCGCU	2	no match			
4196	GACAGAAUAGAGAGGAUCCUCAA	2	no match			
4197	CAGAUCAACAAAGUCCGAUGGGGA	2	1	no annotation		
4198	AUGCUCUCGAAUGCUUGCCCGAUAUAC	2	0	transposon	LINE	R1-element
4199	ACGCGACCUCAGAUCAGACGUGG	2	no match			
4200	UCCUUCGAUUUCCCGUUUUGAGAGGCGCCGUAGU	2	no match			
4201	AUAGGGACGGUCGGGGGCAUCA	2	no match			
4202	AUCUAUAUUAUCCCAGCACUACGAAGC	2	1	no annotation		
4203	CCCAAACGCAGACAUCGAGCGGUCAAUA	2	0	transposon	Low_complexity	F-element
4204	UGCUAACAAGGCGCCGCCCGUGG	2	0	transposon	LINE	G6
4205	UUAUUAUUAUUAAGAUGGGCUUGUAGG	2	0	transposon	LINE	baggins
4206	CUGGCUUUACGGCGAGGACAUUGCG	2	0	transposon	LINE	R1-element
4207	GCAUUCGAGACCCAACGACACGUGAU	2	2	transposon	LINE	INE-1
4208	CUGCUCCGAGCCCCGCGUCGGCAUCC	2	1	coding gene		CG32479-RA|CG32479
4209	CGCAUUGAUUACGUCCCUGCCCU	2	0	ncRNA		rRNA
4210	GCAUUCAUAACGAGGUUGCGCAUUGCA	2	0	transposon	LINE	Doc4
4211	UGCGAGAGGUCCCGGGUUCAUUUCC	2	1	ncRNA		tRNA
4212	AGCGAUCUGAGUGAGUAGAGUGU	2	0	transposon	LINE	RR49783
4213	CUGGGAAACGGGGCGCGGCCGGAAA	2	no match			
4214	UCCGAAGACAACGAGGAGCAGAGCU	2	0	transposon	LTR	1731
4215	GUGGGCCGGUACUAUUACUUUGAACA	2	0	ncRNA		rRNA
4216	AACGAAGAAAAACAAAGAUUGCGCG	2	1	transposon	LTR	McClintock
4217	UGUAAUACAAGAGACCCGAAAACCGU	2	0	transposon	LTR	gypsy4
4218	AACUUCGGGAAAAGGAUUGGCUCU	2	1	ncRNA		rRNA
4219	ACCGAUAAGAGAUUGGCUAAGUUAA	2	0	transposon	LTR	accord
4220	UAUAUCAGUUAUGGUUCCUUAGAUCGU	2	0	ncRNA		rRNA
4221	AACUACUGGACCAGAUGAUAGAGGG	2	0	transposon	Unknown	RR50960
4222	GCUGAGAAGACGGUCGAACUUG	2	no match			
4223	AUGGAGGAGUCCGUGGCAUUCAGGUU	2	0	ncRNA		7SLRNA-RA
4224	AAAACCUGAAGAUGAUCCUAAGUCGC	2	1	transposon	LINE	Helena
4225	UUAGAAGACAUAUCAUACCCAGGAA	2	0	transposon	LTR	Idefix
4226	CGCUAGCUCAUUUCCCGGGAUGUUA	2	1	transposon	LINE	Fw2
4227	AACCAUGAGAAGUAAUUAGGGC	2	no match			
4228	GUUUCAUCGUUAGACGGCUCGGGC	2	0	AT-chX		AT-chX-B
4229	UAUAGCUAAUUAACACAAUCCCGGG	2	0	ncRNA		rRNA
4230	CCAACGGGAUACCUCAUACUCGAG	2	0	transposon	LTR	micropia

4231	GGCUGAUGAAGGAAGACUCGUG	2	no match			
4232	GAUCUUCACCGUUGACUAGCCCCA	2	0	Su(Ste)		Su(Ste)
4233	AAAACCUGAAGAUGAUCCUGUGUCGC	2	1	transposon	LINE	Helena
4234	CGGGAUAAGGAUUGGCUCUAAGGGUC	2	no match			
4235	GAGAUAACCAAGGACGUUCACACGAGG	2	0	transposon	LTR	opus
4236	GCAGUUAUUACGACGAGAAGAUGGCUC	2	0	transposon	Unknown	RR50960
4237	GCCCUUUGUACACACCGCCCGUC	2	0	ncRNA		rRNA
4238	GCGGGAAUAGCUCAGUUGGUAGAGC	2	no match			
4239	GCCCGGCUAGCUCAGUCGGUAGAGCAUG	2	0	ncRNA		tRNA
4240	CUUCCUAAAAUCGAAUUAAAACCCC	2	1	no annotation		
4241	CUGGAGUACUUCAGCCAGACACUGGA	2	0	Su(Ste)		Su(Ste)
4242	UGAGAGGCGCCGUAGUUACCACCCCAC	2	0	no annotation		
4243	AUCAAUCGCUGAUUAGGCACCGGG	2	0	transposon	LTR	INE-1
4244	UCGCAUCGUCGUAGAACAAGCCCGGGG	2	1	Su(Ste)		Su(Ste)
4245	CCGUGGUACGUUCGGAGUGAAAACA	2	0	transposon	Low_complexity	F-element
4246	UAACUUCGGGAUAAGGAUUGGCU	2	0	ncRNA		rRNA
4247	GAUCGUGAUGUUCUUGGCCUUGCCCGUG	2	0	no annotation		
4248	GGUGUCCGAGCGAAUCACCUGCCUGG	2	0	coding gene		CG32512-RA|CG32512
4249	CCGGAAAUGAUCAUCCAGCUGCCAC	2	0	transposon	Low_complexity	F-element
4250	AGCGAAUGAUUAGAGGCCUUAGGGUC	2	0	ncRNA		rRNA
4251	CUCGCCGAAUCCCGGGGCCGAGGGAGCGA	2	no match			
4252	CCGAACUUGAGACCGCCCCUGGACA	2	0	Su(Ste)		Su(Ste)
4253	ACUAUGAUGAUUCACGCAGGCG	2	0	transposon	LTR	Dm88
4254	UGCUUAACUACCGUCACGUUUGGAA	2	0	transposon	LINE	INE-1
4255	GUUAAUUCCGAUAACGAACGAGACUC	2	0	ncRNA		rRNA
4256	GCAUCCUAGAAUCGACCGACCAGC	2	0	transposon	LTR	flea
4257	CAACUCAUUAAUUUUGACCCUAAGCAA	2	0	transposon	Unknown	RR50960
4258	GCCCCGUCGGACCCGCUCUGCCAGCCA	2	0	coding gene		CG31820-RA|CG31820
4259	UGUACACACCGCCCGUCGCUACUAC	2	0	ncRNA		rRNA
4260	CGCGACCGGCUCCGGGACGGCUGGGAA	2	no match			
4261	CCGGCGGGGAAGGUGGCUCGGGGGGC	2	no match			
4262	CUGUAACUCGGGUCCUGUAACUCGGGUC	2	no match			
4263	UUCCGAGGAACCAAUUGGUCCAGCUGC	2	1	Su(Ste)		Su(Ste)
4264	UCUCAUCGUCGUAGAACGAGCCCGAGG	2	1	Su(Ste)		Su(Ste)
4265	UGGAUACUAAGCAGGGACGUUAAUGAA	2	0	coding gene		pseudogene
4266	ACCUGAAGAUGAUCCUGAGUCGC	2	0	transposon	LINE	Helena
4267	UCUCGAAUGUAUGCCCGAUAUACGGA	2	0	transposon	LINE	R1-element
4268	AAAACCUGAAGAUGAUCCCUAGUCGCA	2	2	transposon	LINE	Helena
4269	ACGAGUAAUAUGAGCGCCAGAGU	2	no match			
4270	UGUCUCAUCCGGUUCAAUACAUG	2	1	transposon	LTR	diver2
4271	GUACCGAACCGCGAGACAGCAGAGC	2	1	coding gene		CG31108-RA|CG31108
4272	CCGAAUACGAGGAACAGGCAGAAUCCGA	2	0	transposon	LTR	gypsy4
4273	CUUUGGCUCAGUCACACAACAUU	2	0	transposon	LTR	micropia
4274	ACUGGCAACAUGUCGAGCUCCCAGA	2	no match			
4275	AGGAAUAAUGGAAUAGGACCGCGGUU	2	no match			
4276	GCGCAUGUUGACUGGGUAAGGAUA	2	0	transposon	LTR	opus
4277	CAGCCAGAACGGAGUUACCCUUCGCCC	2	no match			
4278	GAAUGACGGUUUAGACGGCCAGCU	2	0	transposon	LTR	ninja-Dsim-like
4279	GGGAACUCUAAUGGGACUGCCGG	2	no match			
4280	UGGACCGAACACUUUGAACCAAGU	2	0	Su(Ste)		Su(Ste)
4281	AAGAGGGACGGCCGGGGCAUUCGUAUU	2	no match			
4282	UCGGCCCGAACCCUUCUGGUGUGAUU	2	2	transposon	LINE	baggins
4283	AGUCGAGAGUGGACGGGAGCGGCGGGG	2	no match			
4284	GGGUUCGCAUCCCAUCCUCGUCGCCA	2	no match			
4285	UGAGAUGAACAAGAGUGUCACAGC	2	0	transposon	LINE	Circe
4286	GGCCCCGUUCUUUACACCCUGUUC	2	0	transposon	LINE	Doc2
4287	CGUCCGGUGAGCUCUCGCUGGCCCUU	2	no match			
4288	AAGAACAACAGUGCAGACGGCGCCC	2	0	transposon	DNA	H
4289	CAUCCUAGAAUCGACCGACCAGCU	2	0	transposon	LTR	flea
4290	AGCAGUUAUUACGACGAGAAGAUGGC	2	1	transposon	Unknown	RR50960
4291	GCCCGUAUAACGUUAAUGAUUACU	2	0	ncRNA		rRNA
4292	UGUCUGCGAGCAUGAGACCAUGGAU	2	1	transposon	LINE	INE-1
4293	UAGAGGGGGAUGGAAUUCCGCGUG	2	no match			
4294	GAGGCGCCGUAGUUACCACCCCAA	2	1	no annotation		
4295	GGAGCCGGGCGUGGAAUGCGAGUGC	2	no match			
4296	CUCGAGCAGCCACAUUCAGCCGAACGGCC	2	0	coding gene		CG18659-RA|CG18659
4297	ACCCUAAACGCGCCCGAUCUCGU	2	no match			
4298	GUUCCUCGGGAUCAAGGGCAACGAGUUC	2	0	coding gene		CG40635-RA|CG40635
4299	AGAGCUCUGAUGACGAACCCGG	2	0	transposon	LINE	G6
4300	CACGUGCCUGAGACAUGAUGGCUGUGCC	2	no match			
4301	GGAUCUUCAACUGUGAGGGGGAUCCUCAA	2	no match			
4302	UCUGAAGAUAAGUGGAGUUGGAAAG	2	0	transposon	LTR	ninja-Dsim-like
4303	CUGGAUACUAGGUAUCGACGCCGAUA	2	0	no annotation		
4304	GAAUGCCGGAGCGAGGAGGAGGA	2	0	Su(Ste)		Su(Ste)
4305	AGCAUAAGAAAAGAGGAAUACGCGAGA	2	0	transposon	LINE	S
4306	CGAAAAAAUCCCCCGGCUACGACCUU	2	0	transposon	LINE	FW
4307	ACGAGUGAUAUGAGCGCCCAGAGU	2	1	transposon	LTR	opus
4308	ACGCCCGAUACAUCAGGUCAGAGC	2	0	repeat		trf
4309	ACAGGACCAGGGUGUGGCCGCCACCA	2	1	coding gene		CG6451-RA|blue
4310	ACUGGCACGGACUUGGGGAAUCC	2	0	ncRNA		rRNA
4311	AUAUCCGAGGCCCUGUAAUUGGAA	2	0	ncRNA		rRNA
4312	ACUCGCCGAAUCCCGGGCCGAGGGA	2	no match			
4313	ACGACCUCAGAUCAGACGUGGCGACC	2	no match			

4314	UUAAUGCCGGAAGCGAUCCCACUGC	2	0	AT-chX		AT-chX-B
4315	GGUAACGGGGAAUCAGGGUUCGAU	2	0	ncRNA		rRNA
4316	GAAGGCCGAAGUGGAGAAGGGUUC	2	no match			
4317	AACGUCAUGAUCCUCGAGGCCCGU	2	0	coding gene		CG7769-RA|DDB1
4318	CAAGCAUUUGACCGAGUCUGGCUGA	2	0	transposon	LINE	Doc2
4319	ACCCUACUGAUGAAUGUUACCGCAAU	2	no match			
4320	CGACUCGUUCUUAAAUUUCGCAAC	2	1	transposon	LINE	Cr1a
4321	AGUGCCUAACCACCGAGAACCGCAAAC	2	0	coding gene		CG34373-RD|Ect4
4322	CUGGCGAAUAAAAGUAAACGAGCAA	2	0	transposon	Low_complexity	F-element
4323	GAGGUCACCUGUGUACCCUUGACGGA	2	0	transposon	LINE	F
4324	ACCAUGAAAGGCGUUGGUUGCUUA	2	0	ncRNA		rRNA
4325	AACGAUGGAAGAUGAUGAAGAUC	2	0	transposon	Low_complexity	F-element
4326	AGCGAAUGAUUAGAGGUCUUGGGGCC	2	no match			
4327	CUCGGGAUCAAGGGCAACGAGUU	2	0	coding gene		CG40635-RA|CG40635
4328	UGCUCUCGAAUGUAUGCCGAUA	2	no match			
4329	UUCGAAACAAACGAUACUGUUCGGCC	2	no match			
4330	CGGGCCACCCCUCCCACGGCGCGACCGC	2	no match			
4331	GACCAUAUCCGAAGACGCGGCG	2	0	transposon	LINE	Rt1b
4332	AGCCCCGUACGUGAAAUGACCUGUGUG	2	no match			
4333	CUGUAACUCGGGUCAACUGUAACUCGGGU	2	no match			
4334	CCAGUCAGUAGAGCUCCGAGACC	2	no match			
4335	CGGGUAACGGGGAAUCAGGGUU	2	0	ncRNA		rRNA
4336	GAACCUUUUAUGGUUAUAUUCACGGAGG	2	0	transposon	LINE	baggins
4337	GACUUCUUAGAGGGACUAACAGCG	2	no match			
4338	CUGUCCAAAAACAUGACACGCCGGGC	2	1	no annotation		
4339	CAGAGGCGACGAAGGACGUGGUAAGCUG	2	no match			
4340	AGCAGGACGGUGAUCAUGGAAGU	2	0	ncRNA		rRNA
4341	UUCAGCUGAUCGUGAUUUCCGCCGU	2	1	transposon	LTR	Dm88
4342	CGCUGAAUAUACAUAAGCCCGGUCGCGCC	2	1	transposon	LTR	GATE
4343	GCCAAUGAAAGUCACCCCUACACCCG	2	0	transposon	LTR	McClintock
4344	CGGAAUGGAACCGAGCUCUGAGU	2	0	transposon	LINE	X
4345	GGUAGGGAGUUUGACUGGGGCGGUA	2	0	ncRNA		rRNA
4346	CAAACGUAAUGUUACGGUGCCCAAAUU	2	0	ncRNA		rRNA
4347	GCGCCCGAUGCCGACGCUCAUCAGACC	2	no match			
4348	GCGGGCGGCCCGGGUUCGUUUUCCC	2	no match			
4349	CAAUGACAAAAUGCCGACGAGAUUGAGA	2	0	no annotation		
4350	AAAACCCAUGACCUAAACACGAUUCU	2	0	transposon	LTR	Idefix
4351	CGCAGAGUACGGGGCUUACUGAAGUUA	2	0	transposon	LINE	baggins
4352	UCGAGAUGACAACCUUGGACCUGA	2	0	transposon	LTR	micropia
4353	UGCGGGGAUGAACCGAACGCCGGGU	2	no match			
4354	AAGUAAAAGAACCAACCACGGGAAC	2	0	transposon	LTR	diver2
4355	UGCAUAUCAUUCACGCCAAGCUCCGACC	2	0	Su(Ste)		Su(Ste)
4356	GGAGCCCAAGGAGAUGAUUGCCGUGUU	2	0	coding gene		CG18659-RA|CG18659
4357	UUCGAGAGCAUAUAAUCCCCGUAG	2	no match			
4358	AGCUUCCUGAGAUCUGUGCCCGAU	2	0	coding gene		CG31204-RA|CG31204
4359	GAACCGAACGAUUUGUGCUCCGAACA	2	no match			
4360	AGGGCGUGAAACCGUUAAGAGGU	2	no match			
4361	AAAACCAAAGCUUUUGGGCUCCGGGGGA	2	1	ncRNA		rRNA
4362	UGCACAUUAAGUUUUGAUCGCCGAA	2	0	transposon	LTR	invader1
4363	UAUCCGAGGCCCUGUAAUUGGAAUG	2	0	ncRNA		rRNA
4364	AGCGGCGUCCGGUGAGCUCUCGCUGGCCC	2	no match			
4365	UGCUGACUAAGAGGGCUGCACAGU	2	0	transposon	LTR	HMS-Beagle
4366	GGAGUCUAACACGUGCGCGAGUC	2	no match			
4367	GCGAGCACAUGGAUUUACCGAUAAAGA	2	1	ncRNA		Uhg2-RA
4368	CUGUGAGGUCGAUCCUCAA	2	no match			
4369	AAAUAAACAAACUGACCGACACCGUGA	2	0	transposon	LTR	invader3
4370	CUGAGAAGACGGUCGAACUUGACUAUC	2	no match			
4371	UACCCGAACAUCAUCAAUCUGACCA	2	1	coding gene		CG2964-RA|CG2964
4372	GCAUUUAUGUUAGUUACUUGUUCCCCGG	2	0	ncRNA		rRNA
4373	AACCUCCCGUGGAGCAGAAGGUC	2	no match			
4374	GAUAGCAACGAAGACGCCGAUGAG	2	0	transposon	DNA	invader2
4375	ACCGAAAACAGGAUACGGACCCCGCA	2	0	transposon	LINE	TART
4376	GGCGCAAGACGGACCAGAGCGAAA	2	no match			
4377	AGGCCGAGGAGAUGCGCAAGCGCGAGGA	2	0	coding gene		CG4211-RA|nonA
4378	CAAGCAUUUACGCGAUUCACCGGA	2	0	transposon	LINE	INE-1
4379	UCCCCAUGAACGAGGAAUUCCCAGU	2	no match			
4380	UGAGAUAUGCUAGCAGACUGGAGAA	2	0	transposon	LINE	G2
4381	CCACAGAGUACGGGGCUUACUGAAGU	2	0	transposon	LINE	baggins
4382	CCACAGAGUACGGGGCUUACUAA	2	0	transposon	LINE	baggins
4383	ACCCUGAACGCGCCCGAUCUCAU	2	no match			
4384	GGAAGGACGACCACGUUGUAAUGCC	2	0	transposon	LTR	ninja-Dsim-like
4385	AAGGAUACUCGAAGACACCACGGU	2	0	transposon	LTR	ninja-Dsim-like
4386	AAGAGGGACGGCCGGGGGGCAUUCGU	2	no match			
4387	CCAACCUUAAAUUACAGCUUGACAA	2	0	transposon	LTR	McClintock
4388	CAGCUAGACUAUGUAUCCCCGGGU	2	1	transposon	LINE	Doc
4389	CAACCCUGAACGCGCCCGAUCUCGU	2	no match			
4390	GCCCGUAAAAACGACUUGGUUGACAC	2	0	transposon	LTR	gypsy
4391	ACCCUGAACGCGCCCGGUCUCGUC	2	no match			
4392	GCCGGGGCGAGGAGGAGGACUAGCUGU	2	1	Su(Ste)		Su(Ste)
4393	UGGGGUGUAGCCAAGUGGUAAGGCAGC	2	no match			
4394	AUCGUGUCCCAAUAUCUCCUGUGA	2	0	repeat		trf
4395	CCGAACUUGAGACCGCCCCUGGAUG	2	1	Su(Ste)		Su(Ste)
4396	GUCAAAUUUGGGUAUAGGGGCGA	2	no match			

4397	UGGGCCCGAACCCUUCUGGUGUGUUGA	2	1	transposon	LINE	baggins
4398	GAGAAUAGUCUCCAACGCCCAUCCC	2	0	transposon	LINE	G2
4399	AACUAAACUAUGAAGCCGGGGUAA	2	0	transposon	LINE	G2
4400	CAUUCAUAACCGAGCGAUUGUUCAUC	2	2	transposon	LTR	GATE
4401	ACAUUCAUAACCGAGCAACUGUUCAA	2	1	transposon	LTR	GATE
4402	GAACUGUGAGGGGAUCCUCAA	2	no match			
4403	CCCCUAUCCAAGCUACAGCACAGUUGGC	2	0	transposon	Low_complexity	F-element
4404	CUGCCCAGUGCUCUGAAUGUCAAAGU	2	0	ncRNA		rRNA
4405	UGGAUACUAAGCAGGGACGUUAA	2	0	coding gene		pseudogene
4406	UACCUUGGGAAUGGCUUCACAGU	2	1	ncRNA		alphagamma-element
4407	GCAAGAGAUAAAGUGCCUUAAAA	2	0	transposon	DNA	HB
4408	AUAGCGACAACCAGAGUUCCCAGUGU	2	1	transposon	LTR	DM88
4409	GCGAAGCCUUUGCUACUGACACCAG	2	0	transposon	LINE	F
4410	AACAGGCCGUACCAAUAUCCGCA	2	0	ncRNA		rRNA
4411	AUAGAAGCAGUUUGGGGGCAUUAGU	2	0	ncRNA		rRNA
4412	AUAUACGCCGCUCCGAAAUUUGUUGC	2	no match			
4413	UGCCCUCGAAUGUAUGCCCGAUAU	2	1	transposon	LINE	R1-element
4414	GCCGGAGCGAGGAGGAGGACUAUU	2	2	Su(Ste)		Su(Ste)
4415	CCAGACGUAAUACUCCACUACCCAG	2	1	transposon	LINE	HMS-Beagle2
4416	CUGGCAUUUAUACAGUGCAUUGG	2	0	transposon	LINE	baggins
4417	UGCUCUAGAAUGUAUGCCCGAUAUA	2	1	transposon	LINE	R1-element
4418	ACCGAAUUAAUCUCGAGGCUGCAAG	2	1	transposon	LTR	mdg3
4419	AACCAGACAAAAUUUUCGCACAAGUGG	2	no match			
4420	GGAUCCUCAACUGUGAGGUGAUCCUCAA	2	no match			
4421	GCAUUCGAGACCCAACGGCACGUGACAA	2	1	transposon	LINE	INE-1
4422	AUGUUUCAUCGUUAGACGGCUCGGG	2	1	AT-chX		AT-chX-B
4423	AGGAAUUUGAGCUGCGUUGACGCCGAA	2	0	transposon	LTR	ninja-Dsim-like
4424	UGCUUCAUCGUUAGACGGCUCGGCC	2	2	AT-chX		AT-chX-B
4425	AGAGUGAUCGUAUUCCACUGAC	2	no match			
4426	GAGUCAAACAUCCAAGUACCAGAGC	2	0	transposon	LTR	frogger
4427	GGAUAGUCCCGAAGAAACCUAAACCA	2	0	transposon	LTR	opus
4428	CAAGAUAUGACCAAAACAAAGAC	2	0	transposon	DNA	1360
4429	GGAGCAGAAGACGAUUAUACCAGAGC	2	0	coding gene		CG13131-RA|CG13131
4430	AGUUACUAAAGGCGGACAGAGACGGAG	2	1	transposon	LTR	McClintock
4431	ACCGAAGAAAACAAAGCCUCCGUC	2	1	transposon	Low_complexity	F-element
4432	UCCAGAACGAGGACCUGAGGGGCAU	2	0	transposon	LTR	HMS-Beagle
4433	AGACCUCUGUUCUGCUUUCAUU	2	0	ncRNA		rRNA
4434	CGUCUCAUGAGGAAUGGGCCCAGUGU	2	1	transposon	LINE	G2
4435	ACCAAGAUCAUCAACCCGUGUUAU	2	0	transposon	LINE	Cr1a
4436	CGACCAAUAAAACGAUUAAGACCACC	2	1	transposon	LTR	McClintock
4437	CUGGUUGCAACGGGUAACGGGGAAUCA	2	1	ncRNA		rRNA
4438	CUCCGGGGGAAGUAUGGUUGCAA	2	0	ncRNA		rRNA
4439	AACACGGGAAAACUUACCAGGUCCGA	2	0	ncRNA		rRNA
4440	AACUGAAUAACGAAUGGAUGAC	2	no match			
4441	CCCACCCUGAACGCGCCCGAUCUCGUC	2	no match			
4442	GAAAACGUGAAACAAGUACUCUCCGAG	2	0	transposon	LTR	blood
4443	CCCGUAGUGAUGAAGACAGAGCCAC	2	0	transposon	LINE	FW
4444	ACAUUCAUAACCGAGCGACUGUUCAC	2	1	transposon	LTR	GATE
4445	ACUCGCCGAAUCCCGGGGCCGGGGG	2	no match			
4446	GCAUUCGAGACUCAAUGGCACGUCA	2	0	transposon	LINE	Rt1b
4447	CUGGUUAAUUCCGAUAACGAACGAGAC	2	0	ncRNA		rRNA
4448	GACACUAGACAAUAACGACCGGAUAC	2	0	transposon	LTR	RR50207
4449	AGCAGCCAAAACGUGUUGGAGACGC	2	0	no annotation		
4450	UAUUGCACUUGAGACGGCCUGAU	2	0	ncRNA		mir-312-RA
4451	ACUCUAGAUAACCUCGGGCCGAUCGCA	2	no match			
4452	UUCGUCAGGACUCACCAGCGCUC	2	0	transposon	LTR	opus
4453	CACCGGCGGUACAGGAACCAGAACGAG	2	0	coding gene		CG11181-RB|cup
4454	AACGCAUUAAGUCGACCGCCUGGGGAG	2	no match			
4455	UUUGCAAAACGAGAGCGAUAGAGC	2	no match			
4456	ACACACGUUAUGGCAACCCGCUGU	2	0	transposon	LTR	Quasimodo
4457	UGUAGUGUAACGGCUAUCACAUC	2	no match			
4458	ACUAACGGGAUCAAAAGCACUGCUG	2	0	transposon	LTR	accord
4459	AAUGUGAUUUCUGCCCAGUGCUCUGAA	2	0	ncRNA		rRNA
4460	AAUCAAGAACGAAAGUUAGAGGUUC	2	0	ncRNA		rRNA
4461	ACAAAUGAAAUUCAUGAAGCUGUGGCC	2	0	transposon	DNA	RR51476
4462	GGCCCUGAGAGAUGGGCGAGCGCCGUUCC	2	no match			
4463	UGCGAGAAAAAAUGCAAUAACUGUGG	2	0	transposon	LINE	F
4464	ACGUACUAAAACAAAACCUCCGCGU	2	0	transposon	LINE	Rt1b
4465	AAGCACAAGAAAAGUCCUUGCGGGAGC	2	1	coding gene		CG12717-RA|CG12717
4466	UUCAAGGUUAUCCCAGGAUGCUGUGC	2	1	repeat		trf
4467	AAGACAAGUAUACCGUGGCACACC	2	0	transposon	LTR	micropia
4468	AGGUGGUGAACGACGGCGACAA	2	no match			
4469	GCCGCUUUAGCUCACCUGGUAGAGCAUC	2	no match			
4470	UUGAAACUGAAGCACGAAUUGUCCAA	2	0	transposon	LTR	Dm88
4471	GUCGGAGCGUGGCGUGAAUGAUAU	2	0	Su(Ste)		Su(Ste)
4472	GCAUCCUAGAAUCGACCGACCAGCUAU	2	0	transposon	LTR	flea
4473	GAGUCCAAGAGAUUUGCCCCCGAGGA	2	0	coding gene		CG18743-RA|Hsp70Ab
4474	CCCAGUUCUAAAUUAACUCGCAUGCGGC	2	0	transposon	LTR	gypsy8
4475	GAUUCCAAGAAAUAAUAUUACGGCCGA	2	0	transposon	LINE	Doc2
4476	GCCGAUGACAACGCCUUCCUAGCC	2	1	transposon	LINE	G
4477	GAUACUCGAAGACACCACGGUGAAA	2	0	transposon	LTR	ninja-Dsim-like
4478	AGAACCGCACCGAUUUCGCCCUCGC	2	no match			
4479	UGUUUCAUCGCUAGACGGCUCGGGC	2	1	AT-chX		AT-chX-B

4480	UCCAGGAAUAUCUCGACUUAAAGCACA	2	0	transposon	LTR	GATE
4481	UGUCACGCGGGAGACCGGGGUUCA	2	0	ncRNA		tRNA
4482	ACAUUCAUAACCGAGCGACUGUUCUAA	2	2	transposon	LTR	GATE
4483	UACACAAGAAGAUGACGCCAGUGC	2	0	transposon	LINE	G2
4484	GAAACAACAACGUGCGAGAUAUGA	2	0	transposon	LINE	G2
4485	GCACACAGAGGAGAGUAGAUGAGAGCA	2	no match			
4486	AUCGAUGAGAUCAGGGCCUGGAGA	2	0	transposon	LINE	G
4487	AGAUCUUCACCGUUGACUAGCCCCA	2	0	Su(Ste)		Su(Ste)
4488	AUGGCUCUCUCAGUCGCUUCCCGGGGA	2	1	ncRNA		rRNA
4489	UGGCUCAGCGUGUGCCUACCCUACG	2	no match			
4490	CCGAAAACACGAGAGCCGAGCGCUG	2	0	transposon	LTR	opus
4491	GACUGCAUAUCGAGAGGUCCCCGGUUU	2	2	ncRNA		tRNA
4492	CAGCGAGUCUAACCUUGGCCGAGAGGUC	2	no match			
4493	GGAUACUCGAAGACACCACGGUGAA	2	0	transposon	LTR	ninja-Dsim-like
4494	GCGACCUCAGAUCAGACGUGGCGAC	2	no match			
4495	UGCGAUCCACUGAAGCUUAUCCUUU	2	0	ncRNA		rRNA
4496	GCUUCCCGGGAAACCAAAGCUUUUGGGC	2	0	ncRNA		rRNA
4497	GGGCAGUUUGACUGGGGCGGUCGC	2	no match			
4498	CACAUUCUGAUUCCGAUUCGAGUUCAA	2	1	transposon	DNA	transib2
4499	CGCGAGAAGUCGAUGGCACAGUUAUCA	2	0	coding gene		CG16833-RD|CG16833
4500	CGUAGGUGAACCUGCGGAAGGAUCA	2	0	ncRNA		rRNA
4501	GUUCGAUUCCCUGAUGGCGCACCA	2	no match			
4502	AAGUGCGGGUCAUAAGCUUGCGUUGAU	2	no match			
4503	GCUUUAAUCACGAGCAAUGCGGUGAA	2	0	no annotation		
4504	UCGAACUGCACCCUGGAUCGGACUCGC	2	0	transposon	LINE	INE-1
4505	UCGCUGAGGCCCACUGGCAGGGUGUU	2	0	Su(Ste)		Su(Ste)
4506	GGCUCAUCUCAUCCUGGGGCUGUAG	2	no match			
4507	GGAAUCCUCAACUGUGAGGGGAUCCUCAA	2	no match			
4508	AGACAUGGUCGAAGAUGAGCACGUUGC	2	no match			
4509	AGGGAAAGGAGCCGAGGCGUCGA	2	0	transposon	LTR	ninja-Dsim-like
4510	CAAGCUAGACUAUGUAUCCCUGGGU	2	1	transposon	LINE	Doc
4511	ACCACCCUGAAUGCGCCCGAUCUCGUC	2	no match			
4512	AAUGCCGUUCGUCCUGGAGUCCCGAG	2	0	coding gene		CG10252-RA|CG10252
4513	UCACCAAGCGUUGGAUUGUUCACC	2	0	ncRNA		rRNA
4514	AAAACCUUCGUAGACGACCUGCU	2	no match			
4515	ACGGAAUGGAACCGAGCUCUGAG	2	0	transposon	LINE	X
4516	GAUUCCCAGACGGAGUUCAAAAUUUGCC	2	0	coding gene		CG31762-RB|aret
4517	CUCGAGGAAUACGGCCUUACCGC	2	0	transposon	LTR	ninja-Dsim-like
4518	GCCGAACUUGAGACCGUCCCUGGACAA	2	1	Su(Ste)		Su(Ste)
4519	UGCAGACGGCAGGCUUAUGGUCCACCCGA	2	0	no annotation		
4520	GCCCGUUAUAUUUGGCUCGUCAAACG	2	0	transposon	LTR	diver2
4521	ACCCUGAACGCGCCCCGAUCUCGUC	2	no match			
4522	CGACAAACGAUACCACACGAGAA	2	0	transposon	LINE	Rt1b
4523	UCACUGAUCGAGUCGUCCUGCGCGGAAG	2	no match			
4524	UCCGUGAGGUUGCGCCUGAGUCCCGA	2	0	transposon	LTR	gypsy3
4525	UCAGACGCUUAAAUCUGUUGGACGAUCG	2	0	transposon	LTR	G4
4526	CGGGAGAACUUACCAGGUCCGAAC	2	1	ncRNA		rRNA
4527	GGCCGACAUGAAGAUCACCUCUGCA	2	0	coding gene		CG5016-RA|Mst57Db
4528	GAUGUCGUAUCACCCAGGACGCCCGA	2	0	transposon	LINE	Rt1b
4529	GUAUACCGAAGAUGUGAGUGCUUAUU	2	no match			
4530	UACGACGAGAACGAGAUCAUAGU	2	no match			
4531	UGCUCUCGAAUGUAUGCCCGUAUA	2	no match			
4532	UCCGAAACAAGAAGAGGACGACGA	2	1	transposon	Low_complexity	F-element
4533	AAGCCCGAUGAACCUGAAUAUCCGUUAU	2	0	ncRNA		rRNA
4534	ACGAGUAAUAUGAGCGCCCAGAGUGAU	2	0	transposon	LTR	opus
4535	AAAGCCGAAGGCAGUGAUGGUGAC	2	0	no annotation		
4536	CUCGGGUCAAU	2	0	coding gene		CG1227-RA|CG1227
4537	ACUCGCCGAAUCCCUGGGCCGAGGG	2	no match			
4538	ACACCCAAAAAACUCAAAUGCCAUCGU	2	0	transposon	LTR	invader1
4539	ACUCACAAUCUCAGAUAAAACCCGA	2	no match			
4540	CGAGUAAUGGUCUUCGGGGCGUGCAU	2	1	coding gene		CG12717-RA|CG12717
4541	CGUUCAAGGACCACGCGAGCCACGCCAC	2	0	transposon	LINE	Rt1b
4542	CAACGAGCGCAACCCCUGUCACUA	2	no match			
4543	UUCAAGGUUACCCAGGAUGCUG	2	no match			
4544	CAGACAGAGAGGCUCAUUUACGACCAC	2	1	transposon	LTR	McClintock
4545	CAACAAACAACACAUACCCGCUGGCC	2	0	transposon	LTR	GATE
4546	UUAGUGAGGUCUCCGGACGUGAUCACU	2	0	ncRNA		rRNA
4547	UAAACCUGAAGAGAGCCCUGAAUC	2	0	transposon	LINE	Rt1b
4548	CUUGAACGAGUGUUAUUGUGGGCCGGU	2	1	ncRNA		rRNA
4549	AGAGAGAAGACCCUCAGCCUACGAGA	2	0	coding gene		CG41521-RA|CG41521
4550	GGGGACGUAACUCAGUGGAGAGCGCU	2	no match			
4551	GCAUCGGUGGUUCAGUGGUAGAAUGCUC	2	0	ncRNA		tRNA
4552	CACCGUAGAACCAGUAUCGACAUAGC	2	0	no annotation		
4553	AGGGCAUAGUGGGAACUGGGAAA	2	0	transposon	LINE	G2
4554	UGUUUCAUCGUUAGACCGCUCGGGC	2	1	AT-chX		AT-chX-B
4555	CAUGGGUCGAUGAAGAACGCAGCAA	2	0	ncRNA		rRNA
4556	CUGGCAACAUGUCGAGCUCCCAGA	2	no match			
4557	UAGUACAAUAGAUCCGACUGGGCC	2	0	transposon	LINE	baggins
4558	AAAUCUCGCGUCGGGCCGUACCCAU	2	no match			
4559	ACGAACUAGAGCGACAUCGGGAAU	2	0	transposon	LTR	roo
4560	UUGAUAGAAAAAGUACAACAGGGU	2	0	transposon	LINE	Cr1a
4561	UCGAUAUUUUCUCCUCGGACCUUGAAAA	2	0	ncRNA		rRNA
4562	AAGCCGGAUCCGUAACUUCGGGAU	2	no match			

4563	UGAGAUUGCAUGAUACACAGGCGGG	2	0	transposon	LINE	S
4564	UGAGACUUUAUCACACCUUGGCC	2	0	transposon	LTR	gypsy8
4565	ACGAGGCGGAUACGGUGACAGCGGUGG	2	0	coding gene		CG4199-RA|CG4199
4566	AACGCCCAGAUCAGCCCGCCCAAU	2	0	coding gene		CG14561-RA|CG14561
4567	UACAAUUACGUCACGAGCAACACA	2	0	transposon	LTR	DM88
4568	AUCCGCGGCUAGACGGAAAGACCCCGUG	2	no match			
4569	UAAAGGUGGAAAUGAAGAAAGUACA	2	no match			
4570	AACGAGAUUCCCACUGUCCCUAUC	2	1	ncRNA		rRNA
4571	GUACGAGAAGGAUGACAGACUUGA	2	0	coding gene		CG3691-RA|Pof
4572	CGAAUGAGACCUUGGAGGACUGAAGUG	2	0	ncRNA		rRNA
4573	AAAUAGCCCGUAUUGCUUUGGCAGUA	2	no match			
4574	CUUGAUCAAAGUAUCACGAGUUUGUUA	2	0	ncRNA		rRNA
4575	GUCCUGAAGAUCGAUCCCAGCCUAU	2	1	transposon	LTR	Max
4576	ACCCUUGAUAAACGUUUGACUUUCGAUU	2	2	transposon	LINE	BS
4577	ACCGAGCGAAAAAAUCUUUGAAUC	2	0	transposon	LTR	Stalker2
4578	UCAACUGUGAGGAGGAUCCUCAA	2	no match			
4579	CAGACACGAACGAAAUUUUGAUUA	2	0	transposon	LTR	Idefix
4580	GAGAGUGGACGGGAGCGGCGGGGGCAAA	2	no match			
4581	ACCCUGAGCGCGCCCGAUCUCGUC	2	no match			
4582	AAAACCUGAGAUGAUCCUGAGUCG	2	no match			
4583	GAAGAUAAAAACUGGAAAGCCCGGGC	2	0	transposon	LTR	Dm88
4584	AACUUAACUACGUUUUCCCGGGA	2	0	transposon	LINE	Doc3
4585	UCCCACGGAACACCAGAGCAAUCCC	2	0	transposon	LINE	G
4586	GGCCGACAUGAAGAUCACCUCUGCACU	2	0	coding gene		CG5016-RA|Mst57Db
4587	UCCUCGAUAGUAUAGUGGUUAGUAUC	2	0	ncRNA		tRNA
4588	GAGGCCCUGUAAUUGGAAUGAGUACA	2	0	ncRNA		rRNA
4589	ACCGACUACAAAGAUGCCAUACGA	2	0	transposon	Unknown	RR50960
4590	AAUCAACUGUGAGGGGAUCCUCAA	2	no match			
4591	CAUGAAUGAUGGACUGAAAUGGUCUGG	2	no match			
4592	GGAGCCGGGCGUGGAAUGCGAGUG	2	no match			
4593	UUGUUUAAAAUAAUCAAGCUGUUGCCC	2	0	transposon	LINE	Doc
4594	AUGGUGACCACGGGUGACGGGGAAUC	2	no match			
4595	UGAGACUUUAUCACACCUUGGCCUAU	2	0	transposon	LTR	gypsy8
4596	ACUCGCCGAGUCCCGGGGCCGAGGGAGC	2	no match			
4597	UAGUACAAUAGAUCCAACUGGGCC	2	1	transposon	LINE	baggins
4598	GGAGAGGCGCGAGAGGGACGGG	2	no match			
4599	GGCUUUGAGAGGCGCCGUAGUUUA	2	2	no annotation		
4600	CUUACCAGGUCCGAACAUAAGUGUGUA	2	0	ncRNA		rRNA
4601	UGAGGCGCUCGGUGUCGGUGAAGG	2	no match			
4602	AACCAAAUGAAAGACAUGAUUCGA	2	0	transposon	LTR	Quasimodo
4603	ACGCUGCGAUCCUCUCCAUGAGUU	2	0	transposon	LTR	ninja-Dsim-like
4604	CUCCCGGGGAUACGCCUGUCUGAGCGUCGCU	2	no match			
4605	AAAAUCCAUAGCAAGCUUCACACGC	2	2	transposon	Low_complexity	F-element
4606	CUGCGUAAAAUGUGGUGGCCAUCAU	2	1	transposon	LINE	G2
4607	CCGAAAGAUGGUGAACUAUGCCUGGGC	2	no match			
4608	UAUCUGAGGAAGAGUGGACCGGA	2	0	transposon	LTR	1731
4609	CCAACGAGCGGAACAUCCGGGGC	2	0	coding gene		CG34241-RA|CG34241
4610	UCAACUGUGAGGGGAUCUCAA	2	no match			
4611	AGUGGCUUUGAGAGGCGCCGUAGUUA	2	1	no annotation		
4612	GCAUCCGUUACUGUCGAGUCAAAAGAC	2	0	transposon	LTR	gypsy12
4613	AAACCUGAAGAUGAUCCUGAGUUC	2	2	transposon	LINE	Helena
4614	AUCCUCUACACCCUGUUUACAGCAGACC	2	0	transposon	LINE	G2
4615	ACCUAUUGAAGAUGAGGCCCAGAUGU	2	0	transposon	LINE	Circe
4616	AGUCGAGAGUGGACGGGAGCGGCGG	2	no match			
4617	AAAAACCUGAAGAUGAUCCCUGAGUCGC	2	no match			
4618	CGUUCAAGGACCACGCGAGCUACGC	2	0	transposon	LINE	Rt1b
4619	AUUGAAGGAAUCCCAGCACCAGCGUU	2	0	transposon	LTR	flea
4620	GGCAUUGAGAGGCGCCGUAGUUACCACC	2	1	no annotation		
4621	AUGGGAACACCGCGUGUUGUUGGCCU	2	1	ncRNA		5SrRNA
4622	UGCCACACCAAGACAUGUUGUACCC	2	0	transposon	LINE	F
4623	CCAGCCCGUGGACGGUGUGAGGCCGGU	2	no match			
4624	GAGUCCCAAGAACUGGCAGAGCUACGG	2	0	coding gene		CG6008-RA|NP15.6
4625	ACGGGUGACGGGGAAUCAGGGUUC	2	1	ncRNA		rRNA
4626	AUUUUACGUACCGCAGUUUCUCU	2	0	no annotation		
4627	GUGAGACCGUACCCGAAACCGAC	2	no match			
4628	AAACCGGAGGAAAGUGGGGAUGACGU	2	no match			
4629	ACUCUUUCGUCAUGCCGAGG	2	0	transposon	LINE	R1-element
4630	GUGGAGCCGGGCUUGAUUGGGAG	2	no match			
4631	UGUUCGGGAGACACACGGCGGGUGCUA	2	no match			
4632	AGUAUGUGAAUCCGGGGAAUCCAAA	2	0	coding gene		CG9862-RA|Rae1
4633	CUGUGAUGCCCUUAGAUGUCCUGG	2	0	ncRNA		rRNA
4634	CAGAUCAACAAAGUACGAUGGGGAACC	2	0	no annotation		
4635	ACAACCAUAAACACGAAACAGAAUUGCA	2	0	transposon	Unknown	RR50960
4636	AGCGAUAUGUGGCGAGGUUAACCCGU	2	no match			
4637	UGCAAUCGAAACGAGAAACUUCAUC	2	0	coding gene		CG41521-RA|CG41521
4638	UAUCCAUCAAUCCCAGCCCAAGUA	2	0	transposon	LTR	297
4639	UCGGCCCGAACCCUUCUGGUGUGUUU	2	1	transposon	LINE	baggins
4640	ACCCUGGACGCGCCCGAUCUCGU	2	no match			
4641	AAACCUGAAGAUCGAUCCCUGUAGUCGCC	2	no match			
4642	GAGCCCGUUUCCGUUCCCGCUCCGCC	2	0	coding gene		CG7809-RA|Grasp65
4643	ACACCACGAGAGUUGGUUGCACCUGAAGU	2	no match			
4644	GGCACAAUUACCAGAAGAUGUUGACG	2	0	transposon	LTR	roo
4645	GAGAGAUGGGCGAGCGCCGUUCCGA	2	no match			

4646	GAUGAGACGGUGAAGGUGGUUCUGGCC	2	0	coding gene		CG30170-RA|bgcn
4647	CGUAAACCAAAGGCCGUGAGUUCACCC	2	1	ncRNA		tRNA
4648	CGCCCGUCGCUACUACCGAUUGAAUUA	2	0	ncRNA		rRNA
4649	AACGCAGGUGUCCUAAGGCGAGC	2	no match			
4650	GCCGGCGGCUUUGGUGACUCUAGAU	2	no match			
4651	CAUUGAUUGAUAGUGGCUCUCAAA	2	0	transposon	DNA	HB
4652	UCCGGGCCGUAGAGUUGCACGUCUUC	2	0	transposon	LTR	ninja-Dsim-like
4653	ACGUACUAAAACAAAAACCUCCGCGUG	2	0	transposon	LINE	Rt1b
4654	GCCGUUACGAAAACAUCAUUUACG	2	0	no annotation		
4655	CGCCAAGGUGACGCAGGGCUUCUCUG	2	0	coding gene		CG2331-RA|TER94
4656	UAGACUAUGAAGGGACAUUUGCU	2	1	transposon	LINE	Cr1a
4657	UAAGGUGUUCAUCACGUUCCGCGG	2	no match			
4658	UGCUAGAAGACGAUUUUCGGACCGAA	2	0	transposon	LINE	Cr1a
4659	UGCGAGAACAGCGGUUUGCACUAU	2	0	transposon	LTR	diver2
4660	AAGCGACAUGCUCGAGAUGACAACC	2	0	transposon	LTR	micropia
4661	AAAUAGAACCUGAAACCGUAUGCGUAC	2	no match			
4662	ACAUCGAGAAUCUCACGCCCACAGG	2	0	transposon	LINE	Doc3
4663	CAGACACUUGUACCAGCACAGCA	2	1	transposon	LINE	Rt1b
4664	AGGGAGUCUAACACGUGCGCGAGU	2	no match			
4665	AGUGCGGUAACGCGACCGAUUCCGGAGA	2	no match			
4666	GCACACAGAGGAGAGCAGAGAGAGCA	2	no match			
4667	UACCAGGUCCGAACAUAAGUGUGU	2	0	ncRNA		rRNA
4668	AAACCUGAACGAUGAUCCUGAGUCGC	2	no match			
4669	GCUGCUCCGAAAACAAGGCCCGUCGU	2	1	transposon	LINE	G6
4670	UGCUCUCGAAUGUAUGCACGAUA	2	1	transposon	LINE	R1-element
4671	GUCGGGCGCGGUUAGUACUUAGAU	2	0	ncRNA		5SrRNA
4672	CAAUCUCAGAUAAAAACCCGAACGUGG	2	0	transposon	LTR	invader3
4673	UGCUCGAAUGUAUGCCCGAUAU	2	2	transposon	LINE	R1-element
4674	GACGAGAGCGAUCGGCCUGGCGUG	2	1	transposon	LTR	GATE
4675	CCAAGAGUUCCCGUGGUAGGCAGUAGU	2	0	ncRNA		7SLRNA-RA
4676	GGUUCGAGACCGUCAGCACCCACCA	2	no match			
4677	UGCAGACAUCUACUUCCCCCGCUCC	2	0	transposon	LTR	invader1
4678	UUACUUGUUCCCCGGAUAGUUUAGUUA	2	0	ncRNA		rRNA
4679	AUGAGUUUACCGUUAUGGAAGUGCUGU	2	1	coding gene		CG5183-RA|KdelR
4680	AAAUACCAAAGACAACCACGACGA	2	0	no annotation		
4681	AAAACCUGAACAUGAUCCUGAGUCG	2	1	transposon	LINE	Helena
4682	CAGGUCCGAAAAUAAGUGUGUAA	2	1	ncRNA		rRNA
4683	UGUCCGCUAGCUCAUUUCCCGGGA	2	0	transposon	LINE	Fw2
4684	ACGGUCAGACGGAAAGACCCCGUGCACCU	2	no match			
4685	AGAACUCCGAUGGCGAUAGCAACUC	2	0	no annotation		
4686	ACUCGCCGAAUCCCGGGGCCAGGGAGC	2	no match			
4687	CUGGUUAAUUCCGAUAACGAACGAGA	2	0	ncRNA		rRNA
4688	AAUUCCGAUAACGAACGAGACUC	2	0	ncRNA		rRNA
4689	CAACAACGUGGCGGCGUGAAGGGUUC	2	0	coding gene		CG6652-RA|CG6652
4690	GCAUGGCCGUUCUUAGUUCGUGGAG	2	0	ncRNA		rRNA
4691	UUUGCAAAAACGAGAGCGAUAGAGU	2	1	transposon	LTR	micropia
4692	GCAGACCCAAGAAAAAUCCCAGACC	2	0	transposon	LINE	Doc
4693	UUGACAAUAGACGAAAUAUGAUCAGAA	2	0	transposon	LINE	Cr1a
4694	ACGCCCGAUACAUCAAGUCAGAGC	2	0	coding gene		CG42398-RA|SteXh:CG42398
4695	UUUGGGUUCCGGGGGGAGUAUGGUUGC	2	2	ncRNA		rRNA
4696	UCACGCGGGAGACCGGGGUUCAAUUC	2	0	ncRNA		tRNA
4697	AGAGGCGUUUGGAAGGCGGGCGA	2	0	coding gene		CG4434-RA|CG4434
4698	GGAUACUCGAAGACACCACGGUGA	2	0	transposon	LTR	ninja-Dsim-like
4699	UGAGAUAAGUUUCGUUGCCGAGAGGGA	2	no match			
4700	GGAUCCUUAACUGUGAGGUCAA	2	no match			
4701	AGCUUCAUCCAAGCGAAUUGUGCCCGA	2	1	transposon	LINE	R1-element
4702	ACGCACCGCACGUUCGUGGGGAAC	2	no match			
4703	CAGGUCGGAGCGUGGGGAGAAUGA	2	2	Su(Ste)		Su(Ste)
4704	UUGAUUCAGCAAUUCCAGUGUUACUCGA	2	0	no annotation		
4705	CCGUGGCGGCGACGACCCAUUCGAACGU	2	no match			
4706	UACCGGGCGUGAGCAUCCACAAUGCGG	2	0	transposon	LTR	ninja-Dsim-like
4707	CUGUUUCAUCGUUAGACGGCUCGGG	2	0	AT-chX		AT-chX-B
4708	GGUGACGUGUCCGAGCGGCCGAAGGUG	2	no match			
4709	UAGUACAAUAGAUCCGACUGGGCCAU	2	1	transposon	LINE	baggins
4710	UAGACAGAAAAUAAGGGGACACGCCC	2	1	transposon	LTR	opus
4711	GCCAAUAUCACCGCGGAGACUGUCGC	2	1	transposon	LINE	Doc4
4712	GUCGGAGACUGGUAUGGUGACCCUGA	2	0	coding gene		CG9389-RA|CG9389
4713	UCCGAUGUUCUGAAACUGCGUGACC	2	0	transposon	LTR	invader3
4714	CGACUCUACAUAACUUUCCUGAUGA	2	0	transposon	LTR	3S18
4715	CACAGAGUACGGGGCUUACUAAAG	2	0	transposon	LINE	baggins
4716	UACCACAAUAUAAUCCCAGCCCACA	2	1	transposon	Low_complexity	F-element
4717	CGUCCGUUGGAUUAUGCCUGAA	2	0	ncRNA		rRNA
4718	AACGAAACUUUGAAGGCCGAAGU	2	no match			
4719	UGCGAUUCCGGCGCAGGUGACUCCGC	2	0	ncRNA		pseudogene
4720	UAUCAUCGUCGUAGAACAAGCCCGA	2	0	Su(Ste)		Su(Ste)
4721	AAGUUACCGAUGACUUGGGACGUUGA	2	0	transposon	LTR	rooA
4722	CCUACCAAGGCAAUGAUCUAUAGCU	2	no match			
4723	AGCUUGCACCAUCACCAUCACCAU	2	no match			
4724	AUGACAGUUCACGAUUGAUUCCCCACG	2	0	transposon	LTR	opus
4725	UAUGGCUCUCUCAGUCGCUUCCCGGGGA	2	1	ncRNA		rRNA
4726	GCCGUCUACAAGGCCCUGAGCGACCACC	2	0	coding gene		CG6058-RB|Ald
4727	UGCUACGUACGCUGACGAUAUUGCCCU	2	0	transposon	LINE	BS
4728	AUUCCGUAAACGAAGCCCAAAGUU	2	0	transposon	LTR	Rt1c

4729	UCCAGAAUUCGAUUUUGUUGACAU	2	1	transposon	LTR	invader4
4730	CUGCCUAAAAUACAGAUUCCCACU	2	0	transposon	LTR	roo
4731	CAACAAAGCACCAGGACAUGACAUC	2	0	transposon	LINE	X
4732	UCUCAUCGUCGUAGAACAAGCCCU	2	1	Su(Ste)		Su(Ste)
4733	GAACAAACUGAGAUCGAUGUGGAU	2	0	coding gene		CG18397-RA|ssp3
4734	CAAACCAAUAUGCUAUCUUGGCUGA	2	1	transposon	LINE	Doc
4735	AGCGUCGAACGAUUUCUGCCCACCUGA	2	0	transposon	LTR	Quasimodo
4736	AGAGACGUUAGAAAUCCCUUGGUGGA	2	0	transposon	LTR	invader6
4737	GUGUUGACACAAUGUGAUUUCUGCCCA	2	0	ncRNA		rRNA
4738	AUAAGCAAAAUGCAGAACACCGGAGAU	2	0	transposon	LINE	Doc4
4739	AAAACGUCACGAUCCCACGGCUGGA	2	no match			
4740	ACUCGCCGAAUCCCGGGGCCGAGGGU	2	no match			
4741	CCCCUGAUCAAACAUCGUCUGAUGU	2	0	transposon	LINE	Cr1a
4742	UGACUCUCGAAUGUAUGCCCGAUAUAC	2	no match			
4743	CGGUAUCUCGAAGAACAGCCAGACCCGC	2	2	coding gene		CG1873-RA|Ef1alpha100E
4744	UCCGAUAACGAACGAGACUCAAAUAUA	2	0	ncRNA		rRNA
4745	UGAGACAGGGGCACAGACCAGGGGG	2	no match			
4746	UGUACUACCCCUUCAUGACUCAAAGUA	2	1	transposon	LINE	baggins
4747	CCCGUACUCUGUGGUGACCAACA	2	0	transposon	LINE	baggins
4748	ACGAGGCAAACCGGGAGAAGUGAAAC	2	no match			
4749	AGAAGACCCUUUUGAGCUUGACUCU	2	0	repeat		LSU-rRNA_Dme
4750	ACGAGAUUAAGGCGACCCUCGGA	2	0	coding gene		CG12069-RA|CG12069
4751	AUGGCUCUUUCAGUCGCUUCCCGGGAA	2	1	ncRNA		rRNA
4752	CGUCGCCGAGAUGGCCGAGCAGGUUGAUC	2	0	coding gene		CG17927-RA|Mhc
4753	UACUUGUUCCCCGGAUAGUUUAGUU	2	0	ncRNA		rRNA
4754	GGAGCCAGCUACGAUACAUUAGCCCC	2	0	coding gene		CG7184-RA|Mkrn1
4755	ACCUUCUAGACCACAGCCAAUCCC	2	0	transposon	LINE	G2
4756	CAAAGGCAUGCCCGAUGUCCGCGAGAAA	2	2	transposon	LTR	opus
4757	CAGAUCGUUAUCCGAUUGCCCAUG	2	1	transposon	LINE	Doc4
4758	GGCUAUCAGCGGUAUUGGCUAAC	2	0	coding gene		CG6603-RA|Hsc70Cb
4759	ACACUCAUAAGGCCCACGAAGCAU	2	0	transposon	LTR	invader3
4760	GGUUCGUCCUGGGUGACCGCGUACC	2	no match			
4761	GACACCGGAACAUAUGGCUGCGAG	2	0	transposon	LTR	Quasimodo
4762	CUGGGUCGGGGUUUCGUACGUAGC	2	no match			
4763	CGUACCUGUUGGUUUGUCCCAUUAU	2	0	ncRNA		rRNA
4764	AAUGUGGAGACGUCGGCGCGACCCC	2	no match			
4765	UUCGCGAAGAAGGAGGACGAGA	2	1	transposon	LINE	G2
4766	CACUAUCGAAACGACGGAUAAGCCC	2	0	transposon	LTR	gypsy6
4767	CGCCCCAAGCACCCAUGCCACCACCCAC	2	no match			
4768	ACUGCACGAAAUUAACCCGAACGCAC	2	0	transposon	LINE	Doc3
4769	CAGAGAGAUAUGGUAGAUGGGCUA	2	0	ncRNA		rRNA
4770	CCGGGGAGGUAGUGACGAAAAAU	2	1	ncRNA		rRNA
4771	AAAACCUGAGAUGAUCCUGAGUCGC	2	no match			
4772	AAUAACCCCAAAAAUGAUCAUCGAGC	2	0	transposon	LINE	Doc3
4773	AAACCAUUCGUAGACGACCUGCUUCU	2	no match			
4774	CAGAAGAAAGAGACCGCAUUACCCAA	2	0	transposon	LTR	Idefix
4775	CGAGAGGUCCGGGUAAACCGCUG	2	0	ncRNA		rRNA
4776	UUCGAGAAUGUAUGCCCGAUAU	2	no match			
4777	GGAUCCUCAACUGUGAGGGAAUCCUCAA	2	no match			
4778	UGGGUGCGAGAGGCCCUGGGUUCAAU	2	no match			
4779	ACCACCCUGAGCGCGCCCGAUCUCGUC	2	no match			
4780	UUCGAGGUUGGCGAAGAGGAUCCCAAG	2	1	transposon	LINE	G2
4781	GCCGAAUUAAUCGCGAGGCUGCAAG	2	0	transposon	LTR	mdg3
4782	ACAAACAGAGUUCCCGAAUGAGA	2	0	transposon	LINE	Cr1a
4783	AGCUUCAUCCAAGCGAACUGUGGCCGAG	2	0	transposon	LINE	R1
4784	ACCACCCUGAACGCGCCGAUCU	2	no match			
4785	ACCCCUUGACUAAACGGUUGACUUUCGU	2	no match			
4786	GAUGGAAGACCGAAACAGUUCUGAC	2	0	ncRNA		rRNA
4787	UAAUAACCCCGAAAAUGAUCAUCGAGC	2	0	transposon	LINE	Doc3
4788	UCUGCUGCAACCGAAAGCAUAGGGAC	2	1	Su(Ste)		Su(Ste)
4789	GGCUGCAAUUGUUUCCCGUGGGACU	2	0	transposon	LINE	baggins
4790	UCUACACCAAAGUAAGCCGUGCCAG	2	0	coding gene		CG31359-RA|Hsp70Bb
4791	CGUCGCUUUUUGAUCCUUCGAUGUC	2	0	ncRNA		rRNA
4792	UAAACCGUCAUUCCAGCGCAAGUUGCCCCC	2	1	transposon	LTR	ninja-Dsim-like
4793	GACACAACAACAUCCCGGGAAAUGAGC	2	0	transposon	LINE	Fw2
4794	AGAGCAUAAAGACGACAAAAGUGACA	2	1	transposon	LINE	Doc3
4795	CCGUUAAACAUUUGCGAGUAUUCGGA	2	0	transposon	LTR	frogger
4796	AGCUGCGAGAAUUAAUGUGAAUUGC	2	no match			
4797	GAAACCAAAAUACAGAUCAACAAAG	2	0	no annotation		
4798	GAAGAGAUAAGUCGUGCCACCCAG	2	0	transposon	LINE	I
4799	CCUUUACUGAGACAUGGUGACAUCGG	2	0	no annotation		
4800	GGAUCCUCAACUGUGAGGGGAUCCUCGA	2	no match			
4801	UCUGACGAGAACUCGAGCCACGUAGCC	2	1	no annotation		
4802	AACAGGCUGAUAACCCCCAAGAGCUCA	2	no match			
4803	AGCAAGCUAACCCUCAGCUCCCAC	2	0	transposon	LINE	G2
4804	ACUGAAUAAAACGAAAUGGAUGA	2	0	transposon	LTR	invader4
4805	CACAACAACAUCCCGGGAAAUGAGCU	2	0	transposon	LINE	Fw2
4806	GCGACCUCAGAUCAGACGUGGCGA	2	no match			
4807	UCCGAAGAUAACGAGGAGCAGAGCUGG	2	1	transposon	LTR	297
4808	CUCGAACUUAGUUCAGCCCUACGAA	2	0	transposon	LINE	I
4809	UCUCCCCGCGGAAAGACGGAAAGACCCCA	2	no match			
4810	CCCAGAAACACAGUGUUUGACGCCAAG	2	0	coding gene		CG18743-RA|Hsp70Ab
4811	GCAGCAAAUAAUUAAUAGUACACCGCC	2	0	transposon	LTR	diver2

4812	CUUCCAGAGAGCGUUUACCGAAA	2	0	transposon	LTR	Idefix
4813	UACCAUGGUUGCAACGGGUAACGGGGA	2	0	ncRNA		rRNA
4814	CGAGAUAGUACGAAUCGAUUACGAA	2	0	transposon	LINE	I
4815	AACGGAGGUGUCCCAAGGCCAGCUCAGU	2	0	ncRNA		rRNA
4816	AACCCAGGUGUCCUAAGGCGAGCUC	2	no match			
4817	CCAACUAAGGUCCCAAAGCGCGUGCUA	2	no match			
4818	CCCAGAAACACAGUGUUUGACGCCAAU	2	1	coding gene		CG18743-RA|Hsp70Ab
4819	ACUUGAGACCGCCCCUGGACGACCCAC	2	0	Su(Ste)		Su(Ste)
4820	CUGAAUAGAGAGUUAAACAGUACGU	2	0	ncRNA		rRNA
4821	CAUAUCCGAGGCCCUGUAAUUGGAAUGA	2	0	ncRNA		rRNA
4822	CCCUCAAGUAUCGCAGAACUCGCU	2	0	transposon	LTR	DM88
4823	CUGUAUCACUGCUGGGGGUAGAGCACU	2	no match			
4824	GGAAAAUAUGAAACUAUAGACGAAC	2	0	no annotation		
4825	CACCGAAAUUGAUACGGAUGGAUCCGGC	2	1	coding gene		CG5024-RA|CG5024
4826	AUACCACCCUGAACGCGCCCGAUCUC	2	no match			
4827	GCACACAGAGGAGGGCAGAUGAGAGCA	2	no match			
4828	GGCGGGGCGCGGGACAUGUGGCGUA	2	no match			
4829	GCCGGGCGACCUGGGUGUUCUGAUGGAA	2	0	coding gene		CG13577-RB|CG13577
4830	UAUUCGCCAGUCAGAGAGCCACU	2	0	transposon	Low_complexity	F-element
4831	GGAUCUCAACUGUGAGGGGUCCUCAA	2	no match			
4832	CAAUCUCAGAUAAAAACCCGAACGUA	2	0	transposon	LTR	gypsy4
4833	AACAUGUCGAGCUCCCAGAACAAC	2	no match			
4834	AUUGCAUUCACCCCAUUUGACAGA	2	0	transposon	LINE	G2
4835	GCGUCAACGAGGCAUUAACCAUAA	2	1	transposon	LTR	Dm88
4836	AAAACCGCUGAUCGACUCCCAGGAGGA	2	0	coding gene		CG31973-RA|CG31973
4837	AACCAGAGAUCUAGAGGGCCCAA	2	0	no annotation		
4838	AAUUCCGUAAACGAAGCCCAAAGUUCGA	2	0	transposon	LTR	Rt1c
4839	ACGUACUAAAGCAAAAACCUCCGCGU	2	1	transposon	LINE	Rt1b
4840	AAGACAAUCUGGGAGGUGCCCGAGC	2	no match			
4841	UCUCAUCGUCGUAGAAGAAGCCCG	2	1	Su(Ste)		Su(Ste)
4842	UUCUAUAUGGAUGACCUAAUGACUGGA	2	0	transposon	DNA	HB
4843	UGAGAAAGUACUUUGAUCCAGCAA	2	0	transposon	LTR	297
4844	UGAUCCUCAACUGUCAGGGGAUCCUCAA	2	no match			
4845	GACACCAGGAAGAUCGAUGAGGCC	2	0	transposon	LINE	Rt1b
4846	CCCGAAAGUAGAUGAGGUAGCGUACC	2	1	transposon	LINE	FW
4847	GGCUUUGAGAGGCGCCGUAUUACCAC	2	no match			
4848	GCAACUAGCGACGCUGCUAUACCCAA	2	1	coding gene		CG12717-RA|CG12717
4849	CAAAACCGAUAUGCUAUCUUGGCUGA	2	2	transposon	LINE	Doc
4850	UGACACAGUAGAACUAAUCCUGAAGGC	2	1	transposon	LTR	gypsy
4851	GGCCGUGAUCGUCUAGUGGUUAGGACCCC	2	1	ncRNA		tRNA
4852	ACAACCAGAGAUCUAGAGGGCCCAA	2	0	no annotation		
4853	CGAAUCUAUGGGUGGUGGUGCAUGGCC	2	0	ncRNA		rRNA
4854	ACACGUGCUACAAUGGUCGGUACAAA	2	no match			
4855	GGAUCCUCAAGUGAGGGGAUCCUCAA	2	no match			
4856	UGAUGUCGGCUCAUCACAUCCUGGG	2	no match			
4857	GACUAGCUGGCCGAGUAUUUGAUAU	2	0	Su(Ste)		Su(Ste)
4858	AUUCCACGGAACACCCGAGCAAUGCC	2	0	transposon	LINE	G2
4859	AAACUUACCAGGUCCGGACAUAA	2	1	ncRNA		rRNA
4860	AAGGAGUCUAACACGUGCGCGAGAA	2	no match			
4861	CCCGAUGAGAAGAAUGGCCCGUAA	2	0	transposon	LINE	RR49783
4862	ACUCGCCGGAUCCCGGGGGCCGAGGGAGC	2	no match			
4863	GAUGUCGUAUCACCCAGGACGUCCGACC	2	0	transposon	LINE	INE-1
4864	CCGCCCCUGGACGACCCACGAUUUAACA	2	no match			
4865	CUUCAUCCAAGCGAACUGUGCCCGA	2	0	transposon	LINE	R1-element
4866	CGUACACAUAUUUCGCAACGUGACUGU	2	0	repeat		HETRP_DM
4867	CACAAAGUUAUUCUUCUACCGAUGGAG	2	0	transposon	LTR	Rt1c
4868	AGAUCAAGAAUCCCGUGCCAAGA	2	0	transposon	LINE	Circe
4869	AUCAUAAAAAACUCUGACGACAACGAG	2	0	transposon	LTR	RR50207
4870	CACGCAGACGACACCAGCUCAGGG	2	0	coding gene		CG3820-RA|Nup214
4871	UCCGAAAACAAGGCCCGUCGCUAA	2	0	transposon	LINE	G6
4872	CUGUAACUCGGGUCAAUCUGUACCUCGGG	2	no match			
4873	UCACCCAGGACGUUCGACCCACGA	2	0	transposon	LINE	Rt1b
4874	CCCGAAAGCAGACGGGGUAACGUACC	2	1	transposon	Low_complexity	F-element
4875	AGGAGGGAUAAUAAGGAUCCAAUC	2	no match			
4876	AACCAUCAGAGACUGCGGUGGCCAA	2	0	coding gene		CG12717-RA|CG12717
4877	GGAAAACCCAGAAAAACCCGAUCAGAA	2	1	transposon	LINE	Cr1a
4878	UAGACAUGUAGAACACGAGAAUACA	2	1	transposon	LTR	HMS-Beagle
4879	GAUAGCGUACCGGAGUGGACUGCCGUU	2	0	ncRNA		7SLRNA-RA
4880	CACCCCAUGAGAGAUUGAGAAGACG	2	0	no annotation		
4881	GAUGCCAGCUAGCAAUUGGGUGUA	2	0	ncRNA		rRNA
4882	ACCCGGCCCGGACACGGACAGGAUUG	2	no match			
4883	UCAAUUGGGACAGAGUCAUUUCGC	2	0	transposon	LINE	R1-element
4884	UGAGAUGUACAAUGCUGCCGAG	2	0	transposon	LINE	G2
4885	CAGACCCAUCAACGUCAACCAGACGGC	2	0	transposon	LINE	INE-1
4886	GUGGCGAGCGAACCGGGACCAGG	2	no match			
4887	CAUAUCCGAGGCCCUGUAAUUGGAAUG	2	0	ncRNA		rRNA
4888	AAAAAUUACAAAGACGACGGCCCACC	2	0	transposon	LTR	springer
4889	AGAUAGAAGACCCUCAGCCUACGAGA	2	1	coding gene		CG41521-RA|CG41521
4890	ACCGAAUUAAUCGCGAGGCUGCAAU	2	1	transposon	LTR	mdg3
4891	AUUGGCGAGAGACCGAUAGCGAAC	2	no match			
4892	AAAACUGUGAGGGGAUCCUCAA	2	no match			
4893	UUGGUUGAACUCUAGAUAACAUGC	2	0	ncRNA		rRNA
4894	CUCAACUCAUAUGGGACUACCCCC	2	0	ncRNA		rRNA

4895	ACUCGCCAAAUCCCGGGGCCGAGGGAGC	2	no match			
4896	UUCGAGGACAAACGCUACUGUUCGGCC	2	1	transposon	LINE	G2
4897	CAAGAUAUGACCAAAGUAAAGACACU	2	0	transposon	DNA	1360
4898	AUGAAAACGAAGGUGACCUUUUUGG	2	0	transposon	LTR	flea
4899	CACUAUAACAACGGACCAGGGAACU	2	0	transposon	LINE	Doc4
4900	GGAACUAGGUGGAUCUGCCCAGGGU	2	1	coding gene		CG8920-RA|CG8920
4901	CAAGACAUCCGAUUAGUUCAUUUUCGA	2	0	transposon	LINE	Circe
4902	ACCGCAAAAAGUUAAUUUACAUCCUG	2	0	no annotation		
4903	CCCCUGGACCUGUACUGACGCUCAUGCA	2	no match			
4904	AGGACCGCGGUUCUAUUUUGUUGGUU	2	no match			
4905	UGGGCUCCGGGGGAAGUAUGGUUGCAA	2	0	ncRNA		rRNA
4906	GAGAUUCCCACUGUCCCUACCUACUAU	2	2	ncRNA		rRNA
4907	CGCUAAUACCGGAUACGCCCUUU	2	no match			
4908	UGAGACUACAUUGGCUAGCAACGAA	2	0	transposon	LINE	I
4909	ACUGUGAGGAUCCUCAA	2	1	no annotation		
4910	CAAGAUAUGACCAAAACAAAGACACUA	2	0	transposon	DNA	1360
4911	ACGCUCCACAAGUACCUGAUCCGC	2	0	transposon	LINE	G
4912	GAUGACUAUAUCAUUCUCGGGAUUU	2	1	transposon	LTR	gypsy8
4913	GACCUAAAGACGAUUCCAAGCCCAA	2	0	transposon	LINE	G6
4914	UGUUUUCAUCGUUAGACGGCUCGGGC	2	2	no annotation		
4915	AUGACCAAUACCCAUCGCAAAAGCCCA	2	2	coding gene		CG41573-RA|CG41573
4916	GUGAGCCAAACUUAUCAGAUCCGUCGA	2	0	coding gene		CG31287-RA|CG31287
4917	AAUAAACACACAGAACAUGCAGCAGCA	2	1	transposon	LINE	Doc3
4918	AGCGUCCGACGAAUACCCUGGGGGGG	2	no match			
4919	CGAGAAGUUAAGCCAACUAAUCAAG	2	0	transposon	LINE	Cr1a
4920	AUCGAAUGGAGUGCCAGAACCCG	2	0	transposon	LINE	R1-element
4921	GGGAAGAAGACCCUGUUGAGCUUGA	2	2	repeat		LSU-rRNA_Dme
4922	AUAUCCAUCAAUCCCAGCCCAAGUA	2	0	transposon	LTR	297
4923	UGACAGUUCACGAUUGAUUCCCC	2	0	transposon	LTR	opus
4924	AUCUAGGACUACCAUGGUUGCAACG	2	0	ncRNA		rRNA
4925	ACACAGAGGAGAGCAGAUGAGAA	2	no match			
4926	GAGAGAGGGGCCCGUGCCUUGGAAAG	2	no match			
4927	UCUGAUGUUACGCCCGAAGAGAUAACU	2	0	transposon	Low_complexity	F-element
4928	AAAGAUGUCCAACAUGGCGAGUGA	2	0	transposon	LTR	opus
4929	UGCCGGGGACUCAUAGGAGACUGCCG	2	no match			
4930	UCCAAUAUCACCGCGGAGACUGUCGU	2	0	transposon	DNA	1360
4931	UCCGAUAACGAACGAGACUCUGGCAUGC	2	no match			
4932	ACCAUGGUUGCAACGGGUAACGGGGA	2	0	ncRNA		rRNA
4933	ACUCGAAUAACGAGAUAAGACGGGC	2	0	coding gene		CG3354-RA|Mst77F
4934	UCCAAGCGAAUCGAGGAUGCAAAUUC	2	0	transposon	DNA	transib2
4935	ACUCGCUGAAUCCCGGGGCCGAGGGAGC	2	no match			
4936	ACCCUUUCAUGCUUGGGAUUGUGAAC	2	2	ncRNA		rRNA
4937	CACGAGGAGCUCGGGAUUCCAUGG	2	0	transposon	LINE	G2
4938	CCCAGAAACACAGUGUUUGACGCCA	2	0	coding gene		CG18743-RA|Hsp70Ab
4939	AAGACGGAUCACGACCAGGAGCUGUCC	2	0	coding gene		CG5954-RA|l(3)mbt
4940	CUCGGCCGAUCGAAAGGGAGUCGGGUUC	2	no match			
4941	UGGAAUGUUUUCGUCCCACCCAAGU	2	0	transposon	LTR	GATE
4942	UUCGAUUGCAGUACUUUCCUAUAA	2	0	transposon	LINE	Porto1
4943	UGCACCGAAAAUAUAUGCGAGGAAA	2	1	repeat		trf
4944	UCCGCAUCGAUGACAUGGUCGUUA	2	0	transposon	LTR	GATE
4945	GCUUUAAUCACGAGCAAUGCGGUUA	2	1	no annotation		
4946	AACUGAAUAAACGAAUGAUGACAC	2	no match			
4947	UUGGGCCCGAACCCUUCUUGUGUGUU	2	1	transposon	LINE	baggins
4948	AAGUUCAAUAUCCCAAAUUCGGCGUGAC	2	1	no annotation		
4949	UGCGAGAGAUUUGGAGGCAGAAG	2	0	transposon	LTR	ninja-Dsim-like
4950	GUGUAACAACUCACCUGCCGAAGCAA	2	0	ncRNA		rRNA
4951	UGGGGAUAACAGGCUGAUACCGCCCAA	2	no match			
4952	ACAUCGAGAAUCUCACGCCCACAG	2	0	transposon	LINE	Doc3
4953	CAUGGUUACAACGGGUAACGGGGA	2	1	ncRNA		rRNA
4954	GGAUCCUCAACUGUUAGGCUCAA	2	no match			
4955	CCCGAAAAAAUCCCCCGGCUACGACCU	2	0	transposon	LINE	FW
4956	ACACGGGAAAACUUACCAGGUCCGA	2	0	ncRNA		rRNA
4957	GAUCUACUGCGUAGCGGACCAGCUC	2	0	ncRNA		7SLRNA-RA
4958	UGCAUAUCAUUCACGCCACGCUCCGACC	2	0	Su(Ste)		Su(Ste)
4959	AAAAACUGAUAGACAGAUGGACGGAGC	2	1	Su(Ste)		Su(Ste)
4960	UUCAAGGUUACCCAGGAUGCUGU	2	no match			
4961	UGCUCUCGAAUGCUUCCCCGAUAU	2	1	transposon	LINE	R1-element
4962	GCUCUCGAGAGACUACGUUCGGCAGA	2	0	transposon	LTR	invader1
4963	AAGAAAACUUAUUCGAGUAUGCAGCC	2	0	transposon	LINE	Doc3
4964	AGAAGAUCCAGUGGGACGAACCUAU	2	2	transposon	LTR	ninja-Dsim-like
4965	CUAGCGAACUAAUGUCGCCCAAGGC	2	0	coding gene		CG14995-RA|CG14995
4966	UCGGAAUGCCGGGGCGAGGAGGAG	2	0	Su(Ste)		Su(Ste)
4967	CUGCACUGCAUUCCGAAGGGCAGCAA	2	0	transposon	LTR	flea
4968	UGACCCAACAUCCUCCCCGCGUUC	2	1	no annotation		
4969	ACCCAUUACAUCCUAUACGGUCAUUU	2	0	transposon	LTR	GATE
4970	UAGAUACCCGUUACUCAGCUGGAA	2	0	transposon	LINE	INE-1
4971	UAGGAACGAAAGAGGCGAGGACAAC	2	0	coding gene		CG40271-RA|CG40271
4972	UUAAUUCCGAUAACGAACGAGACUCAA	2	0	ncRNA		rRNA
4973	UACCACAUCUAAGGAAGGCAGCA	2	0	ncRNA		rRNA
4974	UACGUAAACAACCCCAGGCUGACUCCG	2	0	transposon	LTR	Max
4975	ACCUCUAGAGCCACGGCGAAAUGUGCCA	2	no match			
4976	CGAGAGGAACCGCAGGUACGGACC	2	0	ncRNA		rRNA
4977	GGGAUAAGGAUUGGCUCUGAAG	2	0	ncRNA		rRNA

4978	UACAAUCUCCGAGCACGACCCCGAU	2	0	transposon	LINE	Circe
4979	UGCAAACGAACAAGAGACGGCCUGU	2	0	transposon	LINE	I
4980	UGAGACUACAAACGGAUUGCUGACCG	2	1	no annotation		
4981	CGCAUCACAAACGAAAUUCGUACCG	2	0	transposon	LINE	Doc2
4982	UGAGACUACAGACGGAUUGCGGACC	2	1	no annotation		
4983	AAAUACCGGCACGAGACCGAUAGUC	2	no match			
4984	AGACUCAACACGAAGACCCAGUAU	2	0	transposon	LTR	Idefix
4985	GACGUACUCGAGCGACAUGCGCCGGCC	2	no match			
4986	AGCUUCAAUAAACGUUAACCCGCUAC	2	0	transposon	LTR	opus
4987	UCCGAAGACAACGAGGAGCAGAGCA	2	1	transposon	LTR	1731
4988	UGAUCCUUCCUGAUGGCCCCGGC	2	0	transposon	LTR	gypsy12
4989	AAGCUUCAUCCAAGCGAACUGUGCCCG	2	1	transposon	LINE	R1-element
4990	UGGGUCGAUGAAGAACGCAGCAAACUG	2	0	ncRNA		rRNA
4991	UGGAGCUAACCGAUACUAAUUGUCC	2	no match			
4992	GGAUCUCAACUGUGAGGGGAUUCUCAA	2	no match			
4993	UAGCCGUGUCGAUCUAGCUAUGUCCG	2	0	transposon	LINE	INE-1
4994	CAAGCUCGAUCGUGUGACUCCGG	2	0	transposon	LINE	Doc2
4995	ACAUUGCUAUGUUGACGCGGAUUGGAC	2	0	transposon	LTR	Dm88
4996	AGCCAGUUACCGAAGCUGCGGAUU	2	no match			
4997	GACAACCAGAGAUCUAGAGGGCCCA	2	0	no annotation		
4998	UCCUGAUGACUCUUCGACAUAUA	2	0	transposon	LTR	Dm88
4999	AAUCCGGACAAACACACAAGACCCGAUC	2	0	transposon	LTR	opus
5000	AUGACUCUGUCCCGAUUCAAGAAGUC	2	0	transposon	LINE	R1-element
5001	AAGGCGAUCGAGCGGCAACAGCGACGA	2	0	coding gene		CG40084-RA|CG40084
5002	ACACACCGCCCGUCGCUACUACCGAUU	2	0	ncRNA		rRNA
5003	UGCUCAUUCCGAACAUACACCGACAAA	2	0	transposon	LINE	Doc
5004	CAAACCGAUAUGCUAUCUUGCCUGAU	2	1	transposon	LINE	Doc
5005	GCUCAUGGGUCGAUGAAGAACGCAGCAA	2	0	ncRNA		rRNA
5006	ACCAUUUACAAAAUGUAUUCCGACC	2	0	transposon	LINE	Doc
5007	ACCGUUUACACAUUUAUAUUUAUC	2	0	transposon	DNA	1360
5008	GCCUCGAAAGCCAGAUAGAGCGGC	2	0	transposon	LTR	roo
5009	AUUGCAGGUUCGUGUCCUGUCACGGUC	2	1	transposon	LINE	Doc
5010	UUUGAGAGGCGCCGUAGUUACCAGA	2	2	no annotation		
5011	AUCAAUCGCUGAUAAGGCACCGGGA	2	0	transposon	LINE	accord2
5012	UGCGCGAGGCAAAGAGACUAGCACUU	2	0	transposon	LINE	accord2
5013	CACUAUCAGACAGCAACAUGACC	2	0	transposon	LTR	McClintock
5014	GAGUCAAACAUCCAAGUACCAGAUA	2	2	transposon	LTR	frogger
5015	AAACAUGAAAAGUGUAUCCAAGGGA	2	0	transposon	LTR	Dm88
5016	UGCCAACACAUUUUGGAGACCCAC	2	0	transposon	LTR	Stalker4
5017	CUCAGCGAUGUAUGGGGCAAGUCA	2	0	coding gene		CG42398-RA|SteXh:CG42398
5018	GGAUCCUCAACUGUGAGGGGAUCUUCAA	2	no match			
5019	GAUACCGCCCUAGUUCUAACCAUAAAC	2	0	ncRNA		rRNA
5020	GCUAGAGAUUGUUUUCGUCGAGCCGA	2	0	transposon	LINE	Cr1a
5021	UUCCGAUAACGAACGAGACUCAAAUAU	2	0	ncRNA		rRNA
5022	CAACCGCACGACCCCCAGCUACGUGGCC	2	no match			
5023	AGACGACGGCGGGGGCCCUGGGUAGA	2	no match			
5024	CAGCGACCGAUAGCGAACCAGU	2	no match			
5025	CCCAGAAACACAGUGUUUGACGCCUAA	2	2	coding gene		CG18743-RA|Hsp70Ab
5026	UGACGAAUUUAGUUACAAGCAGACAU	2	0	no annotation		
5027	GCAACCGAUAACGCCCACCGAGGU	2	1	transposon	LTR	ninja-Dsim-like
5028	GCGGAAAGAGAGCGAUCUUGGCCC	2	2	transposon	DNA	1360
5029	CGAGAGCUGAAACACCAACGGAAUAA	2	1	no annotation		
5030	AAACACGGGAAAACUUACCAGGUCCGAA	2	1	ncRNA		rRNA
5031	CAGAUCAACAAAGUACGAUGGGG	2	0	no annotation		
5032	UGAAACUGAAGCACGAAUUGUCCAACA	2	0	transposon	LTR	Dm88
5033	ACCCGAAGAAGACUCUACUCGCGUACA	2	1	transposon	LINE	Circe
5034	CAUGCAGAUCGUAUGGUCUUGUACC	2	0	ncRNA		rRNA
5035	GAUAACGAACGAGACUCAAAUAUA	2	0	ncRNA		rRNA
5036	GCCAAGCGAAAGGUGCCGGCAGCGGGUG	2	0	coding gene		CG5060-RA|CG5060
5037	AACGAGAAUAUACACAGGGGCUU	2	0	transposon	LINE	Doc3
5038	AACCAAAUGAAAGACAUGAUUCGAC	2	0	transposon	LTR	Quasimodo
5039	GACCCAUUCGAACGUCUGCCCUAUC	2	no match			
5040	CAGCGAUGUGUGGGGCAAGUCAAA	2	1	Su(Ste)		Su(Ste)
5041	UCCUCAACGAGAUUCAGAUGAUCCG	2	0	coding gene		CG9206-RA|GI
5042	CGACCUGCUGCUGAUCAAGAACU	2	0	coding gene		CG30278-RA|CG30278
5043	CAAGUCAAUCGAUUGGAGAGGUG	2	0	transposon	LTR	3S18
5044	AGGGUGCAAGCGUUAAUCGGAAUU	2	no match			
5045	UACCAUGGCGAUCACGGGUAACGGG	2	no match			
5046	CAACCAGAGAUCUAGAGGGCCCAAU	2	1	no annotation		
5047	ACGCGACCUCAGAUCAGACGUGGCGAC	2	no match			
5048	UUUGACAAAAUAUGCCCUUCAGCGUA	2	0	transposon	DNA	1360
5049	GGGUAGUACAAUAGAUCCGACUG	2	0	transposon	LINE	baggins
5050	GGAUACUAAGCAGGGACGUUAAUGAA	2	0	coding gene		pseudogene
5051	ACAUUCGUAACCGAGCGACUGUUC	2	1	transposon	LTR	GATE
5052	AGGGCUGCAAUUGUUUCCCGUGGG	2	0	transposon	LINE	Fw2
5053	UUCGGGAUAAGGAUUGGCUCUGAA	2	0	ncRNA		rRNA
5054	CACCAGGAUUAUCAACCCGUGUUA	2	1	transposon	LINE	Cr1a
5055	UAUGAAUCGAGUAAAAACUGCAGG	2	0	transposon	LTR	diver2
5056	GCAUCGAGAAUAUUACACAGCCAU	2	0	transposon	LINE	Doc
5057	AAGGAAAUGACCCAGUUGGCCAAGGAG	2	0	transposon	LTR	Stalker
5058	AAGGCAAUGAUUGUUGAGUGCUUGA	2	0	transposon	DNA	1360
5059	GGAACGACCAUAAAGAAGCCGUCGAGA	2	0	ncRNA		rRNA
5060	GACCAGAGUCCCAGUUGCCAGAUCCU	2	0	coding gene		CG33092-RA|CG33092

5061	CAAGGACCAAUUGGAGGGCAAGU	2	0	ncRNA		rRNA
5062	GAGCCGUCUAACGAUGAAACAAAGAUC	2	0	no annotation		
5063	AUGCUCUCGAAUGUAUGCCCGAUAUAC	2	0	transposon	LINE	R1-element
5064	UGCAUUUCCAAAAUUGACCUCAAAGA	2	0	transposon	LINE	Circe
5065	CAGAUGCCCGUUACUCAGCUAGACCGA	2	0	transposon	LINE	INE-1
5066	UAAAAGGUGACCGGCAACGACGUU	2	0	no annotation		
5067	UCGGCUCGUCGCAUCCUGGGGCUCG	2	no match			
5068	CUAACCGUAAAAUAAUUGCAGGAGG	2	0	transposon	DNA	Ivk
5069	ACCCUGAACGCGCCCGACCUCGUC	2	no match			
5070	ACCGGACGGGCCUAGAGUUAUUCCCCC	2	0	coding gene		CG40635-RA|CG40635
5071	AAACUGAAAAGUGUAUCCAAGGGA	2	1	transposon	LTR	Dm88
5072	CGGCUAGCUCAGUCGGUAGAGCAUGA	2	0	ncRNA		tRNA
5073	GUAAGCGCGGGUCAACGGCGGGA	2	0	ncRNA		rRNA
5074	AAGAGUACAACGGAACAAGAACUA	2	0	transposon	LTR	gypsy8
5075	AAAAAUUCAAAGCGACGGCCCACC	2	no match			
5076	UGACCGGGGGGCAUUCUCCACUGCGC	2	0	transposon	LINE	G2
5077	AGAUCAGUGAAAUCAUCUUCCGGGAA	2	0	transposon	LTR	micropia
5078	CAUGUCAUAACAACUAAACCCGAAGACC	2	0	transposon	LTR	Quasimodo
5079	AGUUCCGACCAUAAACGAUGCCGACCGG	2	no match			
5080	AACCAUUCGUAGACGACCUGCUUC	2	no match			
5081	UCCCACGGAACACCCGAGCAAUGCCAU	2	1	transposon	LINE	G2
5082	GACGUCAUGACGCUGAGCGACCAC	2	0	transposon	LINE	F
5083	CGUACGCUGACGAUAUUGCCCUGU	2	0	transposon	LINE	BS
5084	AUGGCACCAUGGACGAAUAUUGCAAC	2	0	transposon	LTR	HMS-Beagle
5085	GUAGACAAAAUAAACGAAGUUGA	2	0	transposon	LTR	roo
5086	UUAUCCUUUGCUUGAUGAUUCGA	2	0	ncRNA		rRNA
5087	UGGGAACACCGCGUGUUGUUGGCCUU	2	1	ncRNA		5SrRNA
5088	GAGCGGAAAGAGAGCGAUUUUGGC	2	0	transposon	DNA	1360
5089	AUGUGACGCCAGAGGUAGACGAC	2	0	transposon	LINE	R1-element
5090	AAAACAUGGCCGCAGGAGCUGAUACCC	2	0	coding gene		CG7070-RA|PyK
5091	UGGGGGCAUUAGUAUUACGACGCGA	2	0	ncRNA		rRNA
5092	ACCACCCCUGAACGCGCCCGAUCUCGU	2	no match			
5093	CUCAUCCUGGCAACAGGAACGACCAU	2	0	ncRNA		rRNA
5094	GCUUGGACUACAUAUGGUUGAGGGUUG	2	0	ncRNA		rRNA
5095	AGAGCAGAGAACCCUCUAAUUCGGCCC	2	0	transposon	LINE	baggins
5096	GCCAGAGAACCCUCUAAUUGGGCCCGA	2	0	transposon	LINE	baggins
5097	GGAUCCUCAACUUGAGGGGAUCCUCAA	2	no match			
5098	UCGAGUGAGACCGGGGGGCAUUC	2	0	transposon	LINE	G2
5099	CGGGUAACGGGGAAUCAGGGUUCGAUU	2	0	ncRNA		rRNA
5100	CUGGCAGAGACGGCGUCGCUCGAG	2	0	transposon	LTR	Quasimodo
5101	CAAGCUAGACUAUGUAUCCCCGGGUAG	2	1	transposon	LINE	Doc
5102	UCUGAAACAAGAAGAGGACAACGA	2	1	transposon	Low_complexity	F-element
5103	UGGGCUCCGGGGGAAGUAUGGUUGC	2	0	ncRNA		rRNA
5104	GGAUCCUCAACUGUUAGGGGAUCCUCAA	2	no match			
5105	AAUACAUGCCGACGGGCGCUGACCCCU	2	no match			
5106	AAGACCGUAAACGAAAAGCUAAGAAU	2	0	transposon	LTR	Idefix
5107	AACGUGAAGUGGCAUCGCGAUUGCU	2	0	coding gene		CG31988-RA|CG31988
5108	AGAUUUACAAUUUGGACGACGGAUGC	2	0	transposon	DNA	invader2
5109	ACAUUCAUAACCGAGCGACUGUUCAUA	2	2	transposon	LTR	GATE
5110	ACCUUAACGAAACCAGGGACAUA	2	0	transposon	LTR	gypsy4
5111	CUAAGGAGGGAUGAACCGGGCCA	2	0	ncRNA		7SLRNA-RA
5112	CUCGUCCUAAACUCGGAGGUGGACUCGC	2	0	transposon	LINE	X
5113	ACCGUAGACAGGUGAUGCCGAUGCAA	2	0	transposon	LTR	springer
5114	AGCGAAAUGACUCUGUCCCGAUUCAA	2	0	transposon	LINE	R1-element
5115	ACAUAAGAAAGCCGACAUCCCGACAUGU	2	0	transposon	LTR	GATE
5116	GGAUCUCAACUGUGAGGGGAUCCCCAA	2	no match			
5117	UGCUCGAUCAAAAGGCCCUGGCAAGU	2	0	transposon	LTR	Dm88
5118	UCGGAGUAAUGCCAAUGGAGAGCGA	2	0	transposon	LINE	R1-element
5119	UGUUUCAGAGCGUUUGCCAGUUUCC	2	1	transposon	Low_complexity	F-element
5120	GCACACAGAGGAGAGCAGAUGAAA	2	no match			
5121	CGGGUAACGGGGAAUCAGGGUUCGAU	2	0	ncRNA		rRNA
5122	ACCAGAAUGAGACUAGACUUGGAGGA	2	0	transposon	LINE	RR49783
5123	UUCGGCCCGAACCCUUCUGGUGUGUU	2	0	transposon	LINE	baggins
5124	ACGACGAAUCCCGAAGCCCAGAA	2	no match			
5125	GCCAAGGAGAUGAGCACGGGCAAG	2	0	coding gene		CG18743-RA|Hsp70Ab
5126	ACGACUCGGGUAUUACCUUUUUCGU	2	1	transposon	LTR	opus
5127	ACCAGAAUGAGACUAGACUUGGAG	2	0	transposon	LINE	RR49783
5128	GCGGGACAUGUGGCGUACGGAAGAC	2	no match			
5129	AAACACUUUACCUCGAGGAACCGUUAC	2	0	transposon	LINE	accord2
5130	GAGAAACUGACAAAGGCCCUUGAA	2	0	transposon	LTR	gypsy8
5131	AUGAAACCCAGAACAAUAUAAACGGCA	2	0	transposon	LTR	HMS-Beagle
5132	ACUACAUUGAUGCUUAUCCCGAAGC	2	0	transposon	DNA	Ivk
5133	CUUAAAGGAAUUGACGGAAGGGC	2	0	ncRNA		rRNA
5134	ACUCGCCGAAUCCCGGGGCGAGGGA	2	no match			
5135	UCCUACGAGAUGGAAGCUUAUAUCGCG	2	0	no annotation		
5136	CCGAAGAUGACCAAGUUGGCUACCGAU	2	1	transposon	LTR	gypsy6
5137	AGGGAGUUUGACUGGGGCGGUAC	2	0	ncRNA		rRNA
5138	GAGUACUUCAGCCAGACACUGGA	2	0	Su(Ste)		Su(Ste)
5139	UCAACUGAGGGGAUCCUCAA	2	no match			
5140	GAGCGCACGUGUUAGGACCUGAAA	2	no match			
5141	CAAGAUAUGACCAAAGCAAAGACACA	2	1	transposon	LINE	Cr1a
5142	CAAUAUCCUAUUGAGUUGCACACUGA	2	0	transposon	LTR	micropia
5143	AUAAAUAAAAGGUGACCGGCAACGA	2	0	AT-chX		AT-chX-A

5144	AACUUGAAAAUCCAGGUGAUCCGGC	2	0	transposon	LTR	GATE
5145	ACCGUCAAGGAUGGCAAGACCCUCAC	2	0	coding gene		CG12101-RA|Hsp60
5146	CCCGAAAGCAGACGGGGUAACGUAC	1	1	transposon	Low_complexity	F-element
5147	UACUUCGUUACAACCCAUUUGGAA	1	1	no annotation		
5148	ACCGAAUUAAUCGCGAGGCUGCAAAA	1	1	transposon	LTR	mdg3
5149	AACCGAUGGGCUUGGAGUAUUUC	1	2	repeat		trf
5150	UAUGGCUCUCGUCAGUCGCUUCCCGGGGA	1	no match			
5151	UAAGGAUGAUGAUACCGACUAA	1	0	no annotation		
5152	AACUGUGAGGGAUCCUCUAA	1	no match			
5153	CCUCUCGAGCAUCGAUGCAAGCAG	1	0	transposon	LINE	INE-1
5154	CAAGACGGACGAGUAGAACACGGCAC	1	2	no annotation		
5155	UCGGAAGUGCGCGAGGUUCCUAA	1	0	transposon	LTR	GATE
5156	CAAACCAAUCCCAACCAAACCCGUCU	1	1	transposon	LINE	G2
5157	CAAUCUACAAAGAGCUGAAAUCCG	1	2	transposon	LINE	F
5158	AAGUGAGGUCUCCGGACGUGAUCACUG	1	1	ncRNA		rRNA
5159	UUCUUGAGAAUUUGGACGCCGUUUAU	1	0	transposon	LINE	Cr1a
5160	UACGAGUAGGAUAUCGAUCUGGUGG	1	0	transposon	LINE	G6
5161	CGCCCCCGACAACGCCGCCUUCGAGU	1	no match			
5162	AGCUUCAUCCAAGCGAACUGUCCCCGA	1	1	transposon	LINE	R1-element
5163	GAUUCCGGAGAGGGAGCCUUGAGAAC	1	no match			
5164	AAGCCCCGUACUCUGUGGUGACCAA	1	0	transposon	LINE	baggins
5165	GGAACUCAAAGACAAGCUAUGUAACC	1	0	transposon	LINE	G2
5166	AAAGGAAUUGACGGAAGGGCACCACCAG	1	0	ncRNA		rRNA
5167	GCCGAGAAAAAGGAACACGAAACGA	1	0	transposon	LINE	Rt1a
5168	CCAGAUGAUAUUCCGAAAACAGCCGUCA	1	0	transposon	LINE	Doc4
5169	ACCAAUCCCAACCAAACCCGUCAAA	1	0	transposon	LINE	G2
5170	CCCAGCUCGGGGAGGUAGUGGCGAAA	1	1	ncRNA		rRNA
5171	ACACGCUCUCCCGAGCACCGGUUGCCAU	1	0	coding gene		CG30362-RA|CG30362
5172	GGUUAAUUCCGAUAACGAACGAGACUCA	1	0	ncRNA		rRNA
5173	GGAACACCGCGUGUUGUUGGCCU	1	0	ncRNA		5SrRNA
5174	UUGGUCAACGAAAUGGGUUUUGGU	1	0	transposon	LTR	DM88
5175	GUGAUCGAUACGCCCCCUCAGAGUC	1	0	transposon	LTR	Max
5176	AGUAAUAUGAGCGCCCAGAGUGAUUA	1	1	transposon	LTR	opus
5177	GGAUCCUCAACUGUUAGGGAUCCUAA	1	no match			
5178	GCAAAUUCAUUCACCUUGGGAAUGGC	1	no match			
5179	UCAAGCCAAAGAGGGAGCCCGGAAU	1	2	transposon	LTR	flea
5180	CGUUCACAGACUACCUAAAAGGCGC	1	1	transposon	LINE	Doc4
5181	AGUCCAUCGCCAGAGCCAACGACAUC	1	0	coding gene		CG14200-RA|CG14200
5182	CAAGUUCUCGGCUGAUCAGUACCCCG	1	1	transposon	LTR	Quasimodo
5183	CAAACCAAUCCCAACCAAACCCGUCAAAU	1	1	transposon	LINE	G2
5184	AACCUGAAGAUGAUCCUGAGUCGCUA	1	2	transposon	LINE	Helena
5185	AAACCUAGAAGAUGAUCCUGAGUCG	1	2	transposon	LINE	Helena
5186	AGGAGGGAUAAUAAGGAUCCAAUUC	1	no match			
5187	UUCGGGGCUGAUCCGGGUUGAAGAC	1	no match			
5188	AGCAAAGCACCGAGGUAGAGCUGGC	1	0	coding gene		CG5175-RA|kuk
5189	UCCGCCCGUGGAGAAGUACAUCCCCCG	1	0	coding gene		CG13591-RA|Ssl
5190	AAGACGAGAAGACCCUAUGGUGCUUU	1	no match			
5191	UUCCGAGGAACCAUUGAUCCAGCUG	1	no match			
5192	CUGUAACUCGGGUCUGUAACUCGGGUC	1	no match			
5193	AUGCGCAGCGCUUUGGCCAGGAGGUGU	1	0	coding gene		CG31151-RA|wge
5194	UGUGAUGCCCUUAGAUGUCCUGGG	1	0	ncRNA		rRNA
5195	AAACCAAAGCUUUUUGGGCUCCGGGGGGA	1	no match			
5196	AUGCCGAGAGCUGAUGACGAGCUUU	1	no match			
5197	GAACAAUCAGUGGUUUCCCUAAG	1	1	transposon	LTR	DM88
5198	UGGUUAAUUCCGAUAACGAACGA	1	0	ncRNA		rRNA
5199	UGAUACUAUAAAUUUCGGCCUGCUA	1	0	transposon	LINE	Doc3
5200	ACGCAUCCUAGAAUCGACCGACCA	1	0	transposon	LTR	flea
5201	GGACGGAGCGAUGUUCGGGCCCAG	1	0	repeat		trf
5202	UUCCGACGACUGUGCUACUCCAUCC	1	0	transposon	LTR	GATE
5203	GCGAAGGUGGCGGAAUUGGACA	1	no match			
5204	UGGAUACUAAGCAGGGACGUUGAU	1	0	coding gene		CG12717-RA|CG12717
5205	AUGUUAAACAAUGUACCUGUGAUCA	1	0	transposon	LINE	Cr1a
5206	UUAUGUUAGUUACUUGUUCCCCGG	1	0	ncRNA		rRNA
5207	AGUACUGGGAUGACGUCAUAUUUU	1	0	transposon	LTR	Max
5208	CCGGGGUCCGGUGCGGAGUGCCCUUCG	1	no match			
5209	UACUGUUUGGAGGUGUACUGACCCAG	1	1	transposon	LINE	baggins
5210	UUAUAAAACUGAUUCGGAUCCUGU	1	0	transposon	LINE	Cr1a
5211	AAAACCAGAAGAUGAUCCUGAGUC	1	0	transposon	LINE	Helena
5212	GAAUCUAACAAGGCCCUGCAGGGAAC	1	0	transposon	LTR	diver2
5213	UUAUAAGGAUGAUGAUACCGACUA	1	0	no annotation		
5214	ACCUUGAGCAUAUAUGCUGUGACC	1	0	ncRNA		rRNA
5215	UACAUCGAGAAUCUCACGCCCACAGGAA	1	0	transposon	LINE	Doc3
5216	CACAUACCUGUGACUACUCCUGAGG	1	0	transposon	LINE	TART
5217	GUCAAGUGUACGAGUUUUCCCAGA	1	2	coding gene		CG6502-RA|E(z)
5218	GGCCGAUCGAAAGGGAGUCGGGUUC	1	no match			
5219	AAACUGAACGAGUUACUGCACUUCCCC	1	0	coding gene		CG3213-RA|CG3213
5220	AACAAUUCAAUUUGAGAGCCAGUAC	1	0	transposon	LINE	Cr1a
5221	CAAGCCGAGAGGUGACCCAGUUGAGA	1	no match			
5222	UACCGGACGGUAUGCAACCCUUAUAAGC	1	0	transposon	LTR	gypsy12
5223	UCAACACUGAGAACUGGGACUGCCUG	1	0	transposon	LTR	GATE
5224	AAAGCCUGAAGAUGAUCCUGAGUCG	1	1	transposon	LINE	Helena
5225	UAAUAGUCAUGCGAGGCGAACCGGGUGAA	1	no match			
5226	AAUGGAUUAACGAGAUUCCUACUGUCCCU	1	0	ncRNA		rRNA

5227	GCUGCAGAUCGAGAGGUCCCCGGU	1	1	ncRNA		tRNA
5228	UGCCCCGUACUCUGUGGUGACCGACA	1	1	transposon	LINE	baggins
5229	GCUCCGAAAACAAGGCCCGUCGCUAAA	1	1	transposon	LINE	G6
5230	UACCCUACUGAUGAUGUGUUGUUGCC	1	no match			
5231	GGUGCCAGGCCCGUAUAACGUUAAUGAUU	1	0	ncRNA		rRNA
5232	CAGUGACGGAGCCGGACGACCUCGA	1	0	transposon	LTR	Quasimodo
5233	GACAACCAGAGAUCUAGAGGGCCCAAG	1	0	no annotation		
5234	CUGAAGUUGAGUGGUGAAUCCGUGU	1	1	coding gene		CG41545-RA|CG41545
5235	UCUCAGUAAAAUUCUGUCGAGCGC	1	2	transposon	LTR	invader4
5236	CAUGGUUGCAACGGGUAACGGGGAA	1	0	ncRNA		rRNA
5237	CUCGCCGAACCCCGGGGCCGAGGGAGC	1	no match			
5238	UCCAGCCCAAUUUGGUGACGAACGCGA	1	0	transposon	LTR	HMS-Beagle
5239	UGUAAACGGUAGCUAAUUCGAGC	1	0	transposon	LINE	Cr1a
5240	ACGGAAGCAAGGCUGCAAUUUGACCUC	1	no match			
5241	UGAGACUACAAACGGAUUGCGGACCA	1	1	no annotation		
5242	GUGUUAUUGUGGGCCGGUACUAUU	1	0	ncRNA		rRNA
5243	CAAUCCGAUACAUUUCACGCCGA	1	0	transposon	LINE	INE-1
5244	GAACCCCGUUGGUGUACUGGAGGAG	1	no match			
5245	GUGGCGGCGACGACCCAUUCGAACGU	1	no match			
5246	AGAUGUACCGGGGCUGCACGCGCGCU	1	no match			
5247	UCCUGGGUAUUCUCCUGUUGUCCAG	1	1	transposon	LTR	Dm88
5248	ACCAGAAAAAUCCCCAGACCUGAUCGA	1	no match			
5249	UGACUAUGAAGAGACAUUUGCUCC	1	1	transposon	LINE	Cr1a
5250	UCAAGCGAGAUAGAGCGAUGUUGA	1	0	transposon	LTR	Max
5251	AAACAACGACCCGCUACCGGAGGAC	1	0	transposon	LTR	opus
5252	UCGAUAUUAUCCCAGCACUACGAC	1	1	no annotation		
5253	UGCGAACGAUUUGGCUAGAGCA	1	0	no annotation		
5254	CCCUUAGAUGUCCUGGGCUGCACGCG	1	0	ncRNA		rRNA
5255	UAGUUAAUUAAGAGGAUAUAUAAUA	1	0	transposon	LTR	invader2
5256	GAAGGGGGGCUCGAAGAUGCGGGGUAA	1	2	transposon	LTR	opus
5257	CUGCUUCGACAAGACGGGGACACUUACC	1	1	coding gene		CG6230-RA|CG6230
5258	UCACCAUACCCGACACUUACCCCA	1	0	transposon	LTR	opus
5259	GGAUCCUAAACUCCUCAAACUGUGAGGGUCAA	1	no match			
5260	AAGGACCAAUUGAGGGCAAGUCU	1	no match			
5261	UUCACCGAAAAUAUAUGCGAGGAGA	1	1	repeat		trf
5262	AGCAAUUCUAGAGAACGAAAGCGAU	1	1	transposon	LTR	flea
5263	CACUGAUGAAGAAGAGAUGCCCAG	1	0	transposon	LTR	ninja-Dsim-like
5264	AGUAGAGAGUGAUGCCGAGUAUUAAAC	1	2	no annotation		
5265	UCCUCAACUGUAAGGGGAUCCUCAA	1	no match			
5266	GUAGGUCGAGAUCAGCUGCCGUGUU	1	0	transposon	LINE	G2
5267	UUCCGAUAACGAACGAGACUCAAAUAUA	1	0	ncRNA		rRNA
5268	UGUGCUAGAAGACGAUUUUCGGACC	1	0	transposon	LINE	Cr1a
5269	GAACGAGUGAACUGGCAACAUUUCGAG	1	1	Su(Ste)		Su(Ste)
5270	GGCGUGUGUUGACGCGAUGUGAU	1	no match			
5271	AACGAACUAGAACAAUUGGCUCAGUGU	1	0	transposon	LTR	gypsy6
5272	AGUACCUGAAACCGUGCGCUUACAA	1	no match			
5273	AACCAGCGCGGGUAAACGGCGGAGU	1	no match			
5274	ACCAGCGUUGGAUUGUUCACCCAUG	1	2	ncRNA		rRNA
5275	GAGGAGUGGCGAAAAGAACCCCGGG	1	no match			
5276	AAAGCGAAUGAUUAGAGGUCUU	1	1	ncRNA		rRNA
5277	UCCGGUGCACAUAUACCGAGGCCAACGA	1	no match			
5278	CAAUCUCAGAUAAAAACCCGAAC	1	0	transposon	LTR	297
5279	AGAGGGACUAUCGGUUUCAAGCCGAU	1	no match			
5280	UUUCCUGAUGAAAGUAUACACGAGU	1	0	transposon	LTR	3S18
5281	CAAGAUAUGACCAAAGCAAAGACAU	1	0	transposon	DNA	1360
5282	ACCCUGAACGCGCCCAAUCUCGUC	1	no match			
5283	GAAGCCGGCGGGAGCCCCGGGGAGA	1	no match			
5284	GGUCCGAGAGGUCCCGGGUUCAAAU	1	1	ncRNA		tRNA
5285	UCGAGCGGCAUCGGGACUGUUAU	1	0	transposon	LTR	Dm88
5286	CACGCGGGAGACCGGGGUUCAAUUC	1	0	ncRNA		tRNA
5287	GACACGUUCAACCAGAUGGGCCUGC	1	1	Su(Ste)		Su(Ste)
5288	AGUACGAGAGGACCGAGAUGGAUAUAC	1	no match			
5289	CAAACCAAUCCCAACCAAACCCGUCAAC	1	1	transposon	LINE	G2
5290	CUCACCUGCACGUCGUGUGACCCU	1	0	transposon	LTR	ninja-Dsim-like
5291	UCCGGACAAACACACAAGACCCGA	1	0	transposon	LTR	opus
5292	UGAUCUUCACCGUUGACUUGCCCCG	1	1	Su(Ste)		Su(Ste)
5293	GAAAGAAAUAUAAGAGUUCAUUCUGA	1	0	no annotation		
5294	UCUCAUCGUCGUAGAACUAGCCCGUA	1	2	Su(Ste)		Su(Ste)
5295	CCAGACGUAAUACCCCACUACCCGGA	1	1	transposon	LTR	HMS-Beagle
5296	CAAGCUAAACGUGGAUUCCCGCUGGA	1	2	transposon	LINE	R1-element
5297	UCCUGUUGAUCAUUUGCCCGUGGUU	1	0	transposon	LINE	Cr1a
5298	GAUCAUAUGAAUAGGCUUACGGA	1	1	transposon	LINE	Cr1a
5299	UGACAUAUGAAACAACACCGGAAA	1	0	transposon	LTR	Idefix
5300	AUUCGAGAGAGCCUGGAGCCACCUAUA	1	0	transposon	DNA	1360
5301	CCACUCCCAGCUCGGGGAGGUAGUGA	1	0	ncRNA		rRNA
5302	ACGAGCGCACGUGUUAGGACCCGUAAGA	1	no match			
5303	AGAAUCAAGGACAAACCCGCAGUUG	1	0	transposon	LINE	Cr1a
5304	AUUGAAGGAAUCCCAGCACCAGCGUA	1	1	transposon	LTR	flea
5305	CUGUAACUCGGGUCAAUAGGAAUUCUAG	1	no match			
5306	AGCUAAUUAACACAAUCCCGGGG	1	0	ncRNA		rRNA
5307	GCUAGUUCCAAAAAAUCAUUACCGA	1	2	transposon	LTR	gypsy8
5308	GAAAGUCGGAGGUUCGAAGACGAUCAGA	1	no match			
5309	AAAUCCAUAGCAAGCUUCACACGC	1	2	transposon	Low_complexity	F-element

5310	CAACCAGAGAUCUAGAGGGCCUAA	1	1	no annotation		
5311	CGGACAAACACACAAGACCCGAUCUA	1	0	transposon	LTR	opus
5312	CGCCUGGGAAUACCGGGUGCUGUAGGCU	1	no match			
5313	CCAGCGAAUAAAUUAUUAGGGCCGU	1	0	transposon	Low_complexity	F-element
5314	UCCGGAGAGGGAGCCUGAGAAACGGC	1	0	ncRNA		rRNA
5315	ACAUUCAUAACCGAGCAACUGUUCA	1	1	transposon	LTR	GATE
5316	AACGCCAAGCACCCGAUUCUGAUACC	1	0	transposon	LTR	3S18
5317	ACAUUCAGUAACCGAGCGACUGUUCAAU	1	no match			
5318	UCUCAUCGUCGUAGGACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
5319	UGGAGAACGACGCCGGUGAGAAUG	1	0	coding gene		CG2986-RA|oho23B
5320	CACGGGAAACCUCACCCGGCCCGGA	1	no match			
5321	CGCGACCGGCUCCGGGACGGCUGGGA	1	no match			
5322	UAUCCGAGAUCCCCAGUACAUGUGCU	1	0	transposon	LINE	Doc3
5323	GUUGACUUGAUUCUACCACUGACAA	1	0	transposon	LTR	micropia
5324	CCCGAAAUCAGACGAGGUAACGUACC	1	1	transposon	Low_complexity	F-element
5325	CCCAGAAACACAGUGUUUGACGCCU	1	1	coding gene		CG18743-RA|Hsp70Ab
5326	CCGACUACAAAGAUGCCAUACGA	1	0	transposon	Unknown	RR50960
5327	UACUACAGGAAGUUCAUUCGGGACUA	1	0	transposon	LTR	RR50423
5328	CACUAUUUCACGAUAUACCUUGAGG	1	1	transposon	LINE	Cr1a
5329	CUCCCCGCGGAAAGACGGAAAGACCCC	1	no match			
5330	UACUUGUUCCCCGGAUAGUUUAGUAAC	1	1	ncRNA		rRNA
5331	CAGCGAGCUGGGCCGGCUACGAGAAG	1	0	coding gene		CG9498-RA|CG9498
5332	CCACAGAGUACGGGGCUUACUGAAG	1	0	transposon	LINE	baggins
5333	UAUUCCGAUAAACGACCAUUACACU	1	0	transposon	LTR	gypsy3
5334	CAAGACCAGAACCAAUGGACGUGG	1	0	transposon	LINE	HMS-Beagle2
5335	UGCCGAAUCCCGGGGCCGAGGGAGCGAG	1	no match			
5336	CAGAAUACAAUUCAAAGCGAACGGUC	1	0	no annotation		
5337	GAUGUCCGGACAGCUGGGCCCGAAA	1	1	Su(Ste)		Su(Ste)
5338	GAUCCUGAAGUCCAUGCCCGAACA	1	no match			
5339	AACAGCAUUUGCGAGGCGGGAUGUGAG	1	0	no annotation		
5340	CUGGAUGCAAAGUAUCGACGCCGA	1	0	no annotation		
5341	ACCGAAUUAACCGCGAGGCUGCAAG	1	1	transposon	LTR	mdg3
5342	CAAGCUAGACUAUGUAUCCCCCGGG	1	2	transposon	LINE	Doc
5343	ACUCGCCGAAUCCCGGGCCGAGGGUA	1	no match			
5344	ACCGAAUUAAUCUCGAGGCUGCAA	1	1	transposon	LTR	mdg3
5345	GAGGUAACGAUGGAGUUUGACGGU	1	1	transposon	LTR	flea
5346	CAGUCCAUAGAACUGUCGAACACCAU	1	2	no annotation		
5347	UCCGGAGAGGGAGCCUGAGA	1	0	ncRNA		rRNA
5348	ACGGAAGACCCGCUCCCCGGCGCU	1	no match			
5349	UAUGACGAUGGCGUACAGGAGGCU	1	no match			
5350	GCGAAAGAUUCAGGAUACCUUCGGGAC	1	0	ncRNA		rRNA
5351	UCGGCCCGAACCCUUCUGGUGUU	1	1	transposon	LINE	baggins
5352	UGGAACGAAAUUGGCCUGAUUAGCGGU	1	1	transposon	LTR	GATE
5353	ACCACCCUGAACGCGCCCGAUCUCUC	1	no match			
5354	UGGGCUCCGGGGGAAUAUGGUCGCA	1	no match			
5355	UCGAGUAGAAUGGUGAACGAGAG	1	0	repeat		trf
5356	AACUGAAUAACGAAAUGGAUGACA	1	no match			
5357	UACUCGAUAGUAUAGUGGUUAGUAUCCC	1	1	ncRNA		tRNA
5358	CAUCUAAUUAGUGACGCGCAUGAAUG	1	0	ncRNA		rRNA
5359	UGGCAGCGAACCGGGACUAGGCCAG	1	no match			
5360	UAGACCACACACUUCUUCCAGGACAGUC	1	no match			
5361	ACAUUCAUAACCGAGCGACUGUUCAAC	1	1	transposon	LTR	GATE
5362	UCCAAUAUCACCUCGGAGACUGUCGC	1	1	transposon	LINE	Doc4
5363	UUCGCAAAAAUCACGGAACGAUCGAA	1	0	transposon	LINE	Doc2
5364	UCCGAUGAUAUCACUUGGCCAACAUU	1	1	no annotation		
5365	UGAUUCAUCGUUAGACGGCUCGGGCG	1	1	AT-chX		AT-chX-B
5366	GCACGCAUCGUGACGAUCUUGAACC	1	0	coding gene		CG2247-RA|CG2247
5367	UAGUGACGCGCAUGAAUGGAUGAAC	1	1	ncRNA		rRNA
5368	UCGAGUGUGACCGGGGGGCAUUCUC	1	0	transposon	LINE	G2
5369	UAUCCUGGCUGACAUGAAUCGAUC	1	no match			
5370	AUUCGUAUUGCGCCGCUAGAGG	1	no match			
5371	AAGAACAACAGUGAAGACGGCGCCCGC	1	0	transposon	LTR	diver2
5372	UGAGAUACGCUAGCAGACUGGAGA	1	1	transposon	LINE	G2
5373	CUCAACUUGAGGGGAUCCUCAA	1	no match			
5374	GCGUGAGAUGAUCAGUGCGUCGGAG	1	2	coding gene		CG5798-RA|CG5798
5375	ACCACCCCUGAACGCGCCCCGAUCU	1	no match			
5376	GACCGGACUGGCCCGUGGUGAUGCU	1	0	coding gene		CG2979-RA|Yp2
5377	AAGCCGUCGAGAGAUAUCGGAAGAGUU	1	0	ncRNA		rRNA
5378	ACAGCAACAAUUGUUAUUUUUCGC	1	1	coding gene		CG6627-RA|Dnz1
5379	UGGGUGGUGCAUCAUUUGCGAGGACC	1	0	coding gene		CG17736-RA|schuy
5380	CUACAACACAUCGAGCGACCGCGUA	1	0	transposon	LINE	Doc3
5381	CUACUUGCCGUGUCAGUGGAUGACGC	1	0	transposon	LTR	ninja-Dsim-like
5382	GGCUCUCUCAGUCGCUUCCCGGGUA	1	1	ncRNA		rRNA
5383	GUGAGGUCUCCGGACGGCUCGGG	1	no match			
5384	UCAACCUGAGAGCUUGAUCCUGGCUCA	1	no match			
5385	GCAUGCCCGAUGUCCGCGAGAUUGCCGC	1	0	transposon	LTR	opus
5386	CACUACAUGAAACCGGACGUGAUGCC	1	0	coding gene		CG5548-RA|CG5548
5387	UGUCAUCGUCAACUGCAUCCCGUGAUCC	1	no match			
5388	AAAUGCCGGAUGCAGAAAUCCCGAAUGA	1	0	transposon	LTR	micropia
5389	CGUCACCAUACCCGACACUUACCC	1	0	transposon	LTR	opus
5390	ACCCUCUGAUGAUGUAUUGUUGUCC	1	no match			
5391	GGUUGCAACGGGUAACGGGGAAU	1	0	ncRNA		rRNA
5392	CUCAACUGUGAGGGGGAUCCUCAA	1	no match			

5393	AAACCUGAGAUGAUCCUGAGUCGC	1	no match			
5394	UCGAGAUGGAGAAGCUUCAGAGUGA	1	0	transposon	LTR	ninja-Dsim-like
5395	CAUGGAAUAAAAUAUGGACGAGAAAGAA	1	0	transposon	LTR	diver2
5396	AAAGGAGCAAUUUGGGCCGUCAU	1	no match			
5397	GUUGACUUGAUUCUACCACUGACA	1	0	transposon	LTR	micropia
5398	UUCGAGAACAAACGUUACUGUUUGGCC	1	2	transposon	LINE	G2
5399	UACACACUCAUCAACCAAUGGACGUA	1	1	no annotation		
5400	ACUUGCCCAACAGCGUGCCGAGCUUGA	1	no match			
5401	GUUCGAGCGAACAGUGGUACGAC	1	1	ncRNA		rRNA
5402	UUUGCAAAACGAGAGCGAUACGAGC	1	no match			
5403	AAAAAACCUGAAGAUGAUCCUGAGU	1	2	transposon	LINE	Helena
5404	UGUGAUUUCUGCCCAGUGCUCU	1	0	ncRNA		rRNA
5405	AUCUCAGUCGGAUUAGCCCAUAA	1	0	coding gene		CG41521-RA|CG41521
5406	UGACAAUGGUUCCCAAUUUCGAGU	1	0	transposon	LTR	G4
5407	UGCUGGAGUGUUGUGGGGUCGCAGAG	1	no match			
5408	GAGAGAUUAAAGACUAAGCCUCAGA	1	0	transposon	LINE	Cr1a
5409	CCAGAGACAUGAUUUCACGAAUUUUU	1	0	no annotation		
5410	UAUUACAAUACCCAGAUUUCAGCAAA	1	0	transposon	LTR	gypsy8
5411	AAAGCACCGUUUCUCGUCCGAUCAA	1	no match			
5412	CCCAGACGUAAUACUCCACUACCCGGG	1	2	transposon	LTR	HMS-Beagle
5413	GGAGACGUGGCCGAGUGGCUGAAGGC	1	no match			
5414	GCUUCAUCCAAGCGAACUGUGCCCGU	1	1	transposon	LINE	R1-element
5415	UUUGAAGGCCGAAGUGGAGAAGGGUUCC	1	no match			
5416	AGGGCGGAGGCCGGCGGCGGGUGGGU	1	no match			
5417	AAUCAACCGAGAUUCCCUUAGUAGUGGC	1	no match			
5418	UCUCAUCGUCUAGAACAAGCCCGAGG	1	no match			
5419	ACCACCCUGAACGCGCCUGUCUGAGCGU	1	no match			
5420	UAACUUAAGCGAAAGCAUUUGCCA	1	0	ncRNA		rRNA
5421	AAUCUUGAGAAAUGACAAUGGUC	1	0	transposon	LTR	Dm88
5422	ACCCCUUGAUAAACGGUUGACUUUCG	1	2	transposon	LINE	Rt1b
5423	ACCCUGAACGCGCCGAUCGUCGU	1	no match			
5424	UCUUCUACCACAACACGCACACAU	1	0	transposon	DNA	1360
5425	UUCAGUAAAAUUCUGUCGAGCGCUGC	1	0	transposon	LTR	invader4
5426	UACAUCGAGAAUCUCACGCCCACAGGA	1	0	transposon	LINE	Doc3
5427	CGGAGAAGCCGGCGGGAGCCCCGGA	1	no match			
5428	AAGAACGAAAGUUAGAGGUUCGAA	1	0	ncRNA		rRNA
5429	CUGACAAUGAUAGGGCAGUUUACAGC	1	1	no annotation		
5430	AACGAGAAACUUCAGUUCAGUCAGA	1	0	coding gene		CG12717-RA|CG12717
5431	AUACGCAAAGUUAUACCCGUACCCG	1	0	transposon	LTR	springer
5432	UUCCGGAGAGGGAGCCUGAGAAACGGC	1	0	ncRNA		rRNA
5433	CCCCGCGGAAUGACGGAAAGACCCCA	1	no match			
5434	GAUGUCGUAUCACCCAGGACGUUCGUACC	1	2	transposon	LINE	Rt1b
5435	AGCGAGGAGGAGGACUAGCUGGCCGA	1	0	Su(Ste)		Su(Ste)
5436	AAGCAUCGCGAAGGCCCGCGGCGGGU	1	no match			
5437	UGCUCGAUCAAAGGCCCUGGCA	1	no match			
5438	ACCCCCUGAACGCGCCCGAUCUCGU	1	no match			
5439	UCUGCAUGAAGAUGAUUCCGAAGA	1	0	transposon	LINE	Circe
5440	UGCUCACAAAGGUGGAUCUGGCCGAGG	1	0	coding gene		CG8479-RA|opa1-like
5441	GAGAGCGUUUACCGAAAUUAUAAUCCC	1	1	transposon	LTR	Idefix
5442	AGCCGGACGGUGGCCAUGGAAGUCGGUA	1	no match			
5443	ACCCUUGAUAAACGGUUGACUUUCAU	1	1	transposon	LINE	BS
5444	GCAUUCGAGACCCAACGGCACGUGAAA	1	1	transposon	LINE	INE-1
5445	GUGUCCCAAGGCCAACUCAGUGC	1	0	repeat		LSU-rRNA_Dme
5446	UGGACGGUACAACGAGUUGCGAAGUCG	1	no match			
5447	GAUAACCAAGGACGUUCACACGAGU	1	1	transposon	LTR	opus
5448	GAGUCCAAGAGAUUUGCUCCCGAGG	1	0	coding gene		CG31359-RA|Hsp70Bb
5449	AUAAAUCGGAAGAGACAGCUGGUGGA	1	0	coding gene		pseudogene
5450	CUCGCAAGAAAACGAAUCCGAGUAUGC	1	1	transposon	LINE	Doc
5451	UUGAAUGAAAACCCGGCCUCUGUCAG	1	0	transposon	LTR	micropia
5452	AAUAACCUUGAGACUGAGGGAGGG	1	1	transposon	LTR	opus
5453	UAGAGGUGAAAUUCUUGGACCGGCGC	1	no match			
5454	ACAACUCACCUGCCGAAUGAAC	1	2	ncRNA		rRNA
5455	UAUCAUCUUUCAAGAUAUUUGUCCCA	1	0	transposon	LINE	Cr1a
5456	GGAUCUACAACUGUGAGGGGAUCCUCAA	1	no match			
5457	ACCGUUUACACAUUUAUAUUUUUGU	1	0	transposon	DNA	1360
5458	AUGGACAGGCGAGCGGUGCGGCAAAUG	1	0	coding gene		CG14507-RC|CG14507
5459	AGCACCGUUUCCCGUCCGAUCAA	1	no match			
5460	UUCAACCGGACGGGCCUAGAGUUUA	1	2	coding gene		CG40635-RA|CG40635
5461	CCCCUACAAACAGUACACCUACUA	1	0	transposon	LINE	baggins
5462	UGCCGGGAUCGGAGCCAGUAGCUUCUGC	1	0	coding gene		CG6303-RA|Bruce
5463	UAAUCCUUUAUAAUCCGAUCGUAUUCC	1	0	transposon	LTR	ninja-Dsim-like
5464	UACUUCGGGAUACUGUUCGGC	1	no match			
5465	AAGAUCAUCGAGAAGCUGGAGAGCG	1	0	coding gene		CG8171-RA|dup
5466	AGUUUCAUCCAAGCGAACUGUGUCCCGA	1	no match			
5467	AAAGAAGCCGUCGAGAGAUAUCGGA	1	0	ncRNA		rRNA
5468	AUCAGGUCGGAGCGUGGCGUGAUUGG	1	2	Su(Ste)		Su(Ste)
5469	GGCUUUGAGAGGCGCCGUAGUUAAC	1	1	no annotation		
5470	CACAAGACAUAACGGAACAGCUUGGAA	1	no match			
5471	AAAAACCUGAAGAUGAUCUUGAGUCGC	1	2	transposon	LINE	Helena
5472	GCGAUGAACAGAAAAGAAUCAACCGAUU	1	2	transposon	Unknown	RR50960
5473	GGAUCCUAACUGUGAGGGGAUCCUCAA	1	no match			
5474	AGGAAUUGACGGAAGGGCACCACCGGG	1	1	ncRNA		rRNA
5475	UAGAGUAUUCAUCUUGCGGCGUGU	1	0	transposon	LINE	Cr1a

5476	AAUGCCAAUCACGCCGUUUGAACCC	1	1	transposon	Simple_repeat	I
5477	UACAGAUCAACAAAGUACGAUGGGGAC	1	1	no annotation		
5478	AACGGGUAACGGGGAAUCAGGGU	1	0	ncRNA		rRNA
5479	CCGCUCAAGAUAGGCAGAUGUGCCCGA	1	0	transposon	LINE	Cr1a
5480	UGUAAUUGUCCGCGAGCUUGCCCAG	1	0	transposon	LTR	3S18
5481	GGAUACCAAGCAGGGACGUUAAUGA	1	1	coding gene		pseudogene
5482	ACCUCAGAUCAGACGUGGCGACCCG	1	no match			
5483	UAACCUUGAGACUGAGGGAGGGGCA	1	0	transposon	LTR	opus
5484	GGCGAAGGCAAACUACAAGUAAAUCCGA	1	1	transposon	LINE	Circe
5485	CGGUCUGGUGGUUUGAGCGGUGUGCC	1	no match			
5486	AUUGUUCCAAACACUUACUUGUCCG	1	0	transposon	LTR	opus
5487	UCCUGGAAUACCGAUGUGUAAAG	1	0	transposon	LINE	G6
5488	UGAUAGCCCCGUAGUGAAAUGACCUG	1	no match			
5489	AACAACCGAUAUAAUUCUAACUGAGCACC	1	0	transposon	LTR	G4
5490	UGUUUCAAUCGUUAGACGGCUCGGGC	1	no match			
5491	UAGAAUAACAAGAUGCAUACGAUGUGGC	1	0	transposon	DNA	1360
5492	UCUUAUGUAAAUGAAUCGAGAGCGA	1	0	transposon	LINE	Cr1a
5493	CAGAUCAACAAAAUACGAUGGGGAACC	1	0	no annotation		
5494	ACUCGCCGAAUCCCGGGGCCGAGGGAAA	1	no match			
5495	AUGGCUCUCUCAGUCGCUUCCGGGUA	1	2	ncRNA		rRNA
5496	ACCGUAGAGGAAACCGUGGUGCGA	1	0	coding gene		CG17291-RA|Pp2A-29B
5497	UGCACAAGUUUAGUCUCUUCCGAUU	1	2	transposon	LTR	diver2
5498	CGGAAUGGAAUCGAGCGCUGAGU	1	0	transposon	LINE	X
5499	AGGGAGUUUGACUGGGGCGGUA	1	0	ncRNA		rRNA
5500	UGCAAGAUAAAGACCCUUCUACCUGC	1	0	transposon	LINE	Helena
5501	CCGAACUUGAGACCGCCCCUGGACAAU	1	1	Su(Ste)		Su(Ste)
5502	CCCGAUAGCAGACGAGGUAACGUACC	1	1	transposon	Low_complexity	F-element
5503	CGGAAGGCCCGGGUUCAAUUCCC	1	0	ncRNA		tRNA
5504	AAACUGAAGAUGAUCCUGAGUC	1	1	transposon	LINE	Helena
5505	CGGUAAAUCACCAAUUUUGACCGC	1	1	repeat		RSP
5506	AUUCCGCUGCCCCCUCACCGUCGAGG	1	no match			
5507	ACUGAGCGCAGGGCACGUGGGCAGGGU	1	0	coding gene		CG31538-RA|CG31538
5508	AAAAACCUGAGAUGAUCCUGAGUCGC	1	no match			
5509	AAUACAUGCCGAGGGCGCUGACCCC	1	no match			
5510	CGGCUACCCACCCGACCCGUCUUGAAA	1	no match			
5511	GACAAUCGAACGAAGUCAUAGAAC	1	0	transposon	LTR	gypsy8
5512	CUCUGUUGCACCCGAUGAUGAAAU	1	0	transposon	LTR	accord
5513	UCUUGUUCCCCGGAUAGUUUAGUAC	1	no match			
5514	AGGACAAUGAUGCGACCCGUCUCU	1	0	transposon	LTR	ninja-Dsim-like
5515	GUAUACCUAUACAUUACCGCUAAAG	1	0	ncRNA		rRNA
5516	AUUGAUUACAUACUGGCUCUUUAUGC	1	0	transposon	LTR	McClintock
5517	CCCAAUGAGAUGGCCUUUGGCGAGAACA	1	1	transposon	LTR	diver2
5518	GCCCAUAACCCAGAGGUCCGAGG	1	0	ncRNA		tRNA
5519	ACCGAAACGUUACCUUGCCCUGCUG	1	no match			
5520	UUGCCAGGUAGGGAGUUUGACUGGG	1	0	ncRNA		rRNA
5521	AGAACUGAAAUGGACUACCUGCAAU	1	0	transposon	LINE	Doc2
5522	GGGAGAUAUUAGACCUCGGUUUGG	1	0	repeat		LSU-rRNA_Dme
5523	CAGACUAUGACAAAACCUUUGGCCU	1	0	no annotation		
5524	UGAGCCAGCUACGAUACAUUAGCCCC	1	1	coding gene		CG7184-RA|Mkrn1
5525	CCGAACUUGAGACCGCCCCUGGAGA	1	0	Su(Ste)		Su(Ste)
5526	AAACUCAGGACCUGUUGAACCAAGCG	1	0	transposon	LINE	X
5527	AACACGUGCGCGAGUCGGGGGCU	1	no match			
5528	GAAGAAGAGAUGCCCAGUGAGUUUUGCA	1	no match			
5529	UACUCCAGGCCCAUCUGGUUGAACGUG	1	0	Su(Ste)		Su(Ste)
5530	GAUGAUUAUAUCAUUCUCGGGAUU	1	1	transposon	LTR	gypsy8
5531	CAAAAGGGCAAAUGCUGACUUGAUCUCG	1	0	repeat		LSU-rRNA_Dme
5532	ACGAGCGCACGUGUUAGGACCCGAAAA	1	no match			
5533	ACCGGAGAUGAACGCCAACAGCAGA	1	0	transposon	LTR	McClintock
5534	AAACCUGAGAGUUUGAUCCUGGCUC	1	no match			
5535	UACAGGCGAAAUGAAAAUCCGUUUGU	1	1	transposon	LTR	micropia
5536	CGCCGAACGCUUCUCAGUGUGCCAGC	1	0	no annotation		
5537	GCGACAACGAGGUGACUCAGCAAU	1	1	no annotation		
5538	GCACACAGAGGAAACAGAUGAGAGCA	1	no match			
5539	CGCAUCUACAACAGGCCCGGAGCGUA	1	0	transposon	LTR	invader2
5540	UGAAUCCCAGCACCAGACCAAGUU	1	0	coding gene		CG13449-RA|CG13449
5541	CAAUAAUCCACGGCUGACUCCGGU	1	0	transposon	LTR	GATE
5542	GUGCCAGGCCCGUAUAACGUUAAUGA	1	0	ncRNA		rRNA
5543	CAAGAAAUACAAAACGAGCUUAAAGA	1	0	transposon	LINE	RR49783
5544	UGCGAGGAGAUUUUGGAUCGUGUCCCA	1	0	repeat		trf
5545	AGACCCCAGAAAAGGUGUUGGUUGAUA	1	no match			
5546	CACAGAGUAGGGGGCUUACUAAAGUCAG	1	2	transposon	LINE	baggins
5547	UUGAGAUAGUCGGGCUUGAUUGGGAAA	1	0	ncRNA		rRNA
5548	GUGUACUAUAACCAAUUGUCAUUA	1	0	no annotation		
5549	UAGAGGCAAUCACCGAGAGCAUCAA	1	0	transposon	LINE	F
5550	UGAAGCCGGUGAUCGAUAGUUCCUC	1	0	repeat		trf
5551	ACGCGGAAAGAGACGCAAUUUUGGCC	1	no match			
5552	GACAUCGAGGAGCCACCGACGCCGACG	1	0	coding gene		CG8978-RA|Sop2
5553	AAAUCGCCCAAGCCAAAGAAUAUC	1	0	repeat		trf
5554	AAACCUGAAGAUGAUCCUGAGUCU	1	1	transposon	LINE	Helena
5555	AAGAGUGACGGCCGGGGGCAUUCG	1	no match			
5556	AUGCCCGAUGUCCGCGAGAUUGCC	1	0	transposon	LTR	opus
5557	ACGCGCGCUACACUUACUGGCUCAGC	1	no match			
5558	CGUCCCCGUCCGCCCCCCGUUCCCC	1	no match			

5559	UCCUAAACUACGGGAGAUAUUUGG	1	0	transposon	DNA	hopper
5560	UACUCGAAGACACCACGGUGAAAGUGGG	1	0	transposon	LTR	ninja-Dsim-like
5561	GGCAUUAGUAUUACGACGCGAGA	1	0	ncRNA		rRNA
5562	UGUUUCAUCGUUAGACGGCUCGGACG	1	1	AT-chX		AT-chX-B
5563	GGACGAGGCGCCGCCGCCCCCCCCUAC	1	no match			
5564	AUCUUAUGGUAUUGGGUUUUGAUG	1	0	repeat		LSU-rRNA_Dme
5565	UCCCGUUGGUAAAUGAAGUCGAGUGC	1	0	transposon	LTR	micropia
5566	AGCAAGCCGGGCGCUCCCGUGUUC	1	no match			
5567	UAUUAUUAAGAUGGGCUUGUAGGUGUAC	1	0	transposon	LINE	baggins
5568	UAGGAAUUCAACGAGGUAUGGC	1	0	repeat		SAR_DM
5569	GCUUACAAUAUUAAAAAGUAUGAGA	1	0	transposon	LTR	diver2
5570	GGCAACAUCCAGACCAAACGAGUCCA	1	0	transposon	LINE	Rt1b
5571	UGCACUAAUAUAAGCCCUAACGUCCG	1	1	transposon	LTR	G4
5572	CGGAGAGUGUUGCCAGUACUGGA	1	0	transposon	LINE	G2
5573	UUGGUUGAACUCUAGAUAACAUGCA	1	0	ncRNA		rRNA
5574	UGCUUGGACUACAUAUGGUUGAGGGUUG	1	0	ncRNA		rRNA
5575	CACUCAGUCGCUUCAUCUUACAGACCAA	1	1	transposon	Low_complexity	F-element
5576	CAAAGAAGAACACCGAUAACAUACGA	1	0	transposon	LINE	INE-1
5577	AGAACGAAAGUUAGGGGAUCGAAGAU	1	no match			
5578	AAGGCAAUGAUUGUUGAGUGCUUC	1	0	transposon	DNA	1360
5579	CUGGACCCGUUCAAUCGCGCCCCGU	1	no match			
5580	AAGACAAGUAUACCGUGGCACACCAU	1	0	transposon	LTR	micropia
5581	UAUUAUCCCAGCACUACGAAGCGCCCU	1	0	no annotation		
5582	UGGACAAAAAGACAAAUGACACGGU	1	1	transposon	LTR	gypsy
5583	GAAGACCCUGUUGAGCUUGACUCUA	1	1	repeat		LSU-rRNA_Dme
5584	GCAUUCGAGGCCCAACGGCACGUGAA	1	1	transposon	LINE	Rt1b
5585	GCCUGAGAGACGGCACCUACUUCUAA	1	no match			
5586	GCUCACCAAGGCGACGAUCAGUAGCUGG	1	no match			
5587	GUCCGCCCGAACGCUUCGUCCCCUC	1	no match			
5588	GGAUCUCAACUGUGAGGGAUCCUCAA	1	no match			
5589	ACCCUUAACGCGCCCGAUCUCGUC	1	no match			
5590	AAUAGAAAGUAAGCCCCAUGUAAUCGAA	1	1	transposon	LTR	micropia
5591	CUGAGUUCGAUCGUGUGGCCCUGUACAC	1	0	coding gene		CG7910-RA|CG7910
5592	AACCGACGAGAGCAUAAUGGAGGAGA	1	0	coding gene		CG3002-RB|Gga
5593	UCCGAAUCAAAUGCUACGUAACCAAGC	1	0	transposon	LTR	diver2
5594	CACAGAGUACGGGGCUUACUAUUU	1	no match			
5595	CCAACUCAAAUUGGCACCGAGCGAAA	1	0	transposon	LTR	Stalker2
5596	GGAUCCUCAACUGUGAGGCUCAA	1	no match			
5597	ACGGGAAACUUACCAGGUCCGAA	1	no match			
5598	GCAUAUCGAGAGGUCCCCGGUUCAUACC	1	2	ncRNA		tRNA
5599	AACGUGUCUCUCGAAACUGUUUGA	1	0	transposon	LINE	Doc2
5600	UGAAUUGUUAGACGGAGACAUGUAU	1	1	transposon	LTR	micropia
5601	CUGCCCAAAAUACAGAUUCCCACU	1	1	transposon	LTR	roo
5602	CGCUUCGCUUGUGAGAAGUCCCGGGU	1	1	ncRNA		tRNA
5603	UGCGAGAACAGCGGUUUGCCCUGU	1	0	transposon	LTR	diver2
5604	CCCGAAAAUCGUCUUCUACCACAACACG	1	0	transposon	DNA	1360
5605	UGCUCUCGAAUGUAUGCCCGACA	1	1	transposon	LINE	R1-element
5606	UCACACGGGUAGUACAACUUACAA	1	1	ncRNA		rRNA
5607	UAGAGAGUGAUGCCGGGUAUAAAG	1	1	no annotation		
5608	CCGUUCGUUGACGUUGAUGGCUG	1	0	transposon	LTR	HMS-Beagle
5609	AUGCAAUAGAAACGCGCGAGUCGAGUUGCCC	1	no match			
5610	CCCGUACUCUGUGGUGACCAACACA	1	0	transposon	LINE	baggins
5611	UUGGGUUAAGUCCCGUAACGAGCGCA	1	no match			
5612	AUCCUCAACUGUGAGGGGACCUCAA	1	no match			
5613	GAGCGCACGUGUUAGGACCCUAAAG	1	no match			
5614	UGUUUUCAUCGUUAGACGGCUCGGGGCG	1	no match			
5615	CCGAACUUGAGACCCCAGCCGCG	1	1	repeat		trf
5616	ACCGCGAAAACCCGGGUAGCACCA	1	0	transposon	LTR	GATE
5617	GAGAAGGAGUCGACCGAGGAGAAC	1	0	coding gene		CG4609-RA|fax
5618	UCCGAGGCCCUGUAAUUGGAAUGAG	1	0	ncRNA		rRNA
5619	UGUUUCAUCGUUAGGCGGCUCGGGCG	1	1	AT-chX		AT-chX-B
5620	ACCCGCUUGACCUUACUCAACAUA	1	0	transposon	LINE	G2
5621	UUUGCAAAAACGAGAGCGAUAGAGCUGU	1	0	transposon	LTR	micropia
5622	UCCGAUUUCAAGUACAAGGCGAACAGA	1	0	repeat		RSP
5623	CUGUAACUCGGGUCAAUGGGAAUUCUAG	1	no match			
5624	GGUAACGGGGAAUCAGGGUUCGA	1	0	ncRNA		rRNA
5625	GCAUCUUCCAAAGAACAAUUGACGAUA	1	0	transposon	LTR	blood
5626	ACCGAAAUUACAAAGCUACCGAAUG	1	0	transposon	DNA	transib4
5627	AAAGGAAUUGACGUAAGGGCACCACC	1	1	ncRNA		rRNA
5628	GGGAAUACCGGGUGCUGUAGGC	1	no match			
5629	AUAAUCGUUAAAUGUCCAUUCCUGGGA	1	0	no annotation		
5630	ACACACGAGAACGGAUUGUUGA	1	0	transposon	LTR	springer
5631	GUCCGGCGGCGAGAGGGAGAGAACCCAG	1	no match			
5632	AACACCAAAACACAAGUUCGAAUUA	1	0	transposon	LTR	blood
5633	UCGAGAAGCGAACCCGGCGAACUGAAAC	1	no match			
5634	CCAGAUAGGACGCCGAUUUUCCGC	1	0	no annotation		
5635	AAGAGCAAUGGAUGAAUUUUAAGAGA	1	no match			
5636	AUGGUUGCAACGGUAACGGGGAAUC	1	no match			
5637	GUUCGUUUCCCCGGCCGACGCACCA	1	no match			
5638	GAUGCGGGAUGAACCGAACGUAGAGUU	1	no match			
5639	UAAAUACCUUGGGAACAAUUGAUA	1	0	transposon	LTR	micropia
5640	AUCGCUUGAAACAAGCACAAUUACUCGA	1	0	transposon	LTR	diver2
5641	GCAGUCGUGGCCGAGCGGUUAAA	1	1	ncRNA		tRNA

5642	AAAACCUGAAGAUGAUUCUGAGUCGC	1	1	transposon	LINE	Helena
5643	UUGAUACGAAUGAGACCCGCAGUAAA	1	0	repeat		trf
5644	GGGCUCAAACUGUGAGGGGAGUCGUCAA	1	no match			
5645	AACCGCGAAUUCCCCUACAAUGGCGA	1	0	transposon	LTR	gypsy6
5646	UGGGAUUGGCCUUAUCCCAGGAUCGAAA	1	0	coding gene		CG33495-RA|Dup99B
5647	AAAUACCAUAGCAAGCUUCACACGCA	1	0	transposon	Low_complexity	F-element
5648	AGACCAAUAAAUUACUUCUCUCGAAA	1	1	transposon	LTR	Idefix
5649	CGACGCCGUCUCUACCAGGAAUCAA	1	0	transposon	LTR	diver2
5650	CUGCAAUUACAUCCACGUACGGCGAC	1	2	transposon	LTR	Quasimodo
5651	UCUGACUCUCAACCUUUUUUUGGU	1	0	no annotation		
5652	ACAGAACUGGCGCUGUGGGAUGAA	1	1	ncRNA		rRNA
5653	AGGAGGGCGAUUACCACGGCAGGGUU	1	no match			
5654	CGUGACCUGAUUCAUCGCAGUUCCAGUC	1	1	transposon	LINE	Circe
5655	ACUCGCCGGAUCCCGGGGCCGAGGGA	1	no match			
5656	GAAUCUAUGGGUGGUGGUGCAUGGCC	1	0	ncRNA		rRNA
5657	AACGAUGGAUACUAGGCGUUGACCG	1	no match			
5658	ACCCUGAACGCGCCCGAUCUCGC	1	no match			
5659	CUGUUAAACAAUGUACCUGUGAUCAA	1	1	transposon	LINE	Cr1a
5660	AUCGAAUUAAUCGCGAGGCUGCAAGA	1	1	transposon	LTR	mdg3
5661	GCAACCGAUAACGCCCACCGAGGUCCA	1	1	transposon	LTR	ninja-Dsim-like
5662	CCGCGUUGAUGAGUUGAAUCGUUCU	1	0	ncRNA		snoRNA
5663	GAAACGGGAAACUUCAUCCCAGUCGG	1	1	no annotation		
5664	AACUAUCACAACGGAACUCUGACCC	1	0	transposon	LTR	Dm88
5665	AUGCCCCUGAUGAAGAAGAGAUGCCC	1	0	transposon	LTR	ninja-Dsim-like
5666	ACACCUGAACGCGCCGAUCUCGUUC	1	no match			
5667	GUGAUGUCAACGUUUGAUCCCCUGGGU	1	1	transposon	LTR	ninja-Dsim-like
5668	GGUUGUAAAGCUCUUUUGUCCGGGACGA	1	no match			
5669	UGCUUGCACUACAUAUGGUUGAGGGU	1	1	ncRNA		rRNA
5670	AAAGGAGUCUAACACGUGCGCGAG	1	no match			
5671	ACUGGACGAGGCUGGACUGAUCCCUGA	1	0	transposon	LINE	G2
5672	CUGUUGAUAAGGUAACCAGCGA	1	0	coding gene		CG4141-RA|Pi3K92E
5673	AAGAGGGGCGGCCGGGGGCAUUCGU	1	no match			
5674	AAGAACCUGAAACCGUGUGCUUAC	1	no match			
5675	AAGAGAGCAAUUUUGGCCGUCA	1	0	transposon	LINE	Cr1a
5676	CAGAUGUCCCAAGAGAAGUUGGCCC	1	0	transposon	LTR	flea
5677	GCAGAUAAAAAACAAGCGGUUGAG	1	0	transposon	LTR	flea
5678	UUCUUCGGGACUAUCCAGAUAGGG	1	0	transposon	LTR	opus
5679	ACAACAUCAAGAAAACCGACGACAC	1	2	transposon	LTR	INE-1
5680	AAAACCUGAAGAUGAUCCUGAGUU	1	1	transposon	LINE	Helena
5681	UGGGUUCCGGGGGGAGUAUCGUUGC	1	no match			
5682	AAAAAAGGCUAAAUACCGGCACGAGACCG	1	no match			
5683	GUCCCUGCCCUUUGUACACACCG	1	0	ncRNA		rRNA
5684	UCGUCAAAAGAAGAACGGAACCGUACC	1	no match			
5685	CAGAUCUUUCAAAUGUCUGCCCUAUCA	1	0	ncRNA		rRNA
5686	UGCUUAAAACAGCAGGACGGUGA	1	0	ncRNA		rRNA
5687	ACAUAACGGAACCGCUAGCAGAUCU	1	1	transposon	LTR	opus
5688	UUCAAGGUUACCCGAGGAUGCUGU	1	1	repeat		trf
5689	AUGUGCCCAAGGCGAACCCCCGAUCGCA	1	0	coding gene		CG9450-RA|tud
5690	GCGCGCGAAGGUUGGCUCAGGACGGUU	1	no match			
5691	CCAGACGUAAUACUCCGCUACCCGGAC	1	1	transposon	LTR	HMS-Beagle
5692	ACGAGUGUUAUUGUGGGCCGGU	1	0	ncRNA		rRNA
5693	UCUAUCCAUGGCCAGGAUGAAGGUGCCGU	1	no match			
5694	CUGUAACUCGGGUCAAUUGGGGUCAAU	1	no match			
5695	AGCAGGCGAUGCUUGGACCAGUUA	1	0	no annotation		
5696	UGCUAGAAGACGAUUUUCGGGCC	1	0	transposon	LINE	Cr1a
5697	CUGAAGAUAAGUGGAGUUGGAAAGCC	1	0	transposon	LTR	ninja-Dsim-like
5698	UAGUUUGACAUUGUGAAGAGACAU	1	no match			
5699	UCCGAACAUAUCAAGACAAGACCGG	1	0	transposon	DNA	Tc1-2
5700	AUAGGGCGAUCAGUCGCUGGGUGUA	1	no match			
5701	AUUGCCAGGUAGGGAGUUUGACU	1	0	ncRNA		rRNA
5702	GAUAACCAAGGACGUUCACACGAGG	1	0	transposon	LTR	opus
5703	GAUGUCCGGACAGCUGGGCCCGAAC	1	0	Su(Ste)		Su(Ste)
5704	UUCAAGGUUAUCCCAGGAUGCUGU	1	1	repeat		trf
5705	UAUGGCUCUCUCAGUCGCUUCCCGGA	1	0	ncRNA		rRNA
5706	ACUCGCCGAAUCUCGGGGCCGAGGGAGC	1	no match			
5707	GCUGUACUUAUGCAAAAGAAGGACGAC	1	1	transposon	Unknown	RR50960
5708	UGGAGAACCUGGAGAUGGAGGCCGACA	1	0	coding gene		CG42321-RH|CG42321
5709	GGUUUGAAGCGAACCCGGGGAACUGAAA	1	no match			
5710	GCUCGGGGAGGUAGUGACGAAAA	1	0	ncRNA		rRNA
5711	GGAUCCUCAACUGUGAAGGGGAUCCUCAA	1	no match			
5712	CUGGCUCAGAACGAACGCUUGCGGCGCGC	1	no match			
5713	AAAAUUACAAAGACGACGGCCCUACC	1	no match			
5714	GAGAUUCCCACUGUCCCUACCUACUA	1	2	ncRNA		rRNA
5715	GGACCUCAACUGUGAGGGGAUCCUCUAA	1	no match			
5716	AGCACCGAGAGAUUCGCCUACGCCGGC	1	0	coding gene		CG17210-RA|scpr-B
5717	GACGACGAAAUCUAAAGACACAGUG	1	0	transposon	LINE	G2
5718	CCAAGGCGCUUGAGAGAACUCGGGUGAA	1	no match			
5719	AUGCAAGGUUACCCCAGGUGCUGUGC	1	no match			
5720	ACAGUUCAGUAACCCGAGCGACUGUUCAAU	1	no match			
5721	AAGACCCCCGAAACCAGACGAGCUACC	1	no match			
5722	GCUCUCGAAUGUAUGCCCGAUAUAA	1	1	transposon	LINE	R1-element
5723	AUCUCAUCGUCGUAGAACAAGCCCCGA	1	no match			
5724	CCCCGCGGAAAGACGGAAAGACCC	1	no match			

5725	GAAGCAAAAUUCACCAAGCGUUG	1	0	ncRNA		rRNA
5726	AAGUCCGAGAGAGGCUUCCGAGUC	1	0	transposon	LINE	TART
5727	UGCUCUCGAAUGUAUGCCCGUAUAUACG	1	no match			
5728	UCGGCCCCGAAACCCUUCUGGUGUGUU	1	no match			
5729	CAGGCAAAAAGGACAACAUUCACCCAU	1	1	transposon	LTR	HMS-Beagle
5730	GGUCCGAACAUAAGUGUGUAAGA	1	0	ncRNA		rRNA
5731	UGGAGUGAUUUGUCUGGUUAAUU	1	0	ncRNA		rRNA
5732	GGGGAUGUAGCUCAGAUGGUAGA	1	0	ncRNA		tRNA
5733	CACAGACUCGGAACGCCUCAUUGG	1	0	coding gene		CG31359-RA|Hsp70Bb
5734	CAAGCUAGACUAUGUAUCCCCGGGUAAU	1	1	transposon	LINE	Doc
5735	AAAACCUGAAGAUGAUCCUGAUUCGC	1	1	transposon	LINE	Helena
5736	AAGUACAGUAUCCUGGGAUAUGGAAGU	1	no match			
5737	CGGCACGAGGCCAUCCACGGACCCAG	1	0	coding gene		CG13484-RA|CG13484
5738	AUGCUCGAGAUGACAACCUUGGACCUGA	1	0	transposon	LTR	micropia
5739	GGGGAACUGAGGAACGCAUUUCAAA	1	0	transposon	LTR	invader3
5740	ACCUUCGUCCAAGAGAACUGUGCCCG	1	2	transposon	LINE	R1-element
5741	GUCUGAGAUGGAGGCCCUUGACAAA	1	0	coding gene		CG5968-RA|CG5968
5742	GAUGGUUGGCAGGGAUAACAUCGU	1	0	transposon	LINE	G2
5743	GUGCAACGACGAGAUGACCAACUUGA	1	1	coding gene		CG33233-RC|CG33233
5744	UGAAGCCGGUGAUCGAUAGUUCCUCU	1	0	repeat		trf
5745	GACAUCUUAAAGGGCGAAAAUGGA	1	0	transposon	LTR	springer
5746	GAACAGGUCAACCGAAUCACGUCAG	1	0	transposon	LINE	Doc
5747	ACUCGCCGAAUCCCGGGGCCGGGGA	1	no match			
5748	GGGGAAAGAUAAUCAUGUUGCAGACGC	1	0	transposon	LTR	mdg1
5749	UGUUUCAUCGUUGACGGCUCGGGC	1	no match			
5750	UCCUGACGAAGAGCGCAAGAAGAGGG	1	0	transposon	LTR	opus
5751	CGCGUGAGACCGUACCCGAAACCGAC	1	no match			
5752	CAAGCAUUCGACCGAGUCUGGCUCGU	1	1	transposon	LINE	F
5753	ACCCUGAACGCGCCCGAUCUCUU	1	no match			
5754	CUUCAACUGUGAGGGGAUCCUCAA	1	no match			
5755	AGGAUACUAAGCAGGGACGUUGAUGA	1	1	coding gene		CG12717-RA|CG12717
5756	UGCCGAACAUCCUAUCUUGACGAU	1	no match			
5757	AAACCUCAGAACGUCUUGAUUCG	1	2	transposon	LINE	RR49783
5758	UUUGGUUGCAAAGCUUAGGCUGGGUUA	1	no match			
5759	AUAUCCGAGGCCCUGUAAUUGGAAUG	1	0	ncRNA		rRNA
5760	CCCGUUAGGAAUAAGUAUAUCAUUGC	1	0	transposon	LTR	flea
5761	AACCAUGAGACCGAUAGUAAACAAGUAC	1	0	ncRNA		rRNA
5762	AGAACGCACCCGCUUGACCUUACUC	1	0	transposon	LINE	G2
5763	GAGGACUUGGGCGAUUUAAGUGG	1	0	Su(Ste)		Su(Ste)
5764	UGUUCUUCUAGGGAUGGCGUGAA	1	1	Su(Ste)		Su(Ste)
5765	UCUCCCCGCGGAAAGACGGAAAGACCCCCUA	1	no match			
5766	UGCUCGGACUACAUAUGGUUGAGGGUUGUA	1	1	ncRNA		rRNA
5767	AUAAGGGUUGCACACCGUCCGGUAU	1	0	transposon	LTR	gypsy12
5768	CUCAGCGAUGUGUGGGGCAAGACA	1	1	Su(Ste)		Su(Ste)
5769	ACCCAUGUGAAAUACUUCACAUCGA	1	0	transposon	LTR	opus
5770	UGCGGCUGGAUCACCUCCUUUCU	1	no match			
5771	CGAAGCUAAAACGCUGGAAGGCUCGUA	1	0	transposon	LTR	opus
5772	CCAGCUCGGGGAGGUUGACGAA	1	no match			
5773	GCUAACAAAGACUCGCAGCGGCCACG	1	0	transposon	LTR	gypsy6
5774	GACUGCAGAUCGAGAGGUCCCCGGUA	1	1	ncRNA		tRNA
5775	CUGUGAGGAGGAUCCUCAA	1	2	coding gene		CG8325-RA|l(2)k14710
5776	GCUGCAGAUCGAGAGGUCCCCGGUUU	1	2	ncRNA		tRNA
5777	ACGAGCGCACGUGUUAGGACCCGAAAGUA	1	no match			
5778	CAAGCUAGACUAUGUAUCCCCGUUAA	1	2	transposon	LINE	Doc
5779	UUUAGUUCAAACGAAAGACCGGUGCCC	1	0	transposon	LTR	HMS-Beagle
5780	CUGAGAAUUACGAUCUUUUAUUAGGA	1	0	transposon	LTR	McClintock
5781	AAUUGGAAUGAGUACACUUUAAA	1	0	ncRNA		rRNA
5782	UGAAGGCCGAAGUGGAGAAGGGUUCC	1	no match			
5783	GAAUACUUUAUAUAAUUACCGCAUU	1	0	transposon	LTR	flea
5784	CACCAGCAGAACGCGCAGAGCGGAG	1	0	coding gene		pseudogene
5785	GAGGAAUUUGAGCUGCGUUGACGCCGA	1	0	transposon	LTR	ninja-Dsim-like
5786	CAUACCUUGACUGGCAGUCCCGG	1	0	transposon	LINE	F
5787	CCGAACUUGAGACCGUCCCUGGAA	1	2	Su(Ste)		Su(Ste)
5788	AAUCGAUGACGAAUCGUGAGGACGAGC	1	0	coding gene		CG32697-RA|l(1)G0232
5789	UAGCGGACAACCUAGCCAGUGCAU	1	no match			
5790	ACGUCCCUGCCCUUUGUACACACCGCCC	1	0	ncRNA		rRNA
5791	ACAACAAACAACACAUACCCGCUGGCCG	1	1	transposon	LTR	GATE
5792	UUGUCAAGAAUAGUGUGUUAGAGGGA	1	0	transposon	LTR	invader4
5793	CUGGUGGAAAUGCCAAUGCAGCAGAGC	1	0	coding gene		CG10373-RA|CG10373
5794	AGAGCUAAACUCGAACACGAUUUCGGU	1	1	transposon	LTR	micropia
5795	GCUGGCAGGACGAGGACGAGAAGGC	1	0	coding gene		CG8004-RA|CG8004
5796	AACGAGUCAUGUUAAGGCAGUUACCUUA	1	0	no annotation		
5797	UCGAUAUUAUCCCAGCACUACGAAG	1	0	no annotation		
5798	UUCGUGGAGUGAUUUGUCUGGUU	1	0	ncRNA		rRNA
5799	UCAAGUCAACGAUCGUGGCAGGACGCA	1	0	transposon	LTR	micropia
5800	CAACCUAAUCACGUUCGGUGCCCGUAA	1	2	transposon	LINE	Rt1a
5801	CUGGGAAACGGGGCGCGGCCGGAAAGG	1	no match			
5802	CCCGAAAGCUACGAGGUAACGUA	1	no match			
5803	CCCAACAAGACAUCCGAUUAGUUCAUU	1	0	transposon	LINE	Circe
5804	AAGGAAACCACUGUGGAGGGAACCAGC	1	0	coding gene		CG32473-RA|CG32473
5805	AUGCCUGAACGCUCUAAGGUCGU	1	no match			
5806	GAACGACCAUAAAGAAGCCGUCGAGA	1	0	ncRNA		rRNA
5807	GGAUCAACUGUGAGGGAAUCCUCAA	1	no match			

5808	GCGGGGCGCGGGACAUGUGGCGUACGG	1	no match			
5809	AGUGCGGUAACGCGAUCGAUCCCGGAG	1	no match			
5810	CAUGAUAAAUAGGGGCAAUUUGC	1	no match			
5811	UGGGGAAGCUCCGUUUCAAAGGCCUGAU	1	no match			
5812	AGCUCGAGGAAUACGGCCUUACU	1	0	transposon	LTR	ninja-Dsim-like
5813	CUAGAAAAAACGAGGUUCACCCAA	1	0	transposon	Low_complexity	F-element
5814	GCCUGAUGACCAUGCGAGUUGG	1	2	no annotation		
5815	GCUGACUUGAUCUCGGUGUUCAGUAC	1	0	repeat		LSU-rRNA_Dme
5816	GUUGCAUCCGGUGCGUCCUUAACAGUCC	1	no match			
5817	UUUGAUAAUAACCCCAGGAAUUAAAAA	1	1	transposon	LTR	opus
5818	UUCGAUACUGUUUGGCUUGAGUUUG	1	no match			
5819	GUCGAAGCCCCGGUUGAUGCUCAAGU	1	0	coding gene		CG1099-RA|Dap160
5820	AGCUGGUUCCUGCCGAAGUUUCCCUC	1	1	ncRNA		rRNA
5821	GAGCUCGUUUGAUUUUGACCAGGAU	1	0	transposon	LINE	X
5822	CGGUACCGUCCGUCGCCAGGCCGUCG	1	0	coding gene		CG8922-RA|RpS5a
5823	UACAUUAUUAUUGUGUAUCGACGGGA	1	1	no annotation		
5824	GCAACCCCAGAUGUCCGUGAUCCAGU	1	1	transposon	LTR	ninja-Dsim-like
5825	ACGAGUAAUAUGAGCACCCAGAGU	1	1	transposon	LTR	opus
5826	GGAGACAAGAUGCCAGAUGUACCAA	1	0	no annotation		
5827	AAACCUGAAGUAUGAUCCUGAGUCGC	1	no match			
5828	AACAGGGAACGUGAGCUGGGUUU	1	no match			
5829	AACCGACUACAAAGAUGCCAUACGA	1	0	transposon	Unknown	RR50960
5830	GAUCGAGGCGCUCAACCCAAGCUC	1	0	coding gene		CG32344-RA|CG32344
5831	ACCCUUGAUAAACGUAUGACCUUCGU	1	2	transposon	LINE	BS
5832	UAAUAAUCCACGGCUGACUCCAGU	1	0	transposon	LTR	GATE
5833	UCGAGCUCUCGAGAAAGGGACGCGAC	1	0	coding gene		CG6527-RA|CG6527
5834	AGGCCCAGUGUUCGGUCCACUGCU	1	no match			
5835	CUAGCAAGAACUCGGAGGAGCAUUUAG	1	0	transposon	LTR	flea
5836	GAUGUCGUAUCACCCAGGACGCCCGU	1	1	transposon	LINE	Rt1b
5837	CCAUCAGGUCGGAGCGUGGAAGAACACCC	1	no match			
5838	UAAGGAUGAUGAUACCGACUAAAGGCA	1	0	no annotation		
5839	CUCGCAAUCCCGGGCCGAGGGAGU	1	no match			
5840	CUCAAACUGUGAGGGGGAUCCCUCAA	1	no match			
5841	GUGUGAUUCCCGUGUAGACGUCGCUUG	1	1	coding gene		CG16741-RA|CG16741
5842	GCUAAACCAUUCGUAGACGACCUGC	1	no match			
5843	UCGGGAUAAGGAUUGGCUCUGAAGA	1	0	ncRNA		rRNA
5844	CGACCACGUUGUAAUGCCACCGAGA	1	1	transposon	LTR	ninja-Dsim-like
5845	AACGGAUAAAAGGUACUCCGGGGAUAAC	1	no match			
5846	AAAGGCAUGCCCGAUGUCCGCGAGA	1	0	transposon	LTR	opus
5847	CAAACCAAUCCCAACCAAACCCGC	1	1	transposon	LINE	G2
5848	UGCAACGUUCACCAAUCUUCGUCAGC	1	0	transposon	LTR	3S18
5849	AAAGAGAGCAAUUUUGGCCGUCG	1	0	transposon	DNA	1360
5850	ACAGCUUGUUCACUGCCCGCCUUAUU	1	0	transposon	LTR	diver2
5851	AGAAAGCCGAACUCGACGCGGUUGUC	1	0	no annotation		
5852	UUAACAACGAUGAUGAUUCUGGCCCAU	1	1	transposon	LTR	Max
5853	AGUGGAGCCGUACCUGUUGGUUUGUCCC	1	0	ncRNA		rRNA
5854	GCCGAACGUAGUCUCGAGAGCGA	1	no match			
5855	ACGAGUAAUAUGAGCGCCCAGAGUGU	1	1	transposon	LTR	opus
5856	GGAGAGGUUCCCGAGCGGCCAAAGGGAUCAGA	1	no match			
5857	AAGCAAUUAUCCGAUACCGCUUAUC	1	2	transposon	LTR	flea
5858	GAAGACCAAGGCCAAGCACUUCUCGCCC	1	0	coding gene		CG1193-RA|CG1193
5859	UCCGAGGAAUCUGGGUUCAAUAUUU	1	no match			
5860	UCUGAUGAUGUGUUGUUGCCAUGGU	1	no match			
5861	AGGUCAAAACUCCCGUGCUGAUC	1	no match			
5862	GUAACAACGAAGAUGAUUCUGGCCCAGA	1	0	transposon	LTR	invader4
5863	GAGAAACUGACAAAGGCCCUUGAUGU	1	1	transposon	LTR	Stalker
5864	AUAUGCUAAAAGUUAUCCUUACCCAGUC	1	1	transposon	LTR	opus
5865	UGCGAAAGUAAAAGAUGUCCAGCGAA	1	1	transposon	LINE	Rt1b
5866	UAUGUGAGAACUCGAACUUGCCUA	1	0	transposon	DNA	transib2
5867	CUAGCCUUGCCCACUUCCCGCUAACA	1	0	transposon	LINE	X
5868	UGCUCUCGAAUGUAUGCCCGUAUAU	1	no match			
5869	UCGAGAUGGAGAGGCUUCAGAGUG	1	0	transposon	LTR	ninja-Dsim-like
5870	CAAAGUAUCACGAGUUUGUUAUAU	1	0	ncRNA		rRNA
5871	GCCGAAUGACCAAGGGUUCCUGGGC	1	no match			
5872	CGAGACUUUGAACUAAAACAGUUACC	1	no match			
5873	CGACCUCAGAUCAGACGUGGCGACCCGC	1	no match			
5874	UGAGGCGCUCGGUGUCGGUGA	1	no match			
5875	UCCGGACAAACACACAAGACCCGAUCU	1	0	transposon	LTR	opus
5876	AAACCAUUCGUAGACGGCCUGCU	1	no match			
5877	CACAGAGUACGGGGCUUACUGAG	1	1	transposon	LINE	baggins
5878	ACAGAGAGAAGACUCACCCGACGA	1	0	no annotation		
5879	CAAGGUGUUGUAAGUGGUUGAGC	1	0	ncRNA		rRNA
5880	ACCCUGAACGCGCCCGAUCCCGU	1	no match			
5881	AUAACCUUGAGACUGAGGGAGGGGCAA	1	0	transposon	LTR	opus
5882	GUGCCAGGCCCGUAUAACGUUAAUGAU	1	0	ncRNA		rRNA
5883	ACGGGCUGCCCGAGCGUCACCUGCAC	1	no match			
5884	AGCAGACCAAUCCCAACCAAACGAUACUGUUCGG	1	no match			
5885	AAACCGAGAGGAGGAGGAACGUCGCGU	1	2	coding gene		CG13168-RB|CG13168
5886	GGCUCCUCAACUGUGAGGGGAUCCUCAA	1	no match			
5887	AAUUCCGUAAACGAAGCCCAAAGUUA	1	1	transposon	LTR	Rt1c
5888	UGAUUCCGAGGAACCAAUUGAUCUAG	1	1	Su(Ste)		Su(Ste)
5889	UGGCUCUGAAGAUUGAGAUAGUCGGG	1	0	ncRNA		rRNA
5890	CUCGUCAGAAAACGUUACGCCGA	1	1	repeat		trf

5891	UCUCAUCGUCGUAGAACAAGCCCGAUGA	1	1	Su(Ste)		Su(Ste)
5892	AGCAAAGUCGGCUCCGAGGCCCCUGAC	1	0	coding gene		CG11373-RA|CG11373
5893	AAUCAUGCCGACGGGCGCUGACCCCCUU	1	no match			
5894	AAAGCCGUACCUGUUGGUUUGUCCCAUU	1	2	ncRNA		rRNA
5895	UUCUCAGAGAGCAAAUUGGCAAG	1	0	transposon	LTR	springer
5896	CGAGUUAAUAUGGCACCAAGCUGA	1	0	no annotation		
5897	AAGUGGUACGCGAGCUGGGUUU	1	no match			
5898	AUCCGCACCCGUCUCCCCCACAUCCGUA	1	0	coding gene		CG9821-RA|CG9821
5899	AAGGACACGCAUGCACGAGCGGUGA	1	0	coding gene		CG32849-RA|Hex-t2
5900	CACCAUAGUGAACUGGCUCGAGAU	1	0	transposon	LINE	X
5901	AGGUGGACACUCAGCCUAUUCGCACUC	1	0	coding gene		CG10726-RA|barr
5902	CCUGCAGAGAAUGCACCCGCUUGA	1	1	transposon	LINE	G2
5903	CUGGAGAUCAAGGCUCUCAGCCCGGC	1	0	transposon	LINE	Rt1b
5904	UGCUCUCGAACGUAUGCCCGAUA	1	1	transposon	LINE	R1-element
5905	CAGGAGACACGUCGCAAGAAGGAU	1	1	coding gene		CG11703-RA|CG11703
5906	AAUAGUGUGUUAGAGGGAGAUGA	1	0	transposon	LTR	invader4
5907	ACCACCCUGAACGCACCCGAUCUCGUC	1	no match			
5908	UAGAUGUCUGUACGAAAGCGAGA	1	0	transposon	LTR	Dm88
5909	ACGCAAACUUCUCAACAGGCCGUAC	1	0	ncRNA		rRNA
5910	CGGGAAACCUCACCCGGCCCGGACACGG	1	no match			
5911	AAAAACCCGAAGAUGAUCCUGAGUC	1	2	transposon	LINE	Helena
5912	CACAUCAACAGAAAGCGUAUCCAG	1	0	transposon	LINE	G2
5913	AGAGAGAUAUGGUAGAUGGGCU	1	0	ncRNA		rRNA
5914	ACGAGUAAUAAUGAGCGCCCAGAGUGA	1	no match			
5915	GAUCAUCCAGAGUCUGCUCAACUU	1	0	coding gene		CG9917-RA|CG9917
5916	GUGGAAAUGAAGAACUGUGAGG	1	no match			
5917	UCCCAGAUAUGUUACCUCGUCGGCU	1	0	transposon	LINE	Doc
5918	GGCCAAGAGCUGAUGACGAGUUAACUUU	1	no match			
5919	GCGCCGGCGGCUUUGGUGACUCUAGAU	1	no match			
5920	GAGAAAACUAUUCCUGACCGGUACCC	1	0	transposon	LINE	accord2
5921	UCAACUGUGAGGGGUCCUCAA	1	no match			
5922	GCGCGGGUCAACGGCGGGAGU	1	0	ncRNA		rRNA
5923	GAGCCAAGACCGUCCCCGAGAAGCC	1	no match			
5924	AAACUUCAUCUCAGUCGGAUUAGCCCA	1	0	coding gene		CG41521-RA|CG41521
5925	UGGAUACUAAGCAGGGACGUUGAUGA	1	0	coding gene		CG12717-RA|CG12717
5926	AGGGUAAUACAGGUGACUGGGGCU	1	no match			
5927	GCAUCAUUCAAAGCCAGCUGGAUGCCC	1	1	transposon	LINE	G
5928	CUGAUCGAGGCGAAUAUGACGGUGA	1	0	transposon	LTR	gypsy
5929	UUGCGUAAAGCUGGAUUGACAAUU	1	0	transposon	LINE	Circe
5930	GCCAGAUUGUGAUAAAGUACCGGGC	1	0	coding gene		CG11739-RA|CG11739
5931	CUAACUGGAACGUACCUUGAGCAUAU	1	0	ncRNA		rRNA
5932	GGCGCUAAACCAUUCGUAGACGACC	1	no match			
5933	ACACCCGUUCCCGUCCGAUCACGGCAGU	1	no match			
5934	CGCCAGCUAAGGUCCCUAAAAUUGGCUA	1	no match			
5935	ACGAAACGCAACAACAAAAUGGACGGA	1	0	transposon	LINE	G2
5936	AAAGGGCAAAUGCUGACUUGAU	1	0	ncRNA		rRNA
5937	UCCUGGGGUAACCUUGAACUUCGU	1	0	repeat		trf
5938	CAUGACCUUAUUGACGACUUAUCGC	1	0	transposon	LINE	Cr1a
5939	GUUCCUCGGGAUCAAGGGCAACGAU	1	1	coding gene		CG40635-RA|CG40635
5940	UCCUCGAUAGUAUAGGUGGUUUAGUAUCCC	1	no match			
5941	CGAUGAUCGAUGACGAGCUACGAAGGG	1	0	transposon	LTR	ninja-Dsim-like
5942	UGUAUCGAUAUGAGCAGCAGAACUCCG	1	0	no annotation		
5943	CCCAAACGCAGACAUCGAGCGGUCAAU	1	0	transposon	Low_complexity	F-element
5944	GCCGCCUGGAUACCGCAGCUAGGAAU	1	no match			
5945	UUGUUCCCCGGAUAGUUUAGUUUAC	1	no match			
5946	CUACAUACGAGGCGUAAUUGGCA	1	0	transposon	LTR	Quasimodo
5947	GAGGCAAUCACCGAGAGCAUCAAC	1	0	transposon	LINE	F
5948	GGGGAGAAUGAUAUGCACCGAAAGUA	1	2	Su(Ste)		Su(Ste)
5949	CAACCAAUCCCACCAAACCCGU	1	no match			
5950	UCCAACUGUGAGGGGAUCUCUAA	1	no match			
5951	ACUUGAGACCGCCCCUGGACGACC	1	0	Su(Ste)		Su(Ste)
5952	CUACGCCGAAUCCCGGGGCCGAGGG	1	no match			
5953	GAUGUACUCCGACGAGUGGACCAGUUC	1	0	coding gene		CG5186-RA|slim
5954	CACAAGAAAAGUCCUUGCUCUCCGGGA	1	0	coding gene		CG40609-RA|CG40609
5955	UCUGGUGGGCGAGGAAAAGUUGAGGAA	1	0	coding gene		CG32473-RA|CG32473
5956	GCUUCAUCCAAGCGAACUGUGGCCGA	1	0	transposon	LINE	R1
5957	UGAGAAGACGGUCGAACUUGACUU	1	no match			
5958	GCAUACGAGGAAGCAUACGAAGAAGCAU	1	0	transposon	LTR	flea
5959	UGUGCUAGAAGACGGAUUUACACGAGG	1	1	transposon	DNA	1360
5960	ACCUGAUUGAUAGCGUAGACGACGAU	1	1	transposon	LINE	Rt1a
5961	CGGCCAUAGUGAUCCGGUGGUCCG	1	no match			
5962	CAAACUCACAAUUAGGUGGGUCCCGG	1	0	transposon	LINE	baggins
5963	ACCUGGCGUAGAGGUUAACCGUGAUC	1	0	ncRNA		snRNA
5964	AUCCUGAACGCGCCCGAUCUCGU	1	no match			
5965	GACCACAUGGAAACCAAGGUCCCUU	1	no match			
5966	UGCUCUCGAAUGUAUGCCCAAUAU	1	0	transposon	LINE	R1-element
5967	AAGGCCAUAAAGGAAUACCCAGAGCC	1	0	transposon	LTR	gtwin
5968	GCACCGGGAACAUGUGGCUGCGAG	1	no match			
5969	AAAACCUGAGGAUGAUCCUGAGUCG	1	1	transposon	LINE	Helena
5970	UGAGACUACAAACGGAUUCGCGGACC	1	no match			
5971	AAUUCCGUCAACGAGGCCCAAAGU	1	1	transposon	LTR	Rt1c
5972	CACAUCGACGGGGAGGUUUGGCAC	1	no match			
5973	ACCCAGCGCAAGAUCAUCCCAAUCGA	1	0	coding gene		CG4904-RA|Pros35

5974	ACAACCAAGACGUUUGGCAGAGUG	1	0	transposon	DNA	transib4
5975	AGAGCAAAGACAAGACAGAGUUUACCA	1	0	no annotation		
5976	AAUCAUGCCGACGGGCGCUGACCCCCC	1	no match			
5977	AGCGGGGAAAGAAUACCCUGUUGAGCUUG	1	2	repeat		LSU-rRNA_Dme
5978	AAAACCUGAAGAUGAUCUGAGUCGCAU	1	no match			
5979	AAGGAGAGCAAUUUUGGCCGUCA	1	0	transposon	DNA	1360
5980	AUGAACCGUAUAUAAAGAUGUUGAU	1	0	transposon	LTR	diver2
5981	UGCCGUGGGACGCCGAAGACCAGGAGC	1	no match			
5982	AGUUGAACUUGUGCUUCAUACGGGUAGU	1	0	ncRNA		rRNA
5983	UACCACCCUGAACGCGCCCGAUCUCGU	1	no match			
5984	CGGUACAUCGUUUCGCCGCGUGGCA	1	0	coding gene		CG13591-RA|Ssl
5985	AUCUGGACACCCGGGGCACUGGACU	1	0	coding gene		CG32064-RA|CG32064
5986	UGCGAGAACAGCGGUUUGCCCUUU	1	1	transposon	LTR	diver2
5987	UCAACCGGACGGGCCUAGAGUUAUUCCC	1	0	coding gene		CG40635-RA|CG40635
5988	AAGACCGUAAACGAAAAGCUAAG	1	0	transposon	LTR	Idefix
5989	ACAUUCCGAAGGGCAGCAAACCGU	1	0	transposon	DNA	invader2
5990	CGACGACGAUCAACGGGAGCACGA	1	0	coding gene		CG11265-RC|CG11265
5991	UCGGCCCGAACCCUUCUGGUGUAUU	1	1	transposon	LINE	baggins
5992	CCGAAUACGAAGAACAGGCAGAAA	1	1	transposon	LTR	gypsy12
5993	GCGGAUAUAAAAGCAGCGGGGCGG	1	0	transposon	LTR	opus
5994	AACCGAUAUGGUUGGACCACAAUCUUUUU	1	0	ncRNA		7SLRNA-RA
5995	CAGUCCGAAACACAUCCCCCGAUUA	1	0	no annotation		
5996	UCUCAGUACCCGGAGGAAAGGAC	1	no match			
5997	CCCAUGCCGAGCGCAAAAGUUGUUU	1	2	transposon	LTR	opus
5998	CGAAUAGUGAUUGCGAACCCCCGAA	1	0	transposon	DNA	S
5999	GCAGCCCAGUACUCCGGAGCCCAGGU	1	0	coding gene		CG11901-RA|Ef1gamma
6000	CCUCGUCGGUGUUGUCUCCUGGG	1	0	coding gene		CG12351-RA|deltaTry
6001	UGCCCCGAAGGUCUUCCAGGCUCUGU	1	no match			
6002	UGCAAGCUCUUGACUGAAGUCCCAGU	1	no match			
6003	CACCGAGCGGGGCAACCUGAUCUUCGA	1	0	coding gene		CG2017-RA|CG2017
6004	AGCGAAAUGACUCUGUCCCGAUUCAAG	1	0	transposon	LINE	R1-element
6005	UCUGAUGUUACGCCCGAAGAGAUAAC	1	0	transposon	Low_complexity	F-element
6006	CUUAGAUGGGGGACCGCUUGGG	1	0	ncRNA		5SrRNA
6007	AGUGCCAGGUGGGGAGUUUGACUGGG	1	2	ncRNA		rRNA
6008	UGCUCUACGAAUGUAUGCCCGAUAU	1	no match			
6009	AGAGCCGGGCGUGGAAUGCGAGU	1	no match			
6010	ACGAACUUGAGACCGCCCCUGGACGACC	1	1	Su(Ste)		Su(Ste)
6011	GGAUCGCGGUUCUAUUUUGUUGGUU	1	no match			
6012	UCCCGUAAACCGCCAUCAGCAACACACCC	1	0	transposon	Low_complexity	F-element
6013	UGCCGAACUUGACGACCCACGAUA	1	0	Su(Ste)		Su(Ste)
6014	AUCCAGGACGAAUAGGCGGGGCACCCAGC	1	0	coding gene		CG31642-RA|CG31642
6015	ACGAGUAAAUAGUGAGCGCCCGAGAGU	1	no match			
6016	GAACUUGAGACCGCCCUGGACGA	1	no match			
6017	UUAGAAUGGAUCGGGUUGAAAUUCU	1	0	coding gene		CG41053-RA|CG41053
6018	UCAAUACCUGGCGAGGACGCAUCUUC	1	0	no annotation		
6019	AGAAGACGUAUCUGCCCAGGUUGGUGG	1	0	coding gene		CG13340-RA|CG13340
6020	CAGAUCAGCAAAGUACGAUGGGGA	1	1	no annotation		
6021	AGAACGAAAGUCGGAGGUUCGAAGAC	1	no match			
6022	CGCCAGCUAAGGUCCCUAAAAUUGGCUAA	1	no match			
6023	AGACGACGCCGCCAGUUCGAGAGG	1	1	coding gene		CG3219-RA|Klp59C
6024	CCAAGAAAAAAUCAAAAACGCCCAGA	1	0	transposon	LTR	blood
6025	AUGUUUUCAUCGUUAGACGGCUCGGGCG	1	no match			
6026	CAGACCCAACCAAUACAUAAUACAU	1	0	transposon	LTR	297
6027	UGGGGCCCCGAACCCUUCUGGUGUGU	1	no match			
6028	UCGCUCGAUGUCGUGAGGUGACC	1	0	no annotation		
6029	ACGUAAACAACCCCAGGCUGACA	1	0	transposon	LTR	Rt1c
6030	AAAACCUGAAGAUGAUCCUGAGUCCA	1	2	transposon	LINE	Helena
6031	UCGAUGGCAAACAGGGUUAAUAUUC	1	no match			
6032	AUCCUCAACUGUAGGGGAUCCUCAA	1	no match			
6033	ACUCGCCGAAUCCCGGGGCCGAGGAGC	1	no match			
6034	AAACCUGAACGAUGAUCCUGAGUCGUA	1	no match			
6035	AGCCGUACCUGUUGGUUUGUCCCAU	1	0	ncRNA		rRNA
6036	CCCAAUGAGAUGGCCUUUGGCGAGA	1	0	transposon	LTR	Quasimodo
6037	CUGCGGAAGUUCAAUUCCCGGUAUGGG	1	no match			
6038	AUUAAUACGAAAACUUACAAACAAAUA	1	1	repeat		trf
6039	UAGAUGUCCUGGGCUGCACGCGCGC	1	0	ncRNA		rRNA
6040	UGCUCUCGAAUAUUUGCCCGAUA	1	2	transposon	LINE	R1
6041	GAAACUCAACUGUGAGGGGAUCCUCAA	1	no match			
6042	UACCGAUGGCGAAGGCAGCCCCCUCG	1	no match			
6043	CUGUAAAUAAUUCAGCAGCACUGACCGU	1	0	transposon	LINE	Doc
6044	CAACCGAUAUGCCAUCUUGGCUGA	1	2	transposon	LINE	Doc
6045	UCCAACUGAAGGAGUACGAAUGAUCG	1	0	transposon	LINE	R1-element
6046	GCUAAUACCGAAUAUGACUCCGACCGUC	1	no match			
6047	GGUUCGAUUUCCGGGAGAGGGGAGCCUGAG	1	no match			
6048	UCGAAUCAACAAAUUGCGCCGAUAU	1	0	coding gene		CG41545-RA|CG41545
6049	CGAGAUAGUACGAAUCGGUUACG	1	0	transposon	LINE	I
6050	AGGAGGCUAGGAGAAAACAUUUGGGA	1	0	coding gene		CG12324-RA|RpS15Ab
6051	CCGACGGGGCGGAGUACGAAUGC	1	0	coding gene		CG8184-RB|CG8184
6052	CUCUGAGAAGCGGACGCGUGA	1	no match			
6053	CGAGUUUCGAACGCACCGAAGAUC	1	0	transposon	LTR	INE-1
6054	AACGCUUAUCACCCCGGAAUUGGU	1	no match			
6055	AGAGGUAGACGACCGGGCGAUGGUU	1	2	transposon	LINE	R1
6056	UGUGCGGCCUGCUGGUGUUCACC	1	0	coding gene		CG33931-RA|Rpp20

6057	AUAAGUAGAACACUUUUGUUUUUGG	1	no match			
6058	CGACCAAUGACACCCAAAUGCUGU	1	0	transposon	LTR	Dm88
6059	ACUCUCAUGCCUGGUCGUACUCAU	1	no match			
6060	GCACCCUGGAUCGGACUCGCUAGUCA	1	0	transposon	LINE	INE-1
6061	CUAACCCUAAAAUACGAUAGACACC	1	0	transposon	LTR	INE-1
6062	UACAAUAUCAGAAGUGGGCCGAUCACGA	1	0	transposon	LTR	rooA
6063	CAAACCGACAUGCUAUCUUGGCUGA	1	1	transposon	LINE	Doc
6064	CGCCUUAGACGUAUCGGGAGUCC	1	no match			
6065	UACCGAACGGUGCCUGGUUACGAGCUCC	1	0	coding gene		CG17344-RA|CG17344
6066	GGUAAUCAGCGCCGACACCAGAGUGGA	1	0	no annotation		
6067	AACGAGCGCAACCCUUGUCCUUAGU	1	no match			
6068	UGCUCUCGAACGCUUGCCCGAUAUUAC	1	no match			
6069	AGUCGUCUCAGUUCGGAUUGUCCUCU	1	no match			
6070	CAAGACCCGACGCAACCACCUGACAC	1	0	coding gene		CG11241-RA|CG11241
6071	GUCAUGCGAGGCGAACCGGGUGAACU	1	no match			
6072	AAGACAACUAUGACAUAUGAGACGGA	1	0	transposon	LTR	gypsy3
6073	UAGCCCCGUACGUGAAAUGACCUGUGUG	1	no match			
6074	GUAACAACGAAGAUGACUCUGGCCCAG	1	0	transposon	LTR	invader4
6075	UAAUAAAGGUGACCGGCAACGACGU	1	2	no annotation		
6076	CAGCUAGACUAUGAAUCCCCGGGU	1	2	transposon	LINE	Doc
6077	UUGGAACGAAAUUGGCCUGAUUAGCGG	1	0	transposon	LTR	GATE
6078	ACACGGGAAACUUACCAGGUCCGAA	1	no match			
6079	AGUUCCCGUGGUAGGCAGUAGUGGGAU	1	0	ncRNA		7SLRNA-RA
6080	GAGAAUAUACACAGAGACUUCCAAA	1	0	transposon	LINE	Doc3
6081	CGGGACGUUCGCCAAUUUCUGUGGCC	1	0	coding gene		CG5193-RA|TfIIB
6082	UGGAAAGCUAUCCGUGACGCAGACC	1	no match			
6083	UAGCUCAAAGGGACGGCGGAUGAUCGA	1	1	transposon	LINE	G2
6084	UCCAAUAUCACCGCGGAGACUGUCA	1	1	transposon	LINE	Doc4
6085	ACCCUGAACAAUACACUACACCGCA	1	0	transposon	LTR	mdg3
6086	GUCUCAUCGUCGUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
6087	CAACAAAGCACCACGGACAUGACGUCA	1	no match			
6088	CUGACUGGCUCAGCGUGUGCCUACCCUU	1	no match			
6089	UAGGCAGAUGUGCCCGAGCGUGGGAC	1	0	transposon	LINE	Cr1a
6090	GGUCAAUUCCGAUAACGAACGAG	1	1	ncRNA		rRNA
6091	UACGACGAGAACGAGAUCAGCGCAA	1	1	transposon	LTR	frogger
6092	UCCGCAGAGGGAGCCUGAGAAACGG	1	1	ncRNA		rRNA
6093	UUCAAGGUUACCCCACGGAUGCUGU	1	no match			
6094	AGGAUACUGAGUCGAUUACCGAGGGC	1	0	transposon	LTR	G4
6095	GCAUGAUUCACGCCCGAUACAUCAAGG	1	1	coding gene		CG33247-RB|Ste:CG33247
6096	GCGGCUAGACGGAAAGACCCCGUGA	1	no match			
6097	CACCGGGCGGACCUUGCUGCGGACCCUG	1	0	coding gene		CG17935-RA|Mst84Dd
6098	CACAGAGUACGGGCUUACUGAAGUC	1	no match			
6099	CACACAGAGGGGAGCAGAUGAGAGCA	1	no match			
6100	UCAACCCGUACCGGAUGAUGUUCGCCUCA	1	0	coding gene		CG12225-RA|Spt6
6101	GUCUAACACGUGCGCGAGUCGGGGGC	1	no match			
6102	GAUGUCCUGGGCUGCACGCGCGC	1	0	ncRNA		rRNA
6103	CUGGGUCGGGGUUUCGUACGUAG	1	no match			
6104	AGGGUUCGAUUCCGGAGAGGGAG	1	0	ncRNA		rRNA
6105	CAGCGAUGUAUGGGGCAAGUCAA	1	0	coding gene		CG42398-RA|SteXh:CG42398
6106	GAAGCAUACGAAGAAGCAUACGAUC	1	0	transposon	LTR	invader3
6107	ACUGGGAAGGUGGGACUUGAGU	1	0	transposon	LINE	G2
6108	GCAAAUUAGAUCCGUAACUUCGGG	1	0	ncRNA		rRNA
6109	GGUUAAGCGACUAAGCGUACACGGU	1	no match			
6110	UUUAUACCUGAGCCAGUUUGUUUUGC	1	0	transposon	Low_complexity	F-element
6111	AACUGAAUAAACGAAAUGGAUGACU	1	1	transposon	LTR	invader4
6112	AGAGAACGCACCCGCUUGACCUUAUAAA	1	no match			
6113	GAUCGUUGCCCCGCAUCUCCACCA	1	no match			
6114	ACGCUGGACGUGAGGUGGUUAGGCU	1	no match			
6115	AUGGAAGUCGAAAUCCGCUAAGGAGUGU	1	0	ncRNA		rRNA
6116	AACAAUCGCUGUUAAGGCACCGAG	1	0	transposon	LINE	accord2
6117	UGUGCCCUUGACAGAAACCGAGCUGGA	1	0	coding gene		CG5405-RA|KrT95D
6118	AAUUCCGUAAACGAAGCCCCAAAGUU	1	no match			
6119	GACAGCUUUCUCUACUCUAAAUGGA	1	0	transposon	LTR	opus
6120	AGAUUAAACGAGAUUCCUACUGUCC	1	no match			
6121	CAAGCAAGCAAAUUCUAUUUUUAGA	1	0	transposon	LINE	Cr1a
6122	CGCCCAAGGAUGAGCCCGCGUAGGA	1	no match			
6123	AUCCGCACCCGUCUCCCCACAUCCGUA	1	no match			
6124	AAUGGCUCUCUCAGUCGCUUCCCGG	1	1	ncRNA		rRNA
6125	GACCGACUGGCGGACUGGGACCCU	1	0	coding gene		CG34168-RA|CG34168
6126	UCUCAUCGUCGUAGAACAAGUCCGA	1	1	Su(Ste)		Su(Ste)
6127	ACUUGAUUGCGACGAACUUAGGGAAA	1	1	transposon	LTR	HMS-Beagle
6128	AGCUGGUUCCCUCCGAAGUUUCCCUC	1	1	ncRNA		rRNA
6129	CGUACACGAAAAGUGAAACCUGCGCC	1	0	ncRNA		alphagamma-element
6130	ACCGAAUUAAUCGCGAGGCUGCAGAA	1	2	transposon	LTR	mdg3
6131	AACAUGCAGAUCGUAUGGUCUUGUACCGA	1	0	ncRNA		rRNA
6132	AUGGAACACAUAGGGGCUUCGAGGA	1	0	transposon	LTR	HMS-Beagle
6133	AUGACGGUACCCGGAGAAGAAGCCCCG	1	no match			
6134	UUGGUUAGGACUUGGGCGAUUUAAG	1	0	Su(Ste)		Su(Ste)
6135	CGCCGAAUCCCGGGGCCGAGGGAGC	1	no match			
6136	GGCAACCCCAGAUGUCCGUGAUC	1	0	transposon	LTR	ninja-Dsim-like
6137	GUGGCCUGAUUGCUAUGCACCGAAA	1	0	repeat		trf
6138	CUGACAUUUAUACAGUGCAUUGGACA	1	1	transposon	LINE	baggins
6139	CAAGAUAUGACCAAAAGCAAAGACAC	1	2	transposon	LINE	F

6140	AGAGCUAUCCGAGUCCUGAGAGUCUG	1	0	coding gene		CG31287-RA|CG31287
6141	UCCAACUGUGAGGGGAUCCUCAA	1	no match			
6142	CACAUGAACUUGGAAUUCCCAGUA	1	0	ncRNA		rRNA
6143	GGAUAAAAGGUACUCCGGGGAUAAC	1	no match			
6144	CGGAGGAGUUCGAAGGUACGCUAGUU	1	no match			
6145	UGCUUGGGCUACAUAUGGUUGAGGGU	1	1	ncRNA		rRNA
6146	ACUACUGUCAAAACACGCAACGGGU	1	0	transposon	LTR	invader2
6147	CCAUGACCUAAACACGAUUCUAGAUG	1	0	transposon	LTR	Idefix
6148	UGUUUCAUCGUUAGACGGCUCGGGUU	1	2	AT-chX		AT-chX-B
6149	CCGAGGAGAUCGCCCUGGCCACCGU	1	0	coding gene		CG6058-RB|Ald
6150	CUCCCCGCGGAAAGACGGAAAGACCC	1	no match			
6151	GGCGGCGUCCGGUGAGCUCUCGCUGGCCC	1	no match			
6152	AGAGACAGGAUCAUAUGCCGCCGCCU	1	0	transposon	LTR	springer
6153	GACGUCGGCGCGAGCCCUGGGAGGAGU	1	no match			
6154	CCGAUAACGAACGAGACUCAAAAUA	1	no match			
6155	UGAAGUAUUAGAGGGUGACAGGUAU	1	0	transposon	LTR	micropia
6156	ACCACCCUGAACGCGCCCGAUUUCGU	1	no match			
6157	CAACCAAAGAGGCAUUGCAUUAGCGA	1	0	transposon	Simple_repeat	I
6158	CGACGUACCCGGCCAGAUCAUCGAC	1	0	coding gene		CG11785-RA|bai
6159	UUUCAAGGUUACCCCCAGGAUGCUGUGC	1	no match			
6160	AUGUACAAUGCUGCCGGGUUUGCCAA	1	1	transposon	LINE	G2
6161	AUUCCGAAAAUGCGAGUUCCUCAA	1	0	transposon	LTR	Rt1c
6162	AACCCUUUCAUGCUUGGGAUU	1	2	ncRNA		rRNA
6163	UCCGAUCAGACGAGAUGGCCAACAU	1	1	transposon	LINE	Rt1b
6164	CAGAUCAACAAAGUACGAUGGGGAC	1	1	no annotation		
6165	ACCGAAUACGAAGAACAGGCAGA	1	0	transposon	LTR	gypsy12
6166	AACUGAUAAACGAAAUGGAUGACA	1	no match			
6167	GAUCCAACUGUGAGGGGAUCCUCAA	1	no match			
6168	GAGGCUGAGAUUUGCUGCAGUUCUGA	1	0	coding gene		CG8318-RB|Nf1
6169	CUCGAAUGUAUGCCCGAUAUACGGAGCGU	1	0	transposon	LINE	R1-element
6170	AGCGCGAAGAAGAUGACUGGCCC	1	no match			
6171	AACGGAGUCUAACACGUGCGCGAGU	1	no match			
6172	ACGGUACUGAUCAACUAAAGGAUGCGC	1	0	no annotation		
6173	GGACAAAACAACAUACCGGGAAAUGAG	1	1	transposon	LINE	baggins
6174	CGUAACUCAAACGUCGAGCAUGAAA	1	1	transposon	LINE	R1
6175	GCAUUCGAGACCCAACGGCACGU	1	0	transposon	LINE	INE-1
6176	GACCUACGAAAUGAGCUGUCUGAGU	1	1	coding gene		CG40084-RA|CG40084
6177	GGAGUUAAAAAAGCACGUCAUCCC	1	1	Su(Ste)		Su(Ste)
6178	AACCAAUCCCAACCAAACCCGUCAAAC	1	0	transposon	LINE	G2
6179	CUGUCAGGUACCCCGUUCAGACCUAG	1	0	transposon	LINE	Rt1b
6180	UGCACCAAAAUGGACUUCAACUUGCAGU	1	0	coding gene		CG3517-RA|CG3517
6181	GACAACGCGAGAGUCAGAGUGACCAA	1	0	transposon	LTR	Idefix
6182	ACUUCAUAACCGAGCGACUGUUCCAU	1	2	transposon	LTR	GATE
6183	CCGGCGAAAAUUCGAGAGUCGCAAG	1	0	transposon	Unknown	RR50960
6184	UGGGAAUGGCUUCACAGUUCUGC	1	0	ncRNA		alphagamma-element
6185	GAUAUUCUUAACCCGACACAUAUG	1	1	no annotation		
6186	ACCAGCGGGAUACCUCAUACUCGAGUA	1	1	transposon	LTR	micropia
6187	UAACGGAGGUGUCCCAAGGCCAGC	1	0	ncRNA		rRNA
6188	CUAAGCCCCGAACUGUGACACUGGC	1	0	coding gene		CG31093-RA|CG31093
6189	AAUAGAAGCAUUUGGGGCAUUAGU	1	no match			
6190	UAACCGUCAUUCCAGCGCAAGUUGCCCCC	1	2	transposon	LTR	ninja-Dsim-like
6191	GCGGAGUUAAAAGCGUUUACGUUC	1	0	transposon	LTR	micropia
6192	UCCCCGGAACCGAGAGUCUCCUGCCA	1	0	coding gene		CG34241-RA|CG34241
6193	CCCCGGAUAGUUUAGUUACGUAGC	1	0	ncRNA		rRNA
6194	AGCCUGUGGGAAUACUGCCAGCU	1	no match			
6195	ACCCUGAGGCCAUCGAUGAUGUGAGA	1	no match			
6196	ACCACCCUAAACGCGCCCGAUCUCGUC	1	no match			
6197	UUGGAGCUUUACGGAACUGGUUGAC	1	0	no annotation		
6198	CAGGCAAAUACUUGGAGUUGUGGC	1	0	transposon	DNA	Bari1
6199	UCUUGUACCGACGACAGAUCUUUCAAAU	1	0	ncRNA		rRNA
6200	GACCAAUACCUCGAUUUCCUAAA	1	0	transposon	LINE	Porto1
6201	GCUCGUUCUGGGUAAAUAGAGUUUC	1	0	ncRNA		rRNA
6202	AGCGGGAUUAUUACAGAUGAUGGAUCA	1	0	transposon	DNA	H
6203	CAAGCUAGACUAUGUAUCCCCGGU	1	1	transposon	LINE	Doc
6204	ACCGUUUACACAUUUAUAUUUUUUGU	1	2	transposon	LINE	Cr1a
6205	AUGACAGACAUCCGAGAAAACAAGAGC	1	0	transposon	LTR	gypsy3
6206	UGCCUCGAACUUCCCGAGAUCGUUAC	1	0	coding gene		CG11856-RA|Nup358
6207	UCUCAUCGUCGUGGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
6208	AAGAACAACAACGCAGACGGCGCC	1	0	transposon	LTR	diver2
6209	UUCCGAUAACGAACGAGACUCUGGC	1	no match			
6210	ACCCAAAGACCGUUGAAAGCGAGGUGA	1	0	coding gene		CG6279-RA|CG6279
6211	AAACCAUCAGAGACUGCGGUGGCCAA	1	0	coding gene		CG12717-RA|CG12717
6212	UCACGAGUGAUCCAGUAAAUUACC	1	1	transposon	LINE	Quasimodo
6213	UUGACAAGAAAACGAGAGAACGCUUC	1	0	transposon	LTR	blood
6214	UAGUAAAUAAAACGAUAUUCUCGAAGU	1	0	transposon	LTR	diver2
6215	CUCUCAUCGUCGUAGAACAAGCCCG	1	1	Su(Ste)		Su(Ste)
6216	GAUACUCGAAGACACCACGGUGAA	1	0	transposon	LTR	ninja-Dsim-like
6217	GCCGAACGUAGUCUCUCGAGAGGG	1	1	transposon	LTR	invader1
6218	AGCGAAAUGACUCUGUCCCGAUUCA	1	0	transposon	LINE	R1-element
6219	AAGUCACAAGACCCCUUUGCCCGUGCCC	1	0	no annotation		
6220	GCAAGGACGAUUGAUGCUACAGUAUU	1	no match			
6221	AGAGAUUGGAUGGCCAUUAGAUG	1	1	transposon	LTR	frogger
6222	AAUUAUGACAACAUACCGGUUAAG	1	0	no annotation		

6223	GUGCUAGAAGACGAUUUUCGGGU	1	0	transposon	DNA	1360
6224	CAGCGUCCCGUGCUGAUAAAGACCCACU	1	0	coding gene		CG1710-RA|Hcf
6225	UGGCCUAUCGAUCCUUUAGUCCC	1	no match			
6226	UAGCUAAUUAACACAAUCCCCGGGGGC	1	no match			
6227	GUGGCAAUAUCCCCAAGGUGCAAGA	1	no match			
6228	CAAACUAUCAUGGAAUUGCGCCGG	1	0	transposon	LINE	Doc4
6229	GGAGAACUUUGGAGUGAAGACCGC	1	0	transposon	LTR	ninja-Dsim-like
6230	UGUUUCAUCGUUAGAUGGCUCGGGCG	1	1	AT-chX		AT-chX-B
6231	CAAACCGACACUGGUGCGCGAGAUGAGU	1	no match			
6232	AGCAGUGCCCGGGAUGUCCAGCGACU	1	0	coding gene		CG10561-RA|CG10561
6233	AGCUUCAUCCAAGCGAACUGUGCCG	1	0	transposon	LINE	R1-element
6234	UGGGAACACCGCGUGUUUGUUGGCCU	1	no match			
6235	UCACCGAGGCGGCACUCGCAAUCUGCAA	1	1	transposon	LTR	opus
6236	GUAACAACGUCUACAUCCCGCUACC	1	0	coding gene		CG13591-RA|Ssl
6237	CGUCGCCUACGAGCCGCACCCCGCU	1	0	coding gene		CG11770-RA|lin
6238	GAUGCCGGAGGACGUAAUGGUGCCCG	1	0	coding gene		CG11703-RA|CG11703
6239	CAACUGCCCGAGGUUGACCACGAUGA	1	0	coding gene		CG1365-RA|CecA1
6240	UUUCCCGAAAUCCGAGUGAAUAUGGCA	1	0	no annotation		
6241	GAAUCCUCAACUGUGAGGGAUCCUCAA	1	no match			
6242	ACAGAUUCCGCAGUGCGUUGAUA	1	0	Su(Ste)		Su(Ste)
6243	UGUUUGUAGUUGUACUGACCCAGUCGG	1	1	transposon	LINE	baggins
6244	AACUGAAUAAACGAAAUGGGUGAC	1	1	transposon	LTR	invader4
6245	GAGAAACUUAUCGACUAAUCGACUCGC	1	0	transposon	Unknown	RR50960
6246	UAUCGACGAAUGAUCCAGGCUUGUUU	1	0	no annotation		
6247	UUCAACCGGACGGGCCCUAGAGUUAUUC	1	no match			
6248	ACCGAAUUAAUCGCGAGGCUGCAAGUA	1	2	transposon	LTR	mdg3
6249	UACGAUUGAAAAAGAACUACUAG	1	0	transposon	LINE	McClintock
6250	CAGGUCAUUUCUGGGACUGGCCAGU	1	0	transposon	LTR	FB
6251	CGCAUUGAUUACGUCCCUGCCCUUUGUA	1	0	ncRNA		rRNA
6252	GCCCUAGUUCUAACCAUAAACGAUGCA	1	1	ncRNA		rRNA
6253	GCCCGUAUAACGUUAAUGAUUACUAGAU	1	0	ncRNA		rRNA
6254	UAAUUCCGAUAACGAACGAGACUCAAAU	1	0	ncRNA		rRNA
6255	UGUGAAAAGAUCCUGCUUGAAUAUGACC	1	0	transposon	DNA	S
6256	GCUCAAGAUAGGCAGAUGUGCCCGAU	1	1	transposon	LINE	Cr1a
6257	UGUUACAAUGAUCUGAUCGGGUU	1	0	transposon	LINE	Cr1a
6258	GCCCGAUACAUCAGGUCAGAGCGUGGCC	1	0	repeat		trf
6259	UUCCGUAGUGUAGCGGUAUCACGUGUGC	1	no match			
6260	AAGAAAAACAUGAAGCGCCGUUGU	1	0	transposon	LINE	TART-A
6261	AGCUAGAUAAAAGCUUUUGGGGCGA	1	0	transposon	LINE	Cr1a
6262	CAAGAUAUGACUAAGCAAAGACAC	1	no match			
6263	UGAUACGGAGAAUGACAAUGGUUGA	1	1	transposon	LTR	gypsy3
6264	GUACACAGAGGAGAGCAGAUGAGAGCA	1	no match			
6265	UUCGCAAAAGUUGACCGAACUUGAUU	1	0	ncRNA		rRNA
6266	AUGGCACUCUCAGUCGCUUCCCGGGA	1	1	ncRNA		rRNA
6267	CCCGGGGCGUUCUAUAUAGUUAUGUA	1	0	ncRNA		rRNA
6268	GAGGUAACGAUGGAGUUUGACGGCA	1	0	transposon	LTR	flea
6269	AGAACUGGCGAUGCGGGAUGAACA	1	no match			
6270	AGAACCUGAAACCGUAUGCGUACAAGC	1	no match			
6271	CCGAGCUGUGACCAUCGAGCUAGGAAA	1	0	transposon	LINE	R1-element
6272	AUGGCAUAAAAUAGAUAACCUGCUGAA	1	no match			
6273	CAAGGUCUCUGGAGGAAUUCAAU	1	no match			
6274	GGCAGUCAGAGGCGAUGAAAGACGUGA	1	no match			
6275	CACCUUGGGAAUGGCUUCACAGUU	1	0	ncRNA		alphagamma-element
6276	AGGGUCGAAACGAUCUUAACCUAUUCU	1	0	ncRNA		rRNA
6277	GCGUGAGGAAGGAGGGAUGAUGU	1	no match			
6278	CCUGGGAAACGGGGCGCGGCCGGA	1	no match			
6279	AAUGGAAGGAGGCCUUGAUAAACGAG	1	0	no annotation		
6280	AAAAAAGGAAUUGACGGGGGGCCCGC	1	no match			
6281	CGGAAGCUUCCUUGGCCCAUAUG	1	0	transposon	LTR	ninja-Dsim-like
6282	UUGAAGCAUAUUCCCGAUUCGGAAAAGG	1	0	no annotation		
6283	UACAAUGAUCUGAUCGGGUUUUUCUG	1	0	transposon	LINE	Cr1a
6284	AGGGACAUUAUUGUGGAAAAUGUGA	1	0	no annotation		
6285	AAACUGAUAGACAGAUGGACGGA	1	0	Su(Ste)		Su(Ste)
6286	GCCACCGAGAUAUGAGGACGACGACG	1	0	transposon	LTR	ninja-Dsim-like
6287	AGCUUCAUCCAAGCGAAACUGUCGCCCGA	1	no match			
6288	AUCAAUCGCUGAUAAGGCACCGGGAUAU	1	2	transposon	LINE	accord2
6289	GUCGAGAGAUAUCGGAGGAGUUU	1	1	ncRNA		rRNA
6290	GCGAAGAUGUAACGGGGCUCAAGACAC	1	no match			
6291	CACACAGUGAAGACGAGACUGCAG	1	0	transposon	LTR	frogger
6292	GCCCAGGUGGCGGAAUUGGUAGACGCACU	1	no match			
6293	ACUCAUAUCCGAGGCCCUGUAAU	1	0	ncRNA		rRNA
6294	UAUUGUAUAAAACUUCUUCGUCACAU	1	1	transposon	DNA	hopper
6295	ACCCUGAACGCGCCCGAUCUCGUCUU	1	no match			
6296	AACCCGGAUUGAAGCAACCAGAGAA	1	0	coding gene		CG4704-RA|CG4704
6297	CUGCCGCGUGCCCACCGACUACGUG	1	0	repeat		trf
6298	CAUGCUUGGGAUUGUGAACUGAAACU	1	0	ncRNA		rRNA
6299	GAGACCCGUCGCCGCGCUCUCCCCC	1	no match			
6300	UACCGAUGGCGAAGGCAGCCUUCUGGC	1	no match			
6301	ACUCAUAUCCGAGGCCCUGUAAUUG	1	0	ncRNA		rRNA
6302	CCGGGGGAAGUAUGGUUGCAAAGC	1	0	ncRNA		rRNA
6303	CGCUCUCGAAUGUAUGCCCGAUAU	1	1	transposon	LINE	R1-element
6304	GCACACACAGGAGAGCAGAUGAGAGCA	1	no match			
6305	CUCGCCGAAUCCCGGGGCCGAGGGUGC	1	no match			

6306	ACGAGCGCAACCCUUGUCAUUAGUUGC	1	no match			
6307	GGCCGGCGGGGAAGGUGGCUCGGG	1	no match			
6308	ACCCUGCACGCGCCCGAUCUCGU	1	no match			
6309	AAGCCCGAUGAACCUGAAUAUCCGUUAUG	1	0	ncRNA		rRNA
6310	AGCAACUGAGCCCAAGAAAAUCGCC	1	1	transposon	LINE	Doc
6311	UACAUUGCUAUGUUGACGCGGAUUG	1	0	transposon	LTR	DM88
6312	UGAGAAGUUAUGGUUGGCGAAGUCG	1	0	transposon	LTR	Idefix
6313	GAGGUCUCCGGACGUGAUCACUGUGUGAC	1	no match			
6314	AAGGCAUGCCCGAUGUCCGCGAGA	1	0	transposon	LTR	opus
6315	GUAAAGAUGACCCGUGAUUCCACCGGGGA	1	0	coding gene		CG33322-RA|CG33322
6316	CGGACCGAAAAUCCUUUACUGGA	1	0	transposon	LTR	DM88
6317	CUGAAUAAACGAAAUGGAUGACA	1	0	transposon	LTR	invader4
6318	CCCGCGGUCAGACGGAAAGACCCUA	1	no match			
6319	AGCUUCAUCCAAGCGAACUGUUGCCCGA	1	no match			
6320	CUGAAGGACUUGCUUGCGUUGCAGGA	1	no match			
6321	ACCAGACGUAAUACUCCACUACCCGA	1	2	transposon	LTR	HMS-Beagle
6322	ACCGAAUUAAUCGCGAGGCUGCAAGAU	1	1	transposon	LTR	mdg3
6323	CAUGGAGCUAACCGAUACUAAUUG	1	no match			
6324	ACCAGACGUAAUACUCCACUACCCG	1	1	transposon	LTR	HMS-Beagle
6325	UCCUAGAAUCGACCGACCAGCUGUGCG	1	0	transposon	LTR	invader3
6326	GAUCCACAUCAAUAGUCCCAGCGCC	1	0	coding gene		CG5016-RA|Mst57Db
6327	UGCUCUCGAAUGUAUGCCCGAUUU	1	1	transposon	LINE	R1-element
6328	GGAGUCUAACAUGUGCGCGAGUC	1	no match			
6329	GAUGAUUUAAUUCUAGCGUCCCAGU	1	0	transposon	LTR	1731
6330	UGAUUUGUCUGGUUAAUUCCGAUAACG	1	0	ncRNA		rRNA
6331	AAAACCAUUCGUAGACGACCUGCUUC	1	no match			
6332	UCCGGUUCCGUCACGUAGAGUCAG	1	0	transposon	LINE	Rt1b
6333	GGGGUGCGAGAGGUCCCGGGUUC	1	1	ncRNA		tRNA
6334	CGGAGCCGGACUACCUUGACCUCUU	1	1	transposon	LTR	diver2
6335	ACGGAAUGGAACCGAGCUCUGAGUAAU	1	2	transposon	LINE	X
6336	CAAGACAAUAUAUUGCGACAACGAA	1	0	transposon	LTR	gtwin
6337	UUCACCGGACGUCGAGGACAAGUUCA	1	0	transposon	LINE	INE-1
6338	CCGAACUUGAGACCGCCCCUGGACGUACC	1	2	Su(Ste)		Su(Ste)
6339	UCGCCGAAUCCCGGGGCCGAGGG	1	no match			
6340	UGGAGCGUUCUGCCUUUGGGGUUGA	1	0	transposon	LINE	G2
6341	GUGCUCGUUCUGGGUAAAUAGAGUUUC	1	0	ncRNA		rRNA
6342	CGAGGAGACCUGUUAUACAUCUGCGA	1	0	transposon	LTR	ninja-Dsim-like
6343	GACGCGCAUGAAUGGAUUAACGAG	1	0	ncRNA		rRNA
6344	GUGUAACAACUCACCUGCCCGAAAGCAA	1	no match			
6345	GAGCUUGACUCUAGUCUGGCACGGU	1	no match			
6346	UACUGACGAAUUUAGUUACAAGCA	1	0	no annotation		
6347	GCAAAGAGAACGCGACCAACUGAAUGGC	1	0	no annotation		
6348	ACGAUUUAAGUCCUUCUUAAAUGAGGCC	1	0	ncRNA		rRNA
6349	AAGACCCCCGAAACCAGACGAGCUACCU	1	no match			
6350	AAUCGAAUGAGACCUUGGAGGA	1	0	ncRNA		rRNA
6351	UUGAAAAUUUAUGGUGGGGACACGCA	1	0	ncRNA		rRNA
6352	UAGACAGAGAUAUUACCAGAAGCCAC	1	0	transposon	LTR	gypsy4
6353	ACGGAUGGAAUCGAGCGCUGAGUA	1	no match			
6354	UCCAAUACCACCGCGGAGACUGUCGC	1	1	transposon	LINE	Doc4
6355	ACCCUUUUCACAUUUUCGCCAACCGG	1	0	coding gene		CG41521-RA|CG41521
6356	CGAGAUGACAACCUUGGACCUGAC	1	0	transposon	LTR	micropia
6357	AUUGGACCAUGUGCUCCUGGAGGACCC	1	0	coding gene		CG11719-RA|Mst98Ca
6358	UCAAGUCAGAGCGUGGCGUGAUUGCUA	1	0	coding gene		CG42398-RA|SteXh:CG42398
6359	AAACAGGAUUAGAUACCCUGGUAGUC	1	no match			
6360	GCAACAUGUCGAGACCGAAGUAAC	1	0	Su(Ste)		Su(Ste)
6361	UUGGACACGAGCGACGGCGAGCAGACC	1	1	coding gene		CG9715-RA|CG9715
6362	UCAGAACUGGCACGGACUUGGGGAAU	1	0	ncRNA		rRNA
6363	CAGACCGUGAGCGGGUGCAGCCGAU	1	0	transposon	Low_complexity	F-element
6364	UACAUUUGUACCAACUCCGAGAUAUAC	1	0	coding gene		CG11714-RA|CG11714
6365	GUUGCAAUUAAAACAGAUUCCGCAGUGC	1	no match			
6366	UGCUAACGAAACAGUAACACUAGGC	1	0	transposon	LTR	Idefix
6367	AACUACUUUCAGGACACGUUCAACCAGA	1	0	Su(Ste)		Su(Ste)
6368	ACCGCUUGGACCGAACACUUUGAAA	1	no match			
6369	UACGGGCUCGACAUGUUGCCAGUUCAC	1	0	Su(Ste)		Su(Ste)
6370	CCCUUAGAUGUCCUGGGCUGCACGC	1	0	ncRNA		rRNA
6371	CGACUCGUUUUUAAAUUUCGCAACAA	1	0	transposon	LINE	Cr1a
6372	ACCACCCUGAACGCGCCCCGAUCUCGUCC	1	no match			
6373	CCCUAAACGAAAAGAGUGAUCCACAAA	1	0	transposon	LINE	S
6374	GUGGAGCCGGGCGUGUAAUGCGAGUG	1	no match			
6375	AGCUCUCGAAUGUAUGCCCGAUAUA	1	1	transposon	LINE	R1-element
6376	AACCCGCACUCUUAGACCGACAAGA	1	0	no annotation		
6377	UGCUCUCGAAUGUAUGCCCGAUAUUAC	1	no match			
6378	GAUGCAAACUCCGAAUACCAGCAAGU	1	no match			
6379	ACAAGAGAAAAUAUAACGACCGGUGA	1	1	transposon	LTR	McClintock
6380	ACGGAGAAGGUUAGCUCAGCCGGGU	1	no match			
6381	UUAAAGGAAUUGACGGAAGGGCACCACC	1	0	ncRNA		rRNA
6382	AACGAGAACUUUGAAGGCCGAAGU	1	no match			
6383	GGAUGGCGAUGCCGGACUGAUCACGG	1	0	coding gene		CG15657-RA|CG15657
6384	UUGGGCCCGAACCCUUCUGGUGUGUUG	1	0	transposon	LINE	baggins
6385	CCGAGGUUGAUUGAAUUAGGACUC	1	0	no annotation		
6386	ACAAACUCAAGUCCCACCUUCCCA	1	0	transposon	LINE	G2
6387	UGCCAGCCCCGUAUAACGUUAAUGAU	1	1	ncRNA		rRNA
6388	GCCUGAGAAACGGCUACCACAUCUAA	1	0	ncRNA		rRNA

6389	CCCAUGAGCGAGUCAUUCCGACAAC	1	1	transposon	LTR	blood
6390	AGAUACCGCCCUAGUUCUAACCAUA	1	0	ncRNA		rRNA
6391	CCAAUUCAAGGACAUUGCCAGGUAGGGA	1	0	ncRNA		rRNA
6392	UGAGACUACAAACGGACUGCGGAC	1	1	no annotation		
6393	AAGAGGGACGGUCGGGGGCAUUCG	1	no match			
6394	GUGCACAUAUACCGAGGCCAACGAUCG	1	no match			
6395	CAGGCAUUUAAGUUGUAUUGUGAG	1	0	transposon	LTR	diver2
6396	GAUAGGCCGAUAGCGUACAUUUCC	1	0	transposon	LTR	opus
6397	GCUGAAGAAGGUCCCAAGGGUUCGG	1	no match			
6398	AGGUUGCGUGUUCAUCACGUCGGGG	1	no match			
6399	UGCUCUCGAAUGUAUGCCCGAUAUUUA	1	no match			
6400	UGGAAUCGGGUUUGCAGAUUGCCCU	1	0	transposon	LTR	gypsy12
6401	CUCCGGACAAAAGUUGACCGAACCUGA	1	no match			
6402	UUCCUCGGAAUCAAGGGCAACGAGU	1	0	Su(Ste)		Su(Ste)
6403	UCCUAGUAUUCUUCCGGAUGAUCCACC	1	0	transposon	DNA	S
6404	CGCGAAAGACGGAAAGACCCCAUG	1	no match			
6405	CAGCGAUGUGUGGGGCUAGUCAAA	1	1	Su(Ste)		Su(Ste)
6406	ACGGGGGGCUGGCGGUGCGGCGAC	1	no match			
6407	AAGGGUCGGGUAGUGAGGGCCUU	1	no match			
6408	CACAAACGCACCAAGCAUGUUGACA	1	0	no annotation		
6409	UUCGAUUUCAAGUACCGGGCGAAC	1	0	repeat		RSP
6410	UCGGCCCGAACCCUUCUGGUGUGUUAU	1	2	transposon	LINE	baggins
6411	UGUGCUAGAAGACGAUUUUCGGGAC	1	0	transposon	LTR	HMS-Beagle
6412	UGAGUGUAAAGACGGAGUGUGCCCCAA	1	1	transposon	LTR	ninja-Dsim-like
6413	CUUCGGGAUAAGGAUUGGCUCUUAAG	1	1	ncRNA		rRNA
6414	ACCGUAUUAGAUCAGUUGACUCUCCUCC	1	0	no annotation		
6415	AGCACUUGGGACGCCGUUUACCC	1	0	coding gene		CG12717-RA|CG12717
6416	AAUUCCGUAUAACGAACGUAGACUCUGGC	1	no match			
6417	GUUUGUGGCUUCAGACGGCGAGUGGC	1	0	coding gene		CG10596-RA|Msr-110
6418	AGCUUCAUCCAAGCGAACUGUGGCUGA	1	1	transposon	LINE	R1
6419	GCCAAUUACAUUAGACGAGACGGCCCC	1	1	transposon	LTR	opus
6420	ACGAGUAAUAUGAGCGCCCAGAGUGG	1	1	transposon	LTR	opus
6421	AUAAGUAGAACACUUUUUGUUUUUUGG	1	0	no annotation		
6422	UCUAGGACUACCAUGGUUGCAACGGGU	1	0	ncRNA		rRNA
6423	GUGGCAUGUUUCCGGAUUCCCGAC	1	1	ncRNA		alphagamma-element
6424	UGAACAUCGAGAAUACUGCACAGCC	1	0	transposon	LINE	Doc
6425	UCGGAAUGCCGGGGCGAGGAGGAGG	1	0	Su(Ste)		Su(Ste)
6426	UCACCAAGCGUUGGAUUGUUCACCCAC	1	1	ncRNA		rRNA
6427	GCGGGACAUGUGGCGUACGGAAGACCC	1	no match			
6428	UGAGCGUGAAAUCGUCCGUGACAUCA	1	0	coding gene		CG4027-RA|Act5C
6429	GAUUCAUACCUGAGCACUUUAACA	1	0	transposon	LTR	gypsy8
6430	GACCAUAUCCGAAGACGCGGCGCU	1	0	transposon	LINE	Rt1b
6431	UAAGUUCUCGGCUGAUCAGUACCCA	1	0	transposon	LTR	Quasimodo
6432	GACUGCAUCGCUUGAAUCCAGGAGA	1	no match			
6433	AGCCGUACCUGUUGGUUUGUCCCAUUA	1	0	ncRNA		rRNA
6434	GCAUCGAGAAUAUUACACAGCCAUUA	1	1	transposon	LINE	Doc
6435	GAAGGAAUUGACGGAAGGGCACCA	1	1	ncRNA		rRNA
6436	ACAACCUCAACUCAUAUGGGACUACCCC	1	0	ncRNA		rRNA
6437	AGAUGUCCGGGGCUGCGACGCGCGCUAC	1	no match			
6438	UUGAUACCUUGGCCAGAUUGUUCUA	1	1	transposon	LINE	Doc
6439	AACCGGAGGAAGGUGGGGAUGA	1	no match			
6440	UGCCCGUAAAAGCGACUUGGUUGACAC	1	0	transposon	LTR	gypsy
6441	UUAUGCCUGAGCGCCUCUAAGGUCGU	1	1	ncRNA		rRNA
6442	GAACUGCUCAUUCCGAACAUACACCGA	1	0	transposon	LINE	Doc
6443	UAACGAGAGAAGCAUUUUUGCCGCAAA	1	0	transposon	LTR	gypsy12
6444	AGCGGCGAGCGAAGUGGGACCAGCCC	1	no match			
6445	GCAUCGGUCGUUCAGUGGUAGAAUGCA	1	2	ncRNA		tRNA
6446	AAUGAAGACAAUACUUCUCUAGGGGA	1	1	repeat		trf
6447	UCAUCUUCUUCUCGACGUUGACCA	1	0	transposon	LTR	ninja-Dsim-like
6448	AACUACUGGACCAGAUGAUACGAGGGC	1	no match			
6449	CCGAACUUGAGACCGCCCCUGGAAA	1	1	Su(Ste)		Su(Ste)
6450	ACGAGAAGCCGCCGUAUAGCUAUGCC	1	0	coding gene		CG11799-RA|Mnf
6451	UUCCAAAUCGAGAGACGACCGCCGA	1	1	transposon	LTR	flea
6452	AAGACAAACAACGCACUAGACUCCGGA	1	no match			
6453	GCUGCUCGAACCGGAGGCACAUGCUCGC	1	0	coding gene		CG14305-RA|CG14305
6454	ACCCUUGAUAAACGGUUGACUUUCAGU	1	1	transposon	LINE	BS
6455	ACCUGGAUUGAUAAAACAAAUGAAC	1	0	transposon	LTR	HMS-Beagle
6456	UGCUUUCUCUGAAGCUAUCGAGUGAGA	1	0	transposon	LINE	Cr1a
6457	AAAACCAAUCCCAACCAAACCCGUCAA	1	1	transposon	LINE	G2
6458	UAUUUCAUCGUUAGACGGCUCGGGC	1	1	AT-chX		AT-chX-B
6459	GUGUAUCAUUUCCUAGCCAUUAUACG	1	0	repeat		LSU-rRNA_Dme
6460	AGGUUCCGGGGGGAGUAUGGUUGC	1	no match			
6461	UCCGAAGAUGACCAAGUUGGCUACCAA	1	1	transposon	LTR	gypsy6
6462	CAGAUACUGAAACGAGUUGGCAUU	1	0	transposon	DNA	RR51476
6463	AGUUUAUCCGGUAAAGCGAAUGAUU	1	no match			
6464	UGGGGCUCCGGGGGAAGUAUGGUUGCAA	1	1	ncRNA		rRNA
6465	GAUCACCUGAUGCCGCGCUAGUUU	1	1	ncRNA		rRNA
6466	AGAUACCGCCCUAGUUCUAACCAU	1	0	ncRNA		rRNA
6467	CGGGAUUCCAUGGGUAGACGACGAAA	1	0	transposon	LINE	G2
6468	UCUGGGUCGGGGUUUCGUACGUAGC	1	no match			
6469	AAGGAGUGUGUAACAACUCACCUG	1	0	ncRNA		rRNA
6470	CCACAGAGUACGGGGCUUACUAAAAA	1	2	transposon	LINE	baggins
6471	CCGUUAGAGUACAUUAGGUAAGUGAGC	1	0	transposon	LINE	R1-element

6472	UGGGAGGAUCUUCGAUCACCUGAUGCCG	1	0	ncRNA		rRNA
6473	GUGGAUAACCGGGAUCGACACGGCCUGCGC	1	no match			
6474	ACCCCAGGGACAAGACAGCGCUCAC	1	1	transposon	LINE	G
6475	ACUUGUGCUUCAUACGGGUAGUACAAC	1	0	ncRNA		rRNA
6476	CGUAUACUGAAGGAAGAGAGUUC	1	0	transposon	LINE	G2
6477	GCUCCUGAUGAAGAAGAGAUGCCC	1	0	transposon	LTR	ninja-Dsim-like
6478	AAGGAACGAUCGCAAGGCAGCAGU	1	0	transposon	LTR	flea
6479	ACCACCCAGAGAAGUACCAAAAAAGC	1	no match			
6480	AGAGGGCGUGGAACCGUUAAGAGG	1	no match			
6481	ACGAGCGCAACCCUUGUCAUUAGU	1	no match			
6482	CCAUACUGUACUCUAUUUUCUCCUCAGA	1	1	transposon	LINE	TART-A
6483	GCAAUACUAAUUCGAUCCAGAUGCC	1	1	transposon	LINE	Doc3
6484	UCCGAAGUGACCGACCGAAACAGUCGC	1	0	transposon	LTR	diver2
6485	UGCCCUCGAAUGUAUGCCCGAUAUU	1	2	transposon	LINE	R1-element
6486	UCUCAUCGUCGUACGAACAAGCCCGAGGU	1	no match			
6487	UACGAGAGGAACCGCAGGUACGGACCA	1	0	ncRNA		rRNA
6488	GGAUACUAAGCAGGGACGUUGAUGUA	1	1	coding gene		CG12717-RA|CG12717
6489	AUUCCCCAUGAACGAGAAUUCCCAGU	1	no match			
6490	GCAACAGAUACAAAAACGAGCGGACC	1	no match			
6491	UGGAAAACCUGAAGAUGAUCCUGAG	1	0	transposon	LINE	Helena
6492	UGUUUCAUCGUUAGAGGCUCGGGC	1	no match			
6493	AAUAGGGACGGUCGGGGCGAUC	1	no match			
6494	CGCACUUUAAGCGUUCGUUGACGU	1	0	transposon	LTR	HMS-Beagle
6495	UCGGCCGAAAAUACGACGACCCCAAG	1	0	coding gene		CG18743-RA|Hsp70Ab
6496	CAGAACUUGAAGUCGCCGAGGCAGCCAG	1	1	transposon	LINE	F
6497	GGACCAACUGUGAGGGGAUCCUCAA	1	no match			
6498	GUAGCAAUUACCGACCGGUUUACACGC	1	0	transposon	LTR	G4
6499	AAAUAUACUAUAACCACCAGUUGGGA	1	0	transposon	DNA	S
6500	CGAAAGCAUAGGGACGGACUGCAAG	1	no match			
6501	CCAUCCUAAAACUGAUAGACAGCUGGA	1	1	Su(Ste)		Su(Ste)
6502	UUCGAGGUUACCCCAGGAUGCUGU	1	1	repeat		trf
6503	GCCCCGUACGUGAAAUGACCUGUGUG	1	no match			
6504	GGUCCGGAGAGAUGGGCGAGCGCCGUU	1	no match			
6505	GCCCGGAUAGCUCGGUCUGUAGAGCAUUG	1	2	ncRNA		tRNA
6506	UCGCCGAAUCCCGGGGCCUAGGGAGCGA	1	no match			
6507	ACUCGAAUCCCGGGGCCGAGGGA	1	no match			
6508	UCGAAGGUCUUCGUCCCGGUUUCCA	1	0	transposon	LTR	GATE
6509	UCGGCCCGAACCCUUCUGGUGUUUAAU	1	no match			
6510	AGCCAAUUACAUUAGACGAGACGGCCC	1	1	transposon	LTR	opus
6511	AACUUCAUCCAAGCGAACUGUGCCCGA	1	1	transposon	LINE	R1-element
6512	GAAACGCACGAUGGUGUCCAGGAUAUG	1	0	coding gene		CG13387-RA|emb
6513	AUAAUACACAUCGAUCGAUUACGGACCUA	1	no match			
6514	ACAUUCAUAACCGAGCGACUGUUCC	1	0	transposon	LTR	GATE
6515	GUGAACCUGCGGAAGGAUCAUUA	1	0	ncRNA		rRNA
6516	UCGCCGAAUCCCGGGGCCGAUGGAGC	1	no match			
6517	CAAGCUAGACUAUUUAUCCCCCGGGU	1	no match			
6518	CUUGUAGUCCACCUGCACCUUGGGC	1	2	no annotation		
6519	ACAUUCAUAACCGAGCGACUUUCUAA	1	no match			
6520	CAAAAUGGCCGAGAGCUUGUGCAAUGA	1	0	coding gene		CG8580-RA|akirin
6521	AUCUUCAUCCAAGCGAACUGUGCCCGA	1	1	transposon	LINE	R1-element
6522	GGACCCCCUCUUCCUGACCACUGCC	1	no match			
6523	GAAGUGGAGAAGGGUUUCGUGUGAACA	1	0	ncRNA		rRNA
6524	AAAAGGAGCAAUUUUUGGCCGUCAC	1	no match			
6525	UGUUUCAUCGUUAGACGGCCCGGGC	1	1	AT-chX		AT-chX-B
6526	UCAAAUAGAAAUGGACUCAGAGUC	1	0	transposon	LTR	rover
6527	UGCCUGAAGGACGUAUGCCUGUCGCCC	1	0	coding gene		CG1193-RA|CG1193
6528	CCAAGCUUUUGGGCUCCGGGGGA	1	1	ncRNA		rRNA
6529	CGAGAUAGUACGAAUCGGUUACGAA	1	0	transposon	LINE	I
6530	CCCGAAGAUAACUCGCAGUCGCUACA	1	0	transposon	LTR	GATE
6531	GUACUUGUUCCCCGGAUAGUUUAGUUAC	1	1	ncRNA		rRNA
6532	GUGGGAUAGCGUACCGGAGUGGAC	1	0	ncRNA		7SLRNA-RA
6533	AAAUCUCGCGCCGGGCCGUACCCAUAUC	1	no match			
6534	UGCUCUCGAAUGUAUGCCCAAUA	1	0	transposon	LINE	R1-element
6535	ACCCCAGAACGCGCCCGAUCUCGU	1	no match			
6536	GAUUCCGCAGUGCGUUGAUAUGGUC	1	1	Su(Ste)		Su(Ste)
6537	CAACAAAGCACCUGGACAUGACG	1	0	transposon	Simple_repeat	I
6538	GCCCGUGGACGGUGUGAGGCCGGUAGC	1	no match			
6539	AACCACUUUCAUGCUUGGGAUUGUGAA	1	0	ncRNA		rRNA
6540	UGAUCCUCAA	1	0	transposon	LTR	diver2
6541	CUACACAAAAAGUUGACCAUAAAAA	1	0	transposon	LINE	I
6542	AGAUGCCGCGACGUAAGCCCACCAUUAA	1	0	coding gene		CG10750-RA|CG10750
6543	UCUCAUCGUCCUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
6544	GGAAAACGACAACGACGGAAGAGGA	1	1	coding gene		CG41053-RA|CG41053
6545	CAUCUUGUUAUUCUAGUGUCUUUGC	1	0	transposon	LINE	Cr1a
6546	GAAUCUAUGGGUGGUGGUGCAUGGCCGU	1	0	ncRNA		rRNA
6547	CGCAAAAUUAUACCCGUACCCAAUGG	1	0	transposon	LTR	gypsy6
6548	CCGAAUUAAUCGCGAGGCUGCAAA	1	0	transposon	LTR	mdg3
6549	ACGAGCGCACGUGUUAGGACCCGGA	1	no match			
6550	UGCGCGAAGAAGGAGGACGAGAAGG	1	0	transposon	LINE	G2
6551	GCCUCAACCAACGCUCCGCAGA	1	no match			
6552	CAAUGAGAACGAUUACGAUCCCACG	1	1	transposon	LTR	ninja-Dsim-like
6553	AUAUCCGUUAUGGAAAAUUCAUCAU	1	0	ncRNA		rRNA
6554	GUCGCCAUUGGGACCACUAACUCUC	1	no match			

6555	AUCCCAGCACUACGAAGCGCCCUUC	1	0	no annotation		
6556	UCCGACUUCAAGAACCAGGCGAACAUAU	1	no match			
6557	GCGAUGAACAGAAAAAGAAUCACCCGA	1	no match			
6558	AACUGAAUAAACGAAAUGGAUCACACA	1	1	transposon	LTR	invader4
6559	CAAGAUAUGACCAAAGCAAAGACUAGA	1	1	transposon	DNA	1360
6560	UUGUCCGACAAUUACCGAAGUGAUGC	1	0	transposon	LINE	accord2
6561	AAAAACCUGAAAAAUGAUCCUGAGUCGC	1	no match			
6562	AAUUCCGAUAACGAACGAGACUCAAAU	1	0	ncRNA		rRNA
6563	ACGGUAUAUAUGAGGCAGCCAGAGG	1	0	transposon	LTR	DM88
6564	CCGCUAGAGGUGAAAUUCUUGGACC	1	2	ncRNA		rRNA
6565	CAUUGCCAAGGAGGACCCCCACAA	1	0	coding gene		CG12918-RA|CG12918
6566	UACGACGAGAACGAGAUCACGCGCGA	1	no match			
6567	AUACCACCCUUACUGUCGUUGACG	1	no match			
6568	ACCUACCGGAACAGACGACCAGCCACC	1	0	transposon	LTR	gypsy6
6569	GCUACGGUCAAUGUCUCUGAGGCU	1	0	transposon	LINE	Rt1b
6570	GGGGAAAGGACUCGGACAGUUUCU	1	0	transposon	LINE	accord2
6571	CGGCUCAUCACAUCCUGGGGCUGA	1	no match			
6572	CGGUAAGCGUUGGAUUGUUCACCC	1	no match			
6573	UGUCCUGGGCUGCACGCGCGCUAC	1	0	ncRNA		rRNA
6574	UCCAGAGGCAACUGGAUGCAUUGGA	1	0	transposon	LINE	G2
6575	AAAACCUGAAGAUGAUCCUGAGUCGUU	1	1	transposon	LINE	Helena
6576	AAAAGCCUGAAGAUGAUCCUGAGUCGC	1	2	transposon	LINE	Helena
6577	GGAGCUGCAUGUAGAGAAGCGAGAGGCC	1	1	coding gene		CG31641-RA|stai
6578	CACCGACUACAAAGAUGCCAUACGA	1	1	transposon	Unknown	RR50960
6579	GCGAUCCACUGAAGCUUAUCCUUUGU	1	1	ncRNA		rRNA
6580	ACCCGCUUGACCUUACUCAACACU	1	0	transposon	LINE	G2
6581	CCAGAAGGCCCGGGUUCGAUUCCCG	1	0	ncRNA		tRNA
6582	UCCUUGGGAAUGGCUUCACAGUU	1	1	ncRNA		alphagamma-element
6583	GAAACCGUCAUUCCAGCGCAAGUUGCCCC	1	1	transposon	LTR	ninja-Dsim-like
6584	GAAGUUGAUACACAAUAUAGACAGGCCC	1	0	transposon	LTR	Idefix
6585	UCAACUGUGGGGGGAUCCUCAA	1	no match			
6586	UAGAAGACGAUUUUCGGGCCGAAA	1	0	transposon	LINE	Cr1a
6587	AAAAAACCCUGAAGAUGAUCCUGAGUCGC	1	no match			
6588	CGCAAAAGUUGGCCGAACUUGAUU	1	1	ncRNA		rRNA
6589	GCCUCCUGGGAGCUCGGCGGGCGC	1	no match			
6590	AGCGAAUCCAGCGUAUCCCGAAGG	1	0	transposon	LTR	springer
6591	ACGAGCUCACGUGUUAGGACCCGAAA	1	no match			
6592	UUGAAGUCCACGAAUGUGCCGAGC	1	0	transposon	LINE	R1-element
6593	GACUACAUAUGGUUGAGGGUUGU	1	0	ncRNA		rRNA
6594	AUCUACAUCGUCGUAGAACCAAGCCGA	1	no match			
6595	GAACUUGACGACCCACGAUUAGUAAU	1	1	Su(Ste)		Su(Ste)
6596	AAAUCCGUAAACGAAGCCCAAAGU	1	1	transposon	LTR	Rt1c
6597	AACGUGUUGAACCAGGCUUCCAG	1	no match			
6598	AAUAGAAACGACAGUGUCCCGCGG	1	no match			
6599	CGCAAACCCAAAAUGACAAUCGCACGA	1	no match			
6600	AGAAGCGAUAUCCGAUGCCGAUUG	1	0	transposon	LTR	diver2
6601	AAAACCAUAAAGAUAGCGACAGGUUC	1	2	transposon	LINE	Doc3
6602	CUGAUCACAUUGCCUUCGUGUUUA	1	2	transposon	LINE	Cr1a
6603	ACCACCCUGAACGCCCCGAUCUCGUC	1	no match			
6604	GAUUACGUCCCUGCCCUUUGUACACAC	1	0	ncRNA		rRNA
6605	UAUCUCGAAAGACUCGUGCCGAUC	1	1	transposon	LTR	G4
6606	GUAAUUUCGGGAUAAGGAUUGGCUCU	1	1	ncRNA		rRNA
6607	UGCAAUCGAAACGAGAAACUUCAGU	1	0	coding gene		CG12717-RA|CG12717
6608	GCCCCCUUGAUUUUGAUUUUCAGUGU	1	no match			
6609	UACAGAUUCUAGAGACAAAGACGCAGU	1	0	transposon	LINE	INE-1
6610	GUUGAAAGCGAGGUGAACCCGAAU	1	0	coding gene		CG6279-RA|CG6279
6611	UCCUUGUAGAGGCCUUCUCUUCCCC	1	0	no annotation		
6612	AAAGCCAGUUACCGAAGCUGCGGAU	1	no match			
6613	UAAAAGAACAUUGAAGCGCUUGACU	1	0	Su(Ste)		Su(Ste)
6614	AAAAACCUGAAGAUGGUCCUGAGUCGC	1	2	transposon	LINE	Helena
6615	CUGUAACUCGGGAUAAGGAUUGGCUCU	1	no match			
6616	AGUCUCAACGGCGAGGUAUCAACGGG	1	0	coding gene		CG13255-RA|CG13255
6617	GCACCCGCCCGUUUGACGCCGCCU	1	no match			
6618	GCUAAUUAACACAAUCCCGGGGC	1	0	ncRNA		rRNA
6619	CUCGCCGAAUCCCCGGGGCCGAGGGAGC	1	no match			
6620	AACCAGCAGCAACUCCGAGCAUCAC	1	1	transposon	LINE	Doc4
6621	CACCCUGAACGCGCUCGAUCUCGUC	1	no match			
6622	UCCGACGAGCUGCUACCCGGGGAUUU	1	0	coding gene		CG6230-RA|CG6230
6623	AUCUAGGACAAUGAUGCGACCCGUCU	1	0	transposon	LTR	ninja-Dsim-like
6624	UCCGCUAGCUCAUUUCCCGGGAUGUU	1	0	transposon	LINE	Fw2
6625	UCCAAAGGCAUAGGGCAAGACAGAC	1	0	transposon	LINE	G2
6626	AGCCGGACGGUGGCCAUGGAAGUCGGAA	1	no match			
6627	UCUCCCGCCCACGACUUGACUCCG	1	no match			
6628	AGAGGUCUCCGGACGUGAUCACUG	1	1	ncRNA		rRNA
6629	AUGACAGACAUCCCACGCAUCCAGA	1	0	transposon	LTR	HMS-Beagle
6630	GCCCGUGGACGGUGUGAGGCCGGUAG	1	no match			
6631	AAAACAAAAGCAUUGUGAUGGCCCU	1	no match			
6632	AACACCGAAACGAAAAACGCCGACUGG	1	0	coding gene		CG30425-RA|RpL41
6633	AGACCGUAAACGAAAAGCUAAGAA	1	0	transposon	LTR	Idefix
6634	GAGGAAUUUGAGCUGCGAUGACGCCGA	1	0	transposon	LTR	ninja-Dsim-like
6635	AUCAAUCGCUGAAAGGGCACCGGGU	1	1	transposon	LTR	accord
6636	GCAGUAAUUCCGGGGAGUCAGA	1	0	coding gene		CG9660-RA|toc
6637	ACCGUCAUUCCAGCGCAAGUUGCCC	1	0	transposon	LTR	ninja-Dsim-like

6638	UUCGAGUUUAACAAAGCGCCCUUUGG	1	0	transposon	LTR	flea
6639	CCCCGUACUCUGUGGUGACCGACA	1	1	transposon	LINE	baggins
6640	UUAACAGACAAGGGCCAGUAAAA	1	0	transposon	LTR	Idefix
6641	AGGUGUUGUAAGUGGUUGAGCAGCUGCC	1	0	ncRNA		rRNA
6642	UGUAUAGAUGAGGUAUAAGGUUGGC	1	0	transposon	Low_complexity	F-element
6643	UCCGACCGGGGGCCCGUGGACUGU	1	no match			
6644	CCAUUCGAUUCAUGGUGUGACUACAA	1	0	transposon	LTR	springer
6645	ACGACGCGAGAGGUGAAAUUCUU	1	0	ncRNA		rRNA
6646	GAUCCUCAACUGUGAGGCUCAA	1	no match			
6647	UCCCGCAAUAUGACAAUCGCUCG	1	1	coding gene		CG40609-RA|CG40609
6648	GAAGAAGUUAUGGUUGGCGAGGUCGC	1	0	transposon	LTR	Idefix
6649	AAAGCCGGGACAUCGAAUUGCUGAAA	1	1	transposon	LTR	micropia
6650	GCAAGAGACUCGAGUAGGGCGGGAC	1	no match			
6651	AUGUUCAUCGUUAGACGGCUCGGGC	1	no match			
6652	CCCAGAAACACAGUGUUUGACGCCAAGC	1	0	coding gene		CG18743-RA|Hsp70Ab
6653	AAAACGCGAAUGGCAUCCGGAUGA	1	0	transposon	LTR	flea
6654	CGCAAAAGUUGACCGAACUUGAUUA	1	0	ncRNA		rRNA
6655	AAGGCGGGGACUAUUUACUUAGUU	1	2	transposon	LINE	Cr1a
6656	UGGACUCGCUGACUGCGCACAGUUU	1	2	transposon	LTR	micropia
6657	CCCAUCGCUGAUCGAGAUGAUGCGCUUC	1	0	coding gene		CG15286-RA|CG15286
6658	GAACCACUUUCAUGCUUGGGAUUGUGAA	1	0	ncRNA		rRNA
6659	CUUACAUUCAGUGGCAAGGUUAACC	1	no match			
6660	AGGAGCAUAUAAUGCCCCGUAGGUUGUCCC	1	no match			
6661	CCGAACUUGAGACCGCCCCUGGACGACA	1	1	Su(Ste)		Su(Ste)
6662	AACUGAAUAAAACGAAAAUGGAUGACAC	1	no match			
6663	GUUUUCAAGAAAAGAUGCCCGAACGACU	1	0	transposon	LINE	RR49783
6664	AGCUCUUGAUCGAAGCCCCGGU	1	no match			
6665	CAGCUGCCAUACUGCGAUCCACUG	1	0	ncRNA		rRNA
6666	GCGGUUAUUAGUGGAUUCAGCCUU	1	0	transposon	LTR	Idefix
6667	UGUUUCAUCGUUAGACGGCUCGUGCG	1	1	AT-chX		AT-chX-B
6668	UACAAUACAACGCUGACGACUAAAA	1	0	no annotation		
6669	GAGAAAUUCAACGGUGAUCCAGGC	1	0	transposon	LTR	McClintock
6670	AGCCGACUUAGAACUGGUGCGGACC	1	no match			
6671	UCUUAGAACAACUUAGACGACCAGA	1	0	transposon	LTR	opus
6672	UCCGAAGACAAGUAUACCGUGGCAC	1	0	transposon	LTR	micropia
6673	UGAAACAAGAAGAGGACGACGAUCAC	1	1	transposon	Low_complexity	F-element
6674	UGCUCUCGAAUGCAUGCCCGAUAU	1	1	transposon	LINE	R1
6675	AUCGAUGACGAGCUACGAAGGGA	1	0	transposon	LTR	ninja-Dsim-like
6676	UUCGAUUCCGGAGAGGGAGCCUGAGAA	1	0	ncRNA		rRNA
6677	UCCGAGAACAAACGAUACUGUUCGGC	1	1	transposon	LINE	G2
6678	GACUUGAUCUCGGUGUUCAGUACAC	1	0	repeat		LSU-rRNA_Dme
6679	CACGUCCGAACAUAAGUGUGUAAGA	1	1	ncRNA		rRNA
6680	CAACCAAAGAGGUAUUGCACUAGC	1	0	transposon	LINE	X
6681	AUGCCCUGAAGACCAUUACUCGUC	1	0	coding gene		pseudogene
6682	CCGUACUUUACUUUUGCAACAAAACGA	1	0	transposon	DNA	1360
6683	AGUCACGAUGCUUAGCGUCCGUCGC	1	1	transposon	LTR	GATE
6684	AACGCUUAUACAUAAGCCGAUCGCGGAG	1	0	transposon	DNA	NOF
6685	UGCACCGAAAUAUAUGCGAGGAGA	1	no match			
6686	CACGAAACUGGAACGGGAUUAAAAUC	1	0	no annotation		
6687	GCACUCGCCGAAUCCCGGGGCCGAGGG	1	no match			
6688	CACUGUGUCUCCCUCUUUUGGUCGCGG	1	0	transposon	LTR	DM88
6689	GCGGAGGCGGGCGGCGGGGGGGUU	1	no match			
6690	AAACCAAAGCUUUUGGCUCCGGGGGA	1	no match			
6691	UCAUUGAGCGAACCCGGGGAACUGAAA	1	no match			
6692	ACCACCCUGAACGCGCCCGACCUCGU	1	no match			
6693	GUGGCAUGUUUCCGGAUUCCCGCC	1	0	ncRNA		alphagamma-element
6694	UGCCGAUGACAACGCCUUCCUAGCC	1	1	transposon	LINE	G
6695	AAGAUGAGAUCUGCCUGGGGCCUUGA	1	no match			
6696	ACUCGCCGAAUCCCGGGGGCCGGGGGA	1	no match			
6697	CCAUACUUAGACUAGGAUACUACCCA	1	0	transposon	LINE	G2
6698	AACCUGAUUGAUAGCGUAGACGACG	1	1	transposon	LINE	Rt1a
6699	CACCCAGAAGGCCCGGGUUCGAUUCCC	1	0	ncRNA		tRNA
6700	UCGAAGGGAUAGGGUCGGUUCGUCCGG	1	no match			
6701	AGUCGUGGCCGAGCGGUUAAGGC	1	0	ncRNA		tRNA
6702	CGCCCCUGGACGACCCACGAUUUAACA	1	no match			
6703	CGAACCAGAGGCUGGGUACAUCG	1	0	no annotation		
6704	CCGUUCAAAAUGUUCAUGCAGGAGA	1	no match			
6705	GAACCAGGUGGCGCUGAACCCGC	1	2	coding gene		CG4264-RA|Hsc70-4
6706	CAUUCAUAACCGAGCGACUGUUCUA	1	1	transposon	LTR	GATE
6707	CGCCCGGGCGUCCUCCAGCGAUGAAAA	1	0	transposon	LINE	R1-element
6708	UACAUCGAAAAGCGAGUUGACGUGUUU	1	2	transposon	LTR	Max
6709	CAAACCAAUCCCAGCCAAACCCGUCAAA	1	0	transposon	LINE	G2
6710	UCCUUCAUGAACGAACCGUACCGAG	1	0	no annotation		
6711	UUCAAGGUUACCCCAGGAUGCUGC	1	1	repeat		trf
6712	UCCGAUUUCAAGUACAAGACGAACCGA	1	0	repeat		RSP
6713	UACUUCGAGACUUUUUCACCCGUGUG	1	0	transposon	LTR	297
6714	AAACUGGAUAAACGAAAUGGAUGACAC	1	1	transposon	LTR	invader4
6715	AGAUGGUGCGAGAACCCAGAGUAAU	1	0	coding gene		CG11719-RA|Mst98Ca
6716	GUCGCAGACGACCGGUACUGCAUCGACG	1	no match			
6717	CAUAGUCAUGCGAGGCGAACCGGGUGA	1	no match			
6718	CCAGACGUAAUACUCCUCUACCCGGA	1	1	transposon	LTR	HMS-Beagle
6719	UCUGCCCGAUUAGAAGAAUGGCCCGU	1	1	transposon	LINE	RR49783
6720	GUGGAGCCGGGCGUGGAAUGUGAGUG	1	no match			

6721	ACGGACCAAUGGCACAAUACUUGU	1	0	repeat		LSU-rRNA_Dme
6722	ACAUUCAGUAACCGAGCGACUGUUCA	1	no match			
6723	AUAAGGAUGAUGAUACCGACUAA	1	0	no annotation		
6724	CAGUACGAUAAUAUUUCCCGCGACAAG	1	0	transposon	LTR	gypsy12
6725	ACACGAGAAAACUUACCAGGUCCGAA	1	1	ncRNA		rRNA
6726	AACCAUUCGUAGACGACCUGCUU	1	no match			
6727	GGUAGCCAAAUGCCUCGUCAUC	1	0	ncRNA		rRNA
6728	UAACAAGAUGUAUACGAUGCGGC	1	2	transposon	DNA	1360
6729	GCAACCGAGGGAUCUCAUUACCCGA	1	0	transposon	Simple_repeat	I
6730	UCAGCUAAAAUCCGAGAUGGCUACCC	1	0	coding gene		CG12047-RA|mud
6731	AAAUCGCUCGAUGUCGUGAGGUGACC	1	1	no annotation		
6732	ACGUAAAACAAUUCAAGAUCUGGCCGGC	1	1	transposon	LTR	blood
6733	UGCGAAAGUAUAAGAUGUCCAGCGA	1	0	transposon	LINE	Rt1b
6734	CUUAUCAACAACAAACCCGGGGACGAGC	1	1	transposon	LINE	G6
6735	AGCGUCUAAAAAGAGAUGAGGCCUUGA	1	0	transposon	LTR	GATE
6736	AGAGAGAAGGCUCACCCGACGAGA	1	1	no annotation		
6737	AAAAACCUGAAAGAAUGAUCCUGAGUCG	1	no match			
6738	UGCUCUCGAAUGUAUGCCCCGAUAUACC	1	no match			
6739	GACGCGAUGUGAUUUCUGCCCAGUGCGC	1	no match			
6740	UGUCAAUUCAACGAGGACCGCUAUGAGA	1	0	transposon	LTR	flea
6741	CGGCAACUUCAUCGAUCUUAAACCGCC	1	0	coding gene		CG18628-RA|CG18628
6742	UCGCGAGUUCGAAUCCCGCCACGC	1	no match			
6743	UACAAGCAAGACCAUCACACCCGACGC	1	1	transposon	LTR	gypsy12
6744	GAUGCUCGAAGACACCACGGUGAA	1	1	transposon	LTR	ninja-Dsim-like
6745	CCCGAAAAAUCCCCCGGCUACGACCU	1	2	transposon	LINE	FW
6746	UAGCGAUCCGCGUCCCCAGUUCACC	1	1	transposon	LINE	R1-element
6747	UAAGACUUAAAGAUGAUGAACCCAC	1	1	transposon	DNA	NOF
6748	CUGGACGAGGCCGACAAGAACUCCGAU	1	2	coding gene		CG31359-RA|Hsp70Bb
6749	AGAAGGGCAAAAGCUCGCUUGAUCU	1	no match			
6750	CAACGGGUAACGGGGAAUAAGGGUUCG	1	1	ncRNA		rRNA
6751	GAUUAUGCCUGAACGCCUCUAAGGU	1	0	ncRNA		rRNA
6752	UCCGUCGUUAAUCCAUUCAUGCGCAU	1	0	no annotation		
6753	UAAGAGGGACGGCCGGGGGCAUUC	1	no match			
6754	GUCGUCGAAGGUUUCCCAAUGGA	1	0	transposon	LTR	ninja-Dsim-like
6755	CACGAUGAUCGAUGACGAGCUACGA	1	0	transposon	LTR	ninja-Dsim-like
6756	CUGGCUUAUAACCGUUCACUCUCCAA	1	0	transposon	LINE	Doc3
6757	AAAACCUGAAGAUGAUCCUGAGUCU	1	1	transposon	LINE	Helena
6758	GCUCGUAGCGAUUCUGACGUGCAAA	1	no match			
6759	AGUUAACUAUCCCAAAGGCAUCACGCG	1	0	no annotation		
6760	AACAACAACAACCAGAGCAAAGCAAG	1	0	transposon	Unknown	RR50960
6761	CACACUGGUACUGAGACACGGCC	1	no match			
6762	AACGAGCGCAACCCUUGUCCUUAGUU	1	no match			
6763	ACGUCGGCCAAGCUCCUGCAGCAU	1	no match			
6764	GCACACAGAGGAGUAGCAGAUGAGAGCA	1	no match			
6765	AGACACGCAAGAGCAGCUCCAGC	1	2	transposon	LINE	G6
6766	CCAAACUUAAACAAGACACUAGUCCGA	1	0	no annotation		
6767	CUGGAGCCACAGUAGUAUUUACAUUU	1	no match			
6768	CAAACAACAACAAAAGACGCAAGCGGGA	1	0	transposon	LTR	Rt1c
6769	GAGGAAUUUGAGCUGCGUUGACGCCGAA	1	0	transposon	LTR	ninja-Dsim-like
6770	UGUCAAUUCAACGAGGACCGCUAUG	1	0	transposon	LTR	flea
6771	GCCGAAGUCAUUGUGCGACAGAGUG	1	1	transposon	LINE	X
6772	AAGCAAUUAUCCGAUACCGCUUA	1	2	transposon	LTR	flea
6773	CACAGGUACGGGGCUUACUGAA	1	no match			
6774	UAGUUCAGUCUGAGAAGGGAGGAGUU	1	0	no annotation		
6775	UGAGGGGAUCCUCUAA	1	2	coding gene		CG4584-RA|dUTPase
6776	GUCACCAUACCCGACACUUACCCCAU	1	0	transposon	LTR	opus
6777	ACAUCAAAGACCACAUGCACUUACAU	1	0	transposon	LTR	Stalker2
6778	AUUGGGUGUAGCUACUUUUAUGGCU	1	0	ncRNA		rRNA
6779	AUACCGGAUGAUGUCGUAAGACCAAA	1	no match			
6780	GUUCCCGAUGUUUCUCGUACCCAAAA	1	no match			
6781	AAAAAGCGCGCUUGACUGCGAGACAGAC	1	no match			
6782	CUCAGCGAUGUGUGGACCAAGUCAAC	1	1	Su(Ste)		Su(Ste)
6783	ACCCUGAACGCUCCCGAUCUCGU	1	no match			
6784	UGUUUCAUCGUUAGACGGCUGGGC	1	2	AT-chX		AT-chX-B
6785	GUUUCGCAAAAGUUGACCGAACUUGAU	1	0	ncRNA		rRNA
6786	GCACACAGAGGAGAGCAGAUUGAGAGCA	1	no match			
6787	UUCGUGGAGUGAUUUGUCUGGUUAA	1	0	ncRNA		rRNA
6788	ACUCGAAUGUAACGUGCCCGAAUAUAACGGAGC	1	no match			
6789	ACGGGCCUAGAGUUAUUCCCCUAGA	1	1	coding gene		CG40635-RA|CG40635
6790	AAGCCCGUGGAGAUCGAGUGCCAGGA	1	0	coding gene		CG13891-RA|CG13891
6791	AAUGUAUGUGAAUACUCAGGCCGUUG	1	1	transposon	LTR	blood
6792	UCGGGCGCGGUUAGUACUUAGA	1	0	ncRNA		5SrRNA
6793	UUACCCUACUGAUGAUGUGUUGUU	1	no match			
6794	CGCUAUAACAACGUCUUCGUGCCGCGA	1	0	no annotation		
6795	ACUCUCCGAAUCCCGGGGGCCGAGGGAGC	1	no match			
6796	ACCACCCUGAACUCGCCCGAUCUCGUC	1	no match			
6797	AGCAAGAGACUCGAGUAGGGCGGGA	1	no match			
6798	AAACAUUCCAGAGUCCGAGUAAUA	1	0	transposon	LTR	Stalker4
6799	UAGAGUAGAAGAAGUCGAAAGCGG	1	0	transposon	LINE	accord2
6800	AAACGAGAGAAGCAUUUUUGCCGCA	1	1	transposon	LTR	gypsy12
6801	UGAUCACGAGUUAGUCGGUCCUAAGUU	1	0	ncRNA		rRNA
6802	AAGAGCGCGUGAAACCGUUAAGAGGU	1	no match			
6803	CAAGACAAUGACACUAAUGCAACCAAA	1	0	transposon	LINE	G2

6804	UUCGAGAACAAACGCUACUGUUCGGCC	1	0	transposon	LINE	G2
6805	ACGUGGGGGUCGCAGGUUCAAAUCCU	1	no match			
6806	UAGCUAGAGAUAGUUUUCGUCGAGC	1	0	transposon	LINE	Cr1a
6807	AAGGCCGAGAGCUGAUGACGAGUUA	1	no match			
6808	AAGAGGGCGUGAAACCGUUAAUAGGU	1	no match			
6809	AUCAAUCGCUGAUGAGACACCGGGAU	1	1	transposon	LTR	accord
6810	ACCGAUAUAAUUCUAACUGAGACCCCCC	1	2	transposon	LTR	G4
6811	CUCAUCCAGCAUGUGGAACGGUU	1	0	coding gene		CG1609-RA|Gcn2
6812	UGGAGUCCUGUUGAUCAUUUGCCC	1	0	transposon	LINE	Cr1a
6813	AAGAGUGAGAGCGAAACGUGGAGUA	1	1	transposon	LINE	Cr1a
6814	AACUGAAUAAACGAAAUGGAUGUAC	1	no match			
6815	CGACUUUCCGGCGGUGCCCGGCUCCAG	1	0	coding gene		CG9078-RA|ifc
6816	CGAAUACGAAGAACAGGCAGAAUC	1	0	transposon	LTR	gypsy12
6817	AUCCGGAGAUGGGGUCUUAUGGCUGGA	1	no match			
6818	UAGAGAGUGAUGCCGAGUAUAAAAUA	1	2	repeat		trf
6819	UGAGACUACAAACGGAUUGCGAACC	1	1	no annotation		
6820	CAAUGAGUUACGCGAAUGUAACGAGUA	1	0	transposon	LTR	Idefix
6821	ACAUUCACCUGGGACCGAGGUUCGAAC	1	0	coding gene		CG2304-RA|Trc8
6822	UCCGGACAAACACACAAGACCCGAU	1	0	transposon	LTR	opus
6823	AGGACACUAGCUUCUUAAAUGGA	1	0	ncRNA		rRNA
6824	CAAGAAGGCCGACUACUGCAACGAU	1	0	coding gene		CG12218-RA|mei-P26
6825	CACUGUUCAUACAAUUGAGGCUGGA	1	0	transposon	Low_complexity	F-element
6826	GUUCACGGAGAAGAUUGCGCCGGAGC	1	1	coding gene		CG11719-RA|Mst98Ca
6827	ACUCGAAUCAACAAAUUGCGCCGAUA	1	0	coding gene		CG41545-RA|CG41545
6828	AAUUCCAUCGAAAAGACAUUGGCCA	1	2	coding gene		CG41322-RA|CG41322
6829	AAUUGGCCCAAACGGAUUUCCCUAU	1	1	repeat		SAR_DM
6830	ACGAGCGCAACCCUUGUCCUUAGU	1	no match			
6831	CAAGCAUUCGACAAAGUCUGGCUCGUA	1	2	transposon	Low_complexity	F-element
6832	ACUCGCCGAGUCCCGGGGCCGAGGGAG	1	no match			
6833	AAGACCAAUAACAUUUUAAGGAUA	1	0	transposon	LTR	297
6834	CGCGUGGGUUCGUAUCCCAUCCUCGU	1	1	ncRNA		tRNA
6835	ACGGCCAAUGGCACAAUACUUGU	1	no match			
6836	GAUGCGAGAGGUCCCGGGUUCAUAUC	1	2	ncRNA		tRNA
6837	AAAUGUGAUUUCUGCCCAGUGCUCUGAA	1	1	ncRNA		rRNA
6838	GGCAGCAGGCGCGUAAAUUACCCACU	1	0	ncRNA		rRNA
6839	GAUAACGAACGAGACCUUAACCUACU	1	no match			
6840	GCUACUGUCUUACCAUCAUCGACCGC	1	0	transposon	LINE	Doc4
6841	UCUGGUGGCGAUAGCGAAGAGGACACAC	1	no match			
6842	AUGAGAUUGAGCAAUAACAGGUC	1	0	ncRNA		rRNA
6843	AGAGGGUGAGAAUCCCGUGUGGCGAGGA	1	no match			
6844	GGAACGUGUAUCAUUUCCUAGCCAUUAU	1	0	repeat		LSU-rRNA_Dme
6845	AAGGAAAUGAUUGUUGAGUGCUU	1	0	transposon	DNA	1360
6846	CAGAGAUCCGUGUGGCCUAUCGCUAU	1	0	coding gene		CG32626-RA|CG32626
6847	AUGACCUAAACACGAUUAUAGAUAUAA	1	0	transposon	LTR	Idefix
6848	AGCACCCUGAACGCGCCCGAUCUCGU	1	no match			
6849	CGCUCACAGACUACCUAAAAGGCGCAU	1	1	transposon	LINE	Doc4
6850	AAGUCAGAUAGCGUUAGUUGCGGGA	1	0	transposon	LTR	springer
6851	UGCUCUCGAGUGUAUGCCCGAUAUA	1	1	transposon	LINE	R1-element
6852	CAAAGUGUAAAGGCCCAUAUGUG	1	0	transposon	LTR	flea
6853	GACCUCAAUAAGCUGCCGGACCUGA	1	0	transposon	LINE	F
6854	CAGCAUUCGAUAGAGUAUGGCUGGA	1	1	transposon	LINE	Doc3
6855	AAGUCAAAGACCUUGUUAACGAAGCC	1	0	transposon	LTR	HMS-Beagle
6856	AAACAUGAUAUGUCCCUAGCGUAAGA	1	1	transposon	LINE	Cr1a
6857	AACGGCGAGUGAAGCGGCAAAAGCUC	1	no match			
6858	ACUACUUUCAGGACACGUUCAACCAG	1	0	Su(Ste)		Su(Ste)
6859	ACCCUGAAGCGCCCGAUCUCGU	1	no match			
6860	UACAGACUUAGCCCCGAAGAGAGAGA	1	0	transposon	LTR	micropia
6861	UAGGAGUUCACCGCCACGACGA	1	0	transposon	LTR	GATE
6862	UAUGACCCUUACAAAAAUGCGAA	1	no match			
6863	AGCGAAAUGACUCUGUCCCGAUUCAAU	1	1	transposon	LINE	R1-element
6864	UUGUGGGCCGGUACUAUUACUUUG	1	0	ncRNA		rRNA
6865	GAUCCUCAACUGUGAGGGAUCCUCAA	1	no match			
6866	GCGCCGGGCCGUACCCAUAUCCGC	1	no match			
6867	AGUACUGGAUGACGUCAUAUUUUG	1	no match			
6868	AUAAAACAUACGCCCGGUCGCGCC	1	0	transposon	LTR	GATE
6869	UUCAGCGAGAUAGUGGACGUGAUCCUGA	1	1	repeat		trf
6870	AUGAGAACGAUGACGAUCCCACGGCU	1	0	transposon	LTR	ninja-Dsim-like
6871	GGAAAGGCGAAAAGAACCCCGGG	1	no match			
6872	CUGCCGAACGUAGUCUCUCGAGAGC	1	0	transposon	LTR	invader1
6873	AAAACCUGAAGAUGAUCCUGAGUCC	1	1	transposon	LINE	Helena
6874	CAGAGAGAAGACUCACCCGACGAGA	1	0	no annotation		
6875	CGAGAGAAGACUUUUCACCCGAUGAGA	1	no match			
6876	AAGAGGGACGGCCGGGGGCAUCG	1	no match			
6877	ACCGGAAACAUUAGUCGAGGCGUAAA	1	2	transposon	LINE	Rt1b
6878	ACGAGAAACUUUAUCCCAGUCGGA	1	1	no annotation		
6879	UCCACGGCCGACCGAGAAAUUCCC	1	no match			
6880	CUUACCUUCAACAGCUAAGGGACCA	1	0	transposon	LINE	S
6881	UGUGUUUCCCCGGCGACUAUACAGCC	1	no match			
6882	GAUGUCGGCUCUUCCUAUCAUUGUGAA	1	0	ncRNA		rRNA
6883	CUGGUUAAUCCGAUAACGAACGAGA	1	no match			
6884	GAUGGAAGACCGAAACAGUUCUGA	1	0	ncRNA		rRNA
6885	AGGUGCAGGUGGACUACAAGGG	1	no match			
6886	CACCGACGAAUAUCUGACAAUUGAUGA	1	0	transposon	LTR	Dm88

6887	CAGGGCUGCAAUUGUUUCCCGUGG	1	0	transposon	LINE	Fw2
6888	GUGAAGAAAUUCAAGUAAGCGCGG	1	0	ncRNA		rRNA
6889	ACAUUCGAGACCCAACGGCACGUGAC	1	0	transposon	LINE	Rt1b
6890	GCCUAUACGAGCCGGAGUACCUCA	1	0	transposon	LINE	G2
6891	ACACAAUCGAUCCCGAAAUUUUGA	1	0	transposon	LTR	17.6
6892	CUAGCGGACAACCUAGCCAGGUAA	1	0	transposon	LINE	baggins
6893	UGCUCUCGAAUGUAUGCCCGAUACA	1	1	transposon	LINE	R1-element
6894	UUCCGUAAGUAGUGGCGAGCGAACGCGGAUU	1	no match			
6895	CGGAAGACCCUCUCCCCGGCGCC	1	no match			
6896	UUCAACCGGACGGGCCUAGGGAACUU	1	no match			
6897	AACUAGAAUAAACGAAAUGGAUGACA	1	no match			
6898	ACGAGGACGACCAAGAGAGCUCUGA	1	0	transposon	LINE	G6
6899	UCUCAUCGUCGUAGAACAAGCCCGGGGU	1	2	Su(Ste)		Su(Ste)
6900	UCCGAUCUCCGACUAGGCCCGCGU	1	0	transposon	LTR	GATE
6901	AUCAAUCGCUGAAAAGGCACCGGGAC	1	0	transposon	LTR	accord
6902	GCAAACCAAUCCCAACCAAACCCGU	1	0	transposon	LINE	G2
6903	AGCUACCAUGCUCCGCUAUACCGACCG	1	0	coding gene		CG4836-RA|CG4836
6904	GACAGAUUGAUAGCUCUUUCUCGA	1	0	ncRNA		rRNA
6905	AUCAUUCAAAGCCAGCUGGAUGCCC	1	0	transposon	LINE	G
6906	GGAGCAAACGCAGACGGAGAUGU	1	0	coding gene		CG15307-RA|CG15307
6907	GUGCGUAUUGGGUUGACACGACA	1	0	transposon	LINE	Cr1a
6908	UCCGAAAUGAACGUCUUCAUUGGACCC	1	1	transposon	LTR	gypsy8
6909	CAAGCUGAUCGAGCGCAACUGCCGU	1	1	coding gene		CG18743-RA|Hsp70Ab
6910	UGUUGACAAAAAUGAGGCUGACCAA	1	0	transposon	LTR	roo
6911	UCUAGAGGACGAUAUAUUGACAUAA	1	1	transposon	DNA	transib2
6912	AACCCAGCCGCAGACGUCAGUAAAUC	1	0	coding gene		CG1484-RA|fliI
6913	AUCUGGGAGACAGACGGCGGGUGC	1	no match			
6914	AUGAAAGGUUCGUCCACCCGGGGGCGCCA	1	no match			
6915	UCCGUUCCCGCUAGAGACUACC	1	no match			
6916	AAUUCCGAUAACGAACGAGACUCUG	1	2	ncRNA		rRNA
6917	AAAGUUCUGAGAACCAUCACUGGGGC	1	1	transposon	LINE	F
6918	AAUCGAUGUGUUACUGCUCAGCGUAU	1	2	transposon	LINE	X
6919	UUGAUCACGAGUUAGUCGGUCCUAAGUU	1	0	ncRNA		rRNA
6920	UCCCGUAAACCGCCAUCAGCAACAC	1	0	transposon	Low_complexity	F-element
6921	AUACAUGCCGACGGGCGCUGACCCCCCU	1	no match			
6922	GCACUCGCCGAAUCCCGGGGCCGAGGGAGC	1	no match			
6923	AUAGCGUACCGGAGUGGACUGCCGU	1	0	ncRNA		7SLRNA-RA
6924	UGGGAUUGUACAAUUUCAAUACGGGA	1	0	transposon	LTR	diver2
6925	UAUAGGCUCUCUCAGUCGCUUCCCGGGA	1	no match			
6926	AACAGAAGAAAGAGACCGCAUUACCCA	1	1	transposon	LTR	Idefix
6927	ACUCUGAACGCGCCUGAUCUCGU	1	no match			
6928	UCCUCGAUAGUAUAGUGGUCAGUAUCCC	1	1	ncRNA		tRNA
6929	AACCGCAGCAGGUCUCCAAGGUGAAC	1	no match			
6930	CCGACACUUACCCCAUCCCUGAUA	1	1	transposon	LTR	opus
6931	GCGGAAAGACGGAAAGACCCCAUG	1	no match			
6932	AGUGGCAACUCAUUCCUGCCCGAGAAG	1	0	coding gene		CG4068-RA|CG4068
6933	CGAGCUCGAGGAAUACGGCCUUC	1	1	transposon	LTR	ninja-Dsim-like
6934	CAACCAGAGAUCUGGAGGGCCCA	1	1	no annotation		
6935	UGCUCUCGAAUAUGCCCGAUAU	1	no match			
6936	AAGUACGAAACAUGCUCUGACCUAA	1	0	transposon	LTR	Rt1c
6937	GAACUGAGAACCGUUGGACCAUG	1	0	coding gene		CG17376-RA|CG17376
6938	ACUGACAAUGACAGGGCAAUUUAUAG	1	0	transposon	LTR	gypsy4
6939	AAAACCAUUUAUCCUGCAGUGUGAU	1	1	transposon	LINE	Circe
6940	AGAGUACUUCAGCCAGACACUGGAC	1	1	Su(Ste)		Su(Ste)
6941	UCCUAGGCAGAGACGGCGUCGCUCGAGU	1	no match			
6942	UCACAUGAACUUGGAAUUCCCAGUAAGUU	1	1	ncRNA		rRNA
6943	GACUGCAGAUCGAGAGGUCCCCCGGUU	1	no match			
6944	AGAAGGCCCAAGUUGUGUAUGCAUAU	1	1	no annotation		
6945	GCUCGACGGGCUCAAGAUGUGUGACG	1	0	coding gene		CG17377-RA|CG17377
6946	GGGUUCCGGGGGGAGUGUGGUUGC	1	no match			
6947	UGGGGAAACCCACUCAUCCUGAUUAUC	1	no match			
6948	AAUCCCCCAGGACUUGAUCGCAUCGA	1	2	transposon	LINE	G2
6949	UAAUAGUGGGUCCGAGGAAUCAAUAGC	1	0	transposon	DNA	transib2
6950	AAUGGGCGAAGCUACCAUCUGUGGGAU	1	no match			
6951	UUGUAAACAAAUCGUAAAGCCUGACC	1	0	transposon	LTR	17.6
6952	CCCCAGGAGGACAACCAUCUUCUCUACC	1	no match			
6953	ACAACCAAGACGUUUGGCAGAGUGUGA	1	0	transposon	DNA	transib4
6954	CAAAGAUUAAGCCAUGCAUGUCUAAG	1	0	ncRNA		rRNA
6955	GAAGCCAAUGAAAAACAGCCUGCCGA	1	0	transposon	LINE	G2
6956	ACACGUUCAACCAGAUGGGCCUG	1	0	Su(Ste)		Su(Ste)
6957	UCCAAUUCCAAGGACACUCAAAGA	1	0	transposon	LTR	HMS-Beagle
6958	AAUUUCGGAGCACAAUUUGGCGA	1	0	transposon	LTR	ninja-Dsim-like
6959	ACUUGAGACCGCCCCUGGACGACCCA	1	0	Su(Ste)		Su(Ste)
6960	AGGGAAAGGAGCCGAGGCGUCGUC	1	1	transposon	LTR	ninja-Dsim-like
6961	UCUCCCCGCGGAAAGACGGAAAGACCCC	1	no match			
6962	CGCCCCUGGACAACCCACGUCCCUAU	1	no match			
6963	AAAACCUAGAACAUGAUCCUGAGUCG	1	no match			
6964	GACAGUCCAAACCCCGUAACUCGAACA	1	0	transposon	LTR	GATE
6965	ACCGAACGUAGUCUCUCGAGAGCGA	1	1	transposon	LTR	invader1
6966	CAGUCUGAGAAGGGAGGAGUUACCGCC	1	0	no annotation		
6967	CGGAUGUGCUGACCCCUGCGAUUUCCC	1	no match			
6968	AACCCUGAUGGAAGACCGAAACA	1	0	ncRNA		rRNA
6969	UAACCAAUACUAAUUGCCCGUGAGGC	1	no match			

6970	AACGCAGACAAAUCCUUCCAAACCA	1	1	transposon	LINE	G
6971	ACUCGCCGAAUCCCGGGGGCCGAGGGAGCG	1	no match			
6972	ACUGGGGCGGUACACCUGUCACUGAC	1	no match			
6973	CGAGAGCGAGAGAUAUUCCCGUUG	1	no match			
6974	CGCUAAACACACAAGAUUGCCACC	1	2	transposon	LINE	F
6975	ACAUUCAUAACCGAGCGACUGUUCACC	1	2	transposon	LTR	GATE
6976	CAGUACGUGAAACUGCUUAGAGGU	1	0	ncRNA		rRNA
6977	AAUCGCUCGAUGUCGUAAGGUGAC	1	0	coding gene		CG40609-RA|CG40609
6978	GUCGGCGCGAGCCCUGGGAGGAGUU	1	no match			
6979	AUGAUAUGCACCGAAAGUAUAUGAGA	1	0	Su(Ste)		Su(Ste)
6980	CGCCGGAAACGAGCCCAGGGAUCAGGCCA	1	0	coding gene		CG30460-RA|CG30460
6981	AGGAAUACAACUGCGAACUCAUC	1	0	transposon	LTR	opus
6982	UAAGCCCCGUACUCUGUGGUGACC	1	0	transposon	LINE	baggins
6983	ACCCUGAACGCGCCCGAUUUCGU	1	no match			
6984	AUUGUGGGCCGGUACUAUUACUUUU	1	1	ncRNA		rRNA
6985	AAUAAUGAGAUUGAGCAAUAACAGGUC	1	0	ncRNA		rRNA
6986	GUCCCAAGGCCAGCUCAGUGCGGACAGA	1	0	ncRNA		rRNA
6987	GUGUACGCGACUUUUCUAUUUAAGU	1	0	coding gene		CG10497-RA|Sdc
6988	GCCACCGAGAUCGUGCGAUUUGACCU	1	0	no annotation		
6989	CCCAGCUCGGGAGGUAGUACGACGUAA	1	no match			
6990	CUGACAAUGACAGGGCACUUUAUAG	1	0	transposon	LTR	Burdock
6991	AAAUCAUGCCGACGGGCGCUGACCCCCCC	1	no match			
6992	UAAACCGUCAUCCCAGCGCAAGUUGCCC	1	1	transposon	LTR	ninja-Dsim-like
6993	GACGAGACUUUGGGUGUGCUUAUGU	1	0	coding gene		CG5263-RA|smg
6994	AGCAGCAACCCAACGUCGAUGUGCGACC	1	0	coding gene		CG3213-RA|CG3213
6995	CCCGAUAUUAAUCUAUCCUGAUUU	1	0	transposon	LTR	Idefix
6996	UUAGAUAUGCUAGCAGACUGGAGA	1	1	transposon	LINE	G2
6997	AGUUACGAGACAAAUUGGGUUUGCUGC	1	0	transposon	LINE	Cr1a
6998	AGUACGAUCCGUCUGCAAACUUC	1	0	transposon	LTR	opus
6999	UGUUCAUCGUUAGACGGCUCGGGUA	1	no match			
7000	UCAACUGUGAGGGGACCUCAA	1	no match			
7001	ACGAGCGCACGUGUUAGGACCCGUAAA	1	no match			
7002	AAACCAAGCUUUUGGGCUCCGGGG	1	2	ncRNA		rRNA
7003	UGAGACUACAAACGGAUUCCGGACCG	1	1	no annotation		
7004	CUGACAUUAAUGAAGUGUUGGCC	1	0	transposon	LTR	HMS-Beagle
7005	CAGCUGCCAUACUCCGAUCCACUGAAG	1	1	ncRNA		rRNA
7006	AAGGACCAAUUGGAGGGCAAGUCUGG	1	0	ncRNA		rRNA
7007	AAGACCACGAGCGGUGUAGAACGG	1	0	transposon	LTR	HMS-Beagle
7008	ACCCUCUGAUGAUGUGUUGUUGC	1	no match			
7009	UAUACCUAUACAUUACCGCUAAA	1	0	ncRNA		rRNA
7010	CUCCAUCUCAGAGCGUAGAAUUUGGAC	1	no match			
7011	AACAACAACAGUGCAGACGGCGCCCGC	1	1	transposon	DNA	H
7012	UCGAGAAGAAGAUGAAGCGCAACAAGC	1	0	transposon	LTR	ninja-Dsim-like
7013	GCGAGCGCACGUGUUAGGACCCGAAA	1	no match			
7014	CAUGUCAUAACAACUAAACCCGAAGAC	1	0	transposon	LTR	Quasimodo
7015	AGACGAUGACGAUCCCACGGCUGGAGC	1	2	transposon	LTR	ninja-Dsim-like
7016	AACCGGAGGAAGGUGGGGAUGACGUC	1	no match			
7017	AGGGAAAUUGAGUGGCAAGAGUGA	1	0	transposon	LTR	micropia
7018	GUUCGGACAUCAGUUGGGAUCGCG	1	0	coding gene		CG4825-RA|CG4825
7019	UGGACUCGUUGACUGCGCACAGCU	1	0	transposon	LTR	micropia
7020	AUGAAUGAUGGACUGAAAUGG	1	no match			
7021	CGCAACUUGCUGAACCGCACAAUC	1	0	coding gene		CG13391-RA|Aats-ala
7022	AAACUACCCUGACCCUUGUCCCGCAG	1	1	coding gene		CG30497-RA|CG30497
7023	AAUACAGAUACCCACUUUCUUUGGUGA	1	0	transposon	LINE	Rt1b
7024	ACUUGCCGAAUCCCGGGGGCCGAGGGAG	1	no match			
7025	CCCUCAGAUUUGCUGGCCCUCACAC	1	0	transposon	LTR	Max
7026	ACCCUGAACGCGCCCGAUCUCGGUC	1	no match			
7027	CAUCCAGUUAACUUUGCACAUCAAGCA	1	0	coding gene		CG12717-RA|CG12717
7028	UUCAACCGGACGGGCCUAGAGUUC	1	0	no annotation		
7029	CCGAGCAAGACAUUGAUCACUUUGC	1	0	transposon	LINE	Doc3
7030	CGAGCGACCGGGACCAGGCCAGU	1	no match			
7031	UUAACAACGAUGAUGAUUCUGGCCC	1	0	transposon	LTR	Max
7032	AAUACAAACAGAAUUCCCGAAUGAGAA	1	1	transposon	LINE	Cr1a
7033	CAACCGAUAUGCUAUCUUGGCUGAU	1	1	transposon	LINE	Doc
7034	UUAACAAAGACCCAAGUGCUGAAGUGC	1	0	no annotation		
7035	AAACCUAGAAGAUGAUCCUGAGUCGCA	1	no match			
7036	UUCUAGAACGAAAGCGAGGACGA	1	no match			
7037	CUGUCAGGUACCCCGUUCAGCCCUAGUG	1	0	transposon	LTR	HMS-Beagle
7038	CAGCCUGAGAGGUGAUGCGUAGCCG	1	no match			
7039	AACCUAAACGAUGCCAGCUAGC	1	2	ncRNA		rRNA
7040	AGGCGCGGGUAACCCGUUGAACCCCAUC	1	no match			
7041	AAAAGGUGGAAAUGAAGAAAGUACA	1	no match			
7042	UGCCGCCGGAACUGAUUGUGACCACAC	1	0	coding gene		CG32628-RA|CG32628
7043	CCUUAUGGGACAUGUGCUUUUAUUAGGC	1	0	ncRNA		rRNA
7044	GGAUCAACUGUUGAGGGGAUCCUCAA	1	no match			
7045	UAGUACAAUCUAUCAGACUGAGUU	1	0	transposon	LINE	Fw2
7046	CGAGUAAUAUGAGCGCCAGAGU	1	no match			
7047	UGGCUAUAGCGUCGGUGUCCCACCCG	1	no match			
7048	ACCGACGACAUCCAGACCAAGGAGUA	1	0	coding gene		CG8565-RA|CG8565
7049	CUGACAUUACAUCAAAAUUUUUUUGA	1	no match			
7050	GCCAAUUACAUUAGACGAGACGGCCCA	1	1	transposon	LTR	opus
7051	GGACAAAACAACAUCCCGGGAAAAUGAGC	1	2	transposon	LINE	baggins
7052	AUGGCUCGCUCAGUCGCUUCCCGGGA	1	1	ncRNA		rRNA

7053	AUGCCAAUCACGCCGUUUGAACCCUGC	1	0	transposon	Simple_repeat	I
7054	UGGCAACAUGUCGAGCUCCCAGA	1	no match			
7055	UCCUAGAAUCGACCGACCAGCUGUG	1	0	transposon	LTR	invader3
7056	CCCGGGCGUUCUAUAUAGUUAUGUAUUA	1	2	ncRNA		rRNA
7057	AGCUGGGACUGAGGACUGCGACGUAA	1	no match			
7058	AAUAAACAAACUGACCGACACCGUGA	1	0	transposon	LTR	invader3
7059	GACUAGACGAGACAACGACGGC	1	0	transposon	LINE	Rt1a
7060	AAAGAGAGUAAUUUUGGCCGUCA	1	0	transposon	LINE	Cr1a
7061	GAAUACGUUCCCGGGCCUUGUACACACC	1	no match			
7062	CAAGCUAGACUAUGUAUCCCCCGGGU	1	no match			
7063	UCCGGAGAUGGGGUCUUAUGGCUGGAA	1	no match			
7064	ACGCAGGUGUCCUAAGGCGAGCUC	1	no match			
7065	UGGGCCCGAACCCUUCUGGUGUUAUA	1	no match			
7066	GAGUCCGUGGCAUUCAGGUUGGCUAAG	1	0	ncRNA		7SLRNA-RA
7067	GACCAGAGAGGGUGAUCGGCCACA	1	no match			
7068	AAAGGAGUUUCGACGGAUACGAAUGC	1	0	coding gene		CG8280-RA|Ef1alpha48D
7069	UGCCGGGGCGAGGAGGAGGACUA	1	0	Su(Ste)		Su(Ste)
7070	ACACAGAGGAGAGCAGAUAGAGCA	1	no match			
7071	AUGCCGGGGCGAGGAGGAGGACUAGC	1	0	Su(Ste)		Su(Ste)
7072	AAGGACGUUAAUAAAGAUGCCGUGGU	1	2	coding gene		pseudogene
7073	CACAAAGUUAUUCUUCUACCGAUGGA	1	0	transposon	LTR	Rt1c
7074	AAAUCGAGCAACUCCUCCACGUAAAA	1	1	transposon	Low_complexity	F-element
7075	CAUCCUAGAAUCGACCGACCAGCUA	1	0	transposon	LTR	flea
7076	GGGACAUGUGGCGUACGGAAGAC	1	no match			
7077	AAGGGGUCUAACACGUGCGCGAGU	1	no match			
7078	AACCACCCUGAACGCGCCCCGAUCUCGUC	1	no match			
7079	UACCACAUCUAAGGAAGGCAGCAGGCGC	1	0	ncRNA		rRNA
7080	CGCAAACCCAAAAUAACAAUGCAGGA	1	0	repeat		trf
7081	CAUCGAUGUACUACUGGUAUCAGA	1	0	transposon	LINE	G2
7082	AGGAUCAAUAGCCGAGUCGAACUGGCCA	1	0	transposon	LINE	INE-1
7083	GCCCCGGCUAGCUCAGUCGGUAGAGCAUGA	1	1	ncRNA		tRNA
7084	GCAACAGAGAAAGAGGCGAGGCAG	1	0	coding gene		CG40084-RA|CG40084
7085	CAGAGUUAAAGGCGGCUUCUUUGU	1	0	transposon	LTR	invader3
7086	UCCGAAGAUAACGAGGAGCAGAGCG	1	1	transposon	LTR	297
7087	CCUGGGAAGUCCUCGUGUUGCACCUCU	1	no match			
7088	GCCACCCUGAACGCGCCCGAUCUCGU	1	no match			
7089	AGCACGGGAGACACACGGGGGGGUGCU	1	no match			
7090	ACCCUGAACGCGCCCGAUCU	1	no match			
7091	GACUUCCGAGAAAGCCCGCAGAAUCC	1	2	transposon	LTR	flea
7092	AGCUCUCGAGGCUCAGUGCAGGUGGCG	1	0	coding gene		CG34450-RA|CG34450
7093	AAAGCACCGUUUCCCGUCCGAUC	1	no match			
7094	UGUGUUUGUAUUGCGAGUAGACU	1	1	transposon	LTR	invader1
7095	UUGUGAGUUAUCCGUCUAGGUGAG	1	0	transposon	LTR	micropia
7096	UCCAAGGAAAGAAGAUAUCGCAAAGGA	1	0	transposon	LINE	G2
7097	CUGGGUAGCUAUGUCUGGAACGGA	1	no match			
7098	AGGUGUGUGAGGCGAACCAGGGGAAC	1	no match			
7099	AAAACCUGAAGAUGAUCCUGAGUCGCUA	1	2	transposon	LINE	Helena
7100	GAUACCGUUAAGCGACUAUUUGUAGG	1	0	transposon	Low_complexity	F-element
7101	UGUGCUAGAAGACGAUUUUCGGGUU	1	1	transposon	DNA	1360
7102	ACAAAAUGUAUUCCGACUAAAUCCC	1	1	transposon	LINE	Doc
7103	GGUAACCCAUGCAGCCGAGCGGUCC	1	no match			
7104	AGGAGACGUUAACUCGAGGC	1	2	transposon	LINE	baggins
7105	CUGUGAGGGAUCCUCAACUGUGGGUCAAU	1	no match			
7106	UUAUAAGGAUGAUGAUACCGAUUAA	1	0	no annotation		
7107	AUGGCUCUCUCAGUCCCUUCCCGGGA	1	1	ncRNA		rRNA
7108	CAAAAGUAUUCGAGUUGCGACACAC	1	0	no annotation		
7109	UGUUUCAUCGUCAGACGGCUCGG	1	1	AT-chX		AT-chX-B
7110	UCAGCAUUUACCCUCGGCGGCCCCCCAC	1	0	no annotation		
7111	UGGCGAGCAAUCAUCCAAGGCUGAGUUU	1	2	transposon	LTR	mdg3
7112	AACCGGGAUUGCCUUAGUAACGGC	1	no match			
7113	AACUGAAUAACGAAUGGAUGACAC	1	no match			
7114	AGAUGUCCGGGGCUGCACGCGCG	1	1	ncRNA		rRNA
7115	ACAGCUUAUCACCCCGGAAUUGGUUU	1	no match			
7116	UCUGAUGUUACGCCCGAAGAGAUAA	1	0	transposon	Low_complexity	F-element
7117	UCAACAAACAACAGGUUAUGGGCC	1	0	transposon	LTR	297
7118	UCCCUGGGUACCGAUUUUAGCCGC	1	0	no annotation		
7119	UUGGCAUAAAAACAGUGAUAAAGUGC	1	0	transposon	DNA	1360
7120	CUAGAAUUGAUUGCAUUUUGGAU	1	1	transposon	LTR	gypsy8
7121	GGUUCUACCCGACGACGCUGCGCUAACC	1	no match			
7122	GACUUGAUUAAAACCAGGCCCCUGU	1	1	transposon	LTR	opus
7123	AUCUCGCGCCGGGCCGUACCCAUAU	1	no match			
7124	AGGGUCGGGCCUGGUUAGUACUU	1	no match			
7125	ACCAGAGAUCUAGAGGGCCCAGA	1	2	no annotation		
7126	ACCUUCAUCCAAGCGAACUGUGUCCCGA	1	no match			
7127	UCGCCGAAUCCCGGGGCCGAGGGAGCG	1	no match			
7128	CUGUUCUGCUUUCAUUGGUUUUCAG	1	0	ncRNA		rRNA
7129	UCCUAGAAAAAACGAGGUUCACCCAAU	1	0	transposon	Low_complexity	F-element
7130	GAGCCCAAACGUGAACCAGGUUCGU	1	1	transposon	Unknown	RR50960
7131	CAAGUCGGUCGACGGCGUGCUGAUCA	1	no match			
7132	GCAUAGCUAGAGUACGGUAGAGGGGGA	1	no match			
7133	ACUCGCCGAAUCCCGGGGCCGAGGGAC	1	no match			
7134	GACAGCGAGCCGAACCUUCUUGUCCGC	1	0	coding gene		CG4260-RA|alpha-Adaptin
7135	AAAACCUGAAGAUGAUCCUGAGUCUC	1	1	transposon	LINE	Helena

7136	CCCUUAGAUGUCCUGGGCUGUAACU	1	no match			
7137	UAGGUAAGACGAAUUUAGUCUCUCACUA	1	1	transposon	LTR	G4
7138	GGGCGAAGCUACCAUCUGUGGGAUUU	1	no match			
7139	CGUUCUCUGAAGUUCGACCGGCAC	1	0	transposon	LTR	G4
7140	CACAAACAACACAUACCCCGCUGG	1	no match			
7141	CUUCGUCCUGGGAAACGGGGCGCGGC	1	no match			
7142	UCGAGUGAGACCGGGGGGCAUUCU	1	0	transposon	LINE	G2
7143	ACCGAGGCCCUGUAAUUGGAAUGAGUAC	1	1	ncRNA		rRNA
7144	GCACACGGAGGAGAGCAGAUGAGAGCA	1	no match			
7145	ACGGAACACCCGAGCAAUGCCACCGGC	1	0	transposon	LINE	G2
7146	GAGCGGAGAACCAGAGUUGAUGAGC	1	0	transposon	Low_complexity	F-element
7147	CCAAGAUCAUCAACCCGUGUUAU	1	0	transposon	LINE	Cr1a
7148	CGGUGGGCGAAGAGUUGAGUGGGC	1	0	transposon	Low_complexity	F-element
7149	GAGGAAACGAAAAACCCAUACCGGCACC	1	0	transposon	LTR	HMS-Beagle
7150	UGUACCAAAAAGAGUGCAAUUUAUU	1	0	transposon	LTR	Idefix
7151	CGCAAAGAGAUGGAUGCCGUGAGACA	1	0	transposon	LTR	ninja-Dsim-like
7152	AAAAACCUGAAGAUGAUACUGAGUCGC	1	2	transposon	LINE	Helena
7153	AUAGGGACGGUCGGGGGCAUCAG	1	no match			
7154	UGAGACUACAAACGGAUUGCGGACCGA	1	1	no annotation		
7155	AGGGAAAGGAGCCGAGGCGUCGUA	1	2	transposon	LTR	ninja-Dsim-like
7156	GUCUCGGGAACUUUCAAGGUGCUGGC	1	0	coding gene		CG14690-RA|tomboy20
7157	UCCACCCUGAACGCGCCCGAUCUCGU	1	no match			
7158	UACAAUCUCCGAGCAGACCCCGA	1	no match			
7159	UCGGAAUGCCGGGGCGAGGAGGAGGA	1	0	Su(Ste)		Su(Ste)
7160	ACCGAAAUGAGGUUAGUCGAGUC	1	0	transposon	LTR	gypsy6
7161	CGCUUACUGACGGUGGAUCCCAGAAGAC	1	1	transposon	LTR	rooA
7162	UAGAUCUUCACCGUUGACUAGCC	1	0	Su(Ste)		Su(Ste)
7163	UAAACCGUCAUUCCAGCGCAAGUUGUCU	1	2	transposon	LTR	ninja-Dsim-like
7164	UCUCCGAAAGUUCAUAUGCCUCGGCAUU	1	0	transposon	LTR	flea
7165	GCGCCCGUAGCUCAGCUGGAUAGAGCAC	1	no match			
7166	AAUUCCGAUAACGAACGAGACU	1	0	ncRNA		rRNA
7167	UCAGCGACCGAUAGCGAACCAGUACC	1	no match			
7168	CCCUUGAUAAACGUUUGACUUUCGGC	1	1	transposon	LINE	Rt1b
7169	CGCCCGUUGCGCCCAGUACAAGAA	1	0	coding gene		CG6058-RB|Ald
7170	CGCCAAAUUAUAUCCAUACCCCAUGGG	1	0	transposon	LTR	springer
7171	AACUGAAUAACGAAAUGGAUGACACAC	1	no match			
7172	GUCGGACUCCGAGCUCGAUGGUCAC	1	0	transposon	LINE	R1-element
7173	AGGUGGGGAGUUUGACUGGGGCGGA	1	2	ncRNA		rRNA
7174	AAUCGACCGACCAGCUAUGCGAA	1	0	transposon	LTR	flea
7175	AAGAGGACGUGAAACCGUUAAGAGGUU	1	no match			
7176	UAAUUACCGGUAUAGCACACUUUUG	1	0	coding gene		CG41580-RA|CG41580
7177	UAUUCCGAUAAACGACCAUUACACUU	1	0	transposon	LTR	gypsy3
7178	AACGAGCUCCAGAUGAUGGGGCCAGUUC	1	no match			
7179	UAAGGCCGAGAGCUGAUGACUAGUUGUC	1	no match			
7180	UGAUGGAAGACCGAAACAGUUCUGAC	1	0	ncRNA		rRNA
7181	CAACGGGUAACGGGGAAUCAGGGUU	1	0	ncRNA		rRNA
7182	UGAGCGAACAAACCCAACAUAGUGCAA	1	1	transposon	LTR	McClintock
7183	GACUUGAUCUCGGUGUUCAGUACA	1	0	repeat		LSU-rRNA_Dme
7184	CAUACCAGUAAUAUCCUCUGGUCGAGA	1	0	coding gene		CG3506-RA|vas
7185	CCAGAUGAUAUUCCGAAAACAGCCAAAA	1	no match			
7186	AAAUAGGGACGGUCGGGGGCAUCAGU	1	no match			
7187	UGCCGCGCUAGUUACAUAUAAAAGC	1	0	ncRNA		rRNA
7188	CGGCGCGGCGAAAUUGGCAGGCG	1	no match			
7189	CCAGGAGACACGCUAGGUUGGCC	1	0	transposon	LINE	G2
7190	CUCGCCGAAUCCCCGGGGGCCGAGGGAGCC	1	no match			
7191	CGAGCAGGAACCCAGCGACACCGAGUCCA	1	0	coding gene		CG42390-RB|CG42390
7192	CGGGAUAAGGAUUGGCUCUGAAGAUU	1	0	ncRNA		rRNA
7193	GUGUAACAACUCACCUGCCGAAUGU	1	no match			
7194	UCAGACAAUCCUGCGACUCGAUG	1	0	transposon	LTR	gtwin
7195	UCCUUGAGGAGCUGGAACCUGGUGA	1	0	transposon	LINE	G2
7196	ACGCGCCGCCGCCCCCCCACGCCC	1	no match			
7197	UUGCAAAAACGAGAGCGAUAGAGU	1	1	transposon	LTR	micropia
7198	AAAGACCAUGUGACGACGAGGCCCGUGC	1	no match			
7199	ACACAUCCGAACGAGCUAGGCAACCAGU	1	0	coding gene		CG31882-RA|CG31882
7200	UGACGCCUUGCGUGUUACGGUUGUUUCG	1	0	ncRNA		rRNA
7201	UAUCUUCAGGAUUAUGGUGCUGAAGCU	1	0	ncRNA		rRNA
7202	GUUUGAGAAGCGCAGAGAACGGCCCU	1	0	no annotation		
7203	UUGAACGUGUCCUGAAAGUAGUUGGU	1	0	Su(Ste)		Su(Ste)
7204	AACCCGUGAACCCCCCUUGCCCUCGCA	1	no match			
7205	UUCGAUUUCAAGUACCGGGCGAA	1	0	repeat		RSP
7206	CGCGACGCGAAGUGGCUGGGCACAAGU	1	0	transposon	LINE	G6
7207	UCCGGGAACAUGCCACCUCCGUUGUUG	1	0	coding gene		CG1763-RA|nod
7208	GCAUCCUAGAAUCGACCGACCAGCU	1	0	transposon	LTR	flea
7209	ACGCGGGAGACCGGGGUUCGAUUC	1	1	ncRNA		tRNA
7210	CUGUUUAUCGACCCUGGAAAGCUGUUA	1	0	coding gene		CG4038-RA|CG4038
7211	AACCCUUGAUAAAACGGUUGACUUUCG	1	no match			
7212	AACUCGCCGAAUCCCGGGGCCGGAGGGAGC	1	no match			
7213	AAUGGCCAUCUCUCGUUGAUCUUAAA	1	0	transposon	LTR	opus
7214	UCUCAUCGUCGUAGAACUAGCCCGAGGU	1	1	Su(Ste)		Su(Ste)
7215	CCAACCGGGACAUUUCUCCAACCAGU	1	0	transposon	LTR	micropia
7216	CUCGCCUUGAGAUGAGAUUUCCCGG	1	no match			
7217	GUAACACCCGAUGGCGAAGAGGAA	1	1	coding gene		CG6238-RA|ssh
7218	AAACACAAAAUAAGAAUCGGGGCGC	1	0	transposon	LINE	Cr1a

7219	ACGGACUUGGGGAAUCCGACUGUCUAAU	1	0	ncRNA		rRNA
7220	UGCAGUAGAUUCCUGGCAGAGACGGC	1	0	transposon	LTR	diver2
7221	UCUCAUCGUCGUAGAACAAGCCCGAUU	1	2	Su(Ste)		Su(Ste)
7222	AAAAACCCGAAGAUGAUCCUGAGUCGC	1	2	transposon	LINE	Helena
7223	AGCGUCGAACGAUUUCUGCCCACCUGU	1	1	transposon	LTR	Quasimodo
7224	GCCACAUCUAUCAACAUUCAGCUGGA	1	0	transposon	Low_complexity	F-element
7225	GCCAAUUACAUUAGGCGAGACGGCCC	1	1	transposon	LTR	opus
7226	UACGGUCGACCAACCGCAACCCCUGGU	1	0	transposon	LTR	opus
7227	ACUCGCCGAAUCCCUGGGCCGAGGC	1	no match			
7228	GGAUCCUCAACUGCGAGGGGAUCCUCAA	1	no match			
7229	AAUACCGAGAAGACGUUGAACGCAC	1	1	transposon	LTR	flea
7230	GAGGCCGAGCUGUGACCAUCCGAGC	1	no match			
7231	UACUUGUUCCCCGGAUAGUUUAGUUG	1	1	ncRNA		rRNA
7232	UGGUCUGAGAGGAUGAUCAGUCACA	1	no match			
7233	AGCGCCGGGCCGUACCCAUAUCCGCAGC	1	no match			
7234	AUUGUCAGAAUUGAGUAUAGGGGCGA	1	0	ncRNA		rRNA
7235	GACUCUUUCGUCUAGUGGUUAGGACACC	1	no match			
7236	GUCGCAAAAGACGCGAGAGUGCUUCCGG	1	0	coding gene		CG6184-RA|CG6184
7237	AUGGAACACAUAGGGGCUUCGAGGAAC	1	0	transposon	LTR	HMS-Beagle
7238	ACGGGAGACGCGAGAUCACUACAAG	1	0	coding gene		CG13855-RA|CG13855
7239	GGGGAAUGUCCCGAGCGGCAAAGGGG	1	no match			
7240	UACCGCCCUAGUUCUAACCAUAAAC	1	0	ncRNA		rRNA
7241	CCAGGAGACACGCUAGGUUGGC	1	0	transposon	LINE	G2
7242	ACGACGCGAGAGGUGAAAUUCUUGGUACC	1	2	ncRNA		rRNA
7243	GCGACGACGAGCGAGAUCCGGAUG	1	0	transposon	LTR	ninja-Dsim-like
7244	GGAGUGCCCUUCGUCCUGGGAAACGGGG	1	no match			
7245	AACGAAAUGCGUCGCGACCCCUCGUAUC	1	0	coding gene		CG9641-RA|CG9641
7246	CACUCUAAAAAAAGCUUCGGGACGAGA	1	no match			
7247	CAACGGACCGUCCACCGGUGUUCCC	1	0	coding gene		CG11129-RA|Yp3
7248	GACGUAGCAGCCCGGGCGUCCUCCAG	1	0	transposon	LINE	R1-element
7249	UCGCCGAAUCCCGGGGCCGAGGGGGC	1	no match			
7250	CAAGCCGAGAGGUGACCCAGUGCAU	1	1	transposon	LTR	diver2
7251	GUCACCUACAGGUCCUGCUGCGAA	1	0	transposon	LINE	BS3
7252	ACUCGCCGAAUCCCCGGGGGCCGAGGGAGC	1	no match			
7253	CCGAAAGAUGGUGAACUAUACUUGAUC	1	0	ncRNA		rRNA
7254	UCAACUGUGAGGGGAUCCUCUUAA	1	no match			
7255	AGUUAUGGUUCCUUAGAUCGUUAAC	1	0	ncRNA		rRNA
7256	GGAGGCGGCUCAUCAGUUGCUGUCC	1	0	coding gene		CG31289-RA|Dph5
7257	AGAUCACCCAGACAUCGUGCAGCAGAC	1	1	no annotation		
7258	CAGUUCAUUCUUGAGGCCUUCGA	1	0	transposon	LINE	G2
7259	CCCGUACUCUGUGGUGACCAACACACC	1	0	transposon	LINE	baggins
7260	AACAUGAAAAGUGUAUCCAAGGGA	1	0	transposon	LTR	Dm88
7261	CCUGAGAAACGGCUACCACAUCU	1	0	ncRNA		rRNA
7262	CAAGAAUAACAGGACCAUCCAAGAGU	1	0	no annotation		
7263	GACUAGGGAUUGGCGGCAGUUCAU	1	no match			
7264	AUGGUGCUUACCAAGAUGAAGGAGACC	1	0	coding gene		CG4264-RA|Hsc70-4
7265	AACCAAGCGCGGGUAAACGGCAGGAGU	1	no match			
7266	CAACCGGAGCACCGAAAUAAUUGAUCC	1	1	coding gene		CG32628-RA|CG32628
7267	AACGACGAAAACUAAUAAAGAGCAU	1	0	transposon	LTR	opus
7268	GUCUGGUUAAUUCCGAUAACGAA	1	0	ncRNA		rRNA
7269	GCUGCGGGAUGAACCGAACGCCGG	1	no match			
7270	GAACUUGAGACCCCAGCCGCUCC	1	0	no annotation		
7271	GAUGUCGUAUCACCCAGGACGCCCG	1	0	transposon	LINE	Rt1b
7272	UCAGGUCGGAGCGUGGCGUGAAUGA	1	0	Su(Ste)		Su(Ste)
7273	ACCGGCCGAAUGAACUAGCCCUGAA	1	no match			
7274	CCAAGAUCAUCAACCCGUGUUUA	1	1	transposon	LINE	Cr1a
7275	AUGGCUCUCUCAGUCGCCUCCCGGGA	1	1	ncRNA		rRNA
7276	GAUCGAGCGUCUGGAGGAUGUGGACCA	1	0	coding gene		CG10333-RA|CG10333
7277	GUUACUAUGAUGACGAGUCUCAAGCUG	1	0	transposon	LTR	opus
7278	UGCAUUCGCGUCGAGGCCCAGGAUUG	1	0	transposon	LINE	R1-element
7279	CAGAGUGGUUACAUUAAAGACGAAA	1	0	transposon	LINE	INE-1
7280	AACUGAAUAAACGAAAUGGAUAGACA	1	no match			
7281	CAAGAUAUGACCAAAACCCAGACACC	1	1	transposon	DNA	1360
7282	AGAUCCCCGAAUCCGGAGUGGCGG	1	no match			
7283	GCGAGCACAUGGAUUUACCGAUAAAGUA	1	2	ncRNA		Uhg2-RA
7284	GGAACACGAAACGAAAGCCCCUGU	1	2	transposon	LINE	Rt1a
7285	CCCCCUCCUGUCUAAGGAAUGUCCCGA	1	2	coding gene		CG13591-RA|Ssl
7286	CCCUGAACAAUACACUACACCGCAGC	1	0	transposon	LTR	mdg3
7287	GAGGCCGAGCUGUGACCAUCGAGC	1	0	transposon	LINE	R1-element
7288	GAGAGAAGACUAACCCGACGAGA	1	1	no annotation		
7289	UCCCCUGAUCAAACAUCGUCUGU	1	0	transposon	LINE	Cr1a
7290	CAGAUCGAAAUCGUCACUGCUGA	1	0	transposon	LTR	gypsy
7291	ACCGUUUACACAUUUAUAUUUUGUU	1	2	transposon	LINE	Cr1a
7292	ACUCGCCGAAUCCCGGGGCGAGGGAGC	1	no match			
7293	UAGUACAAUAGAUCCGACUGGGCCAG	1	0	transposon	LINE	baggins
7294	UCUCGCCUCCUCUGGACGUCUUCG	1	0	AT-chX		AT-chX-B
7295	AACCGAGAUUCCCUUAGUAGUGGCGAGC	1	no match			
7296	GCAUUCAUAUUAUAAUUGCCUAUCCCAGA	1	1	transposon	LTR	gypsy
7297	CAAACCGAUGUGCUAUCUUGGCUGA	1	1	transposon	LINE	Doc
7298	ACCCUCCAAGCCUAAAUACUCGUAC	1	no match			
7299	CGUUCCGGAGAGGGAGCCUGAGAAA	1	2	ncRNA		rRNA
7300	AACCAACCGGGAUUGCCUUAGUAAC	1	no match			
7301	AGAGAUACGAUAGAUGCAAAACGA	1	0	transposon	LINE	Circe

7302	GUGCCAGGCUCGUAUAACGUUAAUGAU	1	1	ncRNA		rRNA
7303	AGCCAGGAGGCGAGCCAGGUUAUACCGCA	1	0	coding gene		CG7014-RA|RpS5b
7304	CAGCUCGGGGAGGUAGUGACGA	1	0	ncRNA		rRNA
7305	AUGAGGAGUGUCCCUGCGACUGUGA	1	0	coding gene		CG31948-RA|CG31948
7306	CCAAUGUGGAUUUUCUUGCCCCCGUAGUUGCUCU	1	no match			
7307	AACUUCGGGGUAAGGAUUGGCUCUGA	1	1	ncRNA		rRNA
7308	UGAUACGAGGGAGAAUUGUGAGUCGG	1	0	transposon	LTR	Rt1c
7309	AGUUCGUUCCUGAGUGCGAGCUCCA	1	no match			
7310	CACGAGUUCCUCUGCCACAUGCCC	1	2	Su(Ste)		Su(Ste)
7311	AAACAACACAAUCGACCCCGAAAUUUUU	1	2	transposon	LTR	17.6
7312	AAGCCCAAGAUCGAGGUGACCUA	1	0	coding gene		CG4264-RA|Hsc70-4
7313	CUGCCCAGCGCCUCGGAUGGAAGCCG	1	no match			
7314	CACUGAGUACGGGGCUUACUGAAGUCA	1	1	transposon	LINE	baggins
7315	GGGGUAAAGGCCCACUAAGGCGACGAU	1	no match			
7316	UGAUUACGUCCCUGCCCUUUGUAC	1	0	ncRNA		rRNA
7317	ACCCUGAACAAUACACAACACCGC	1	0	transposon	Unknown	RR50960
7318	GUUGUCCUAUGCACGCCCAUUUGCCCA	1	1	no annotation		
7319	GUAAUUGGAAUGAGUACACUUUAAAU	1	0	ncRNA		rRNA
7320	GUAUCUCGAAGAACGGCCAGACC	1	0	coding gene		CG8280-RA|Ef1alpha48D
7321	UGCCUCAACAUUUGGUGACCUCGACGUU	1	1	no annotation		
7322	AGACCGUAAACGAAAAGCUAAGA	1	0	transposon	LTR	Idefix
7323	AGAGACUUAAAUAUCCCAUCAGUUACC	1	0	transposon	Low_complexity	F-element
7324	AGAGAGAAGACUCUCACCCGACGAGA	1	0	coding gene		CG40609-RA|CG40609
7325	UUAAUUCCGAUAACGAACGAGACUCA	1	0	ncRNA		rRNA
7326	ACCCGGAACAACAAAUGUCGUAAC	1	0	no annotation		
7327	GUUGGUGGAGCGAUUUGUCUGGUU	1	2	ncRNA		rRNA
7328	AAACCUGAAGAUGAUCCUGAGUCGCAU	1	1	transposon	LINE	Helena
7329	GCGUUCGUUGACGUUGAUGGCUGCGG	1	0	transposon	LTR	HMS-Beagle
7330	CAACGAGUUCCUCUGCCGCGUGCCUA	1	0	coding gene		CG40719-RA|CG40719
7331	CGAGCGAACAGUGGUAUGACGCUACGU	1	0	ncRNA		rRNA
7332	ACUGACUAAAUGUCUCAAAAAGGGA	1	0	transposon	LTR	RR50207
7333	UUCUCAACAGGCCGUACCAAUAUC	1	0	ncRNA		rRNA
7334	UGAGACUACAAACGGGAUUGCGGACCC	1	no match			
7335	GACGUCAUUAUUUUUUCCGAAAACGAA	1	0	transposon	LTR	invader3
7336	AAGACCUAUAAACACACUCAACAGAC	1	no match			
7337	ACGGCUACCACAUCUAAGGUAGGCA	1	1	ncRNA		rRNA
7338	UAGAUGAUCUCGGGAACUUACUUAC	1	no match			
7339	AGCGACGUUGUGACCACUGGACU	1	1	coding gene		CG33525-RE|CoRest
7340	UGCUUCUGUUGUGUGGAGUCCUGGGA	1	0	coding gene		CG4986-RA|Mst57Dc
7341	AAAUUCCGUAAACGAAGCCCAAAGUUC	1	1	transposon	LTR	Rt1c
7342	ACUCGCCGAAUCCCGGGGCCAAGGGAG	1	no match			
7343	UUUGUAAGGGUUACAGGGUCAUCAA	1	1	repeat		SAR_DM
7344	AAGGUACAAUGGUAUGCGUAGA	1	0	ncRNA		rRNA
7345	UGCCGGAGCGAGGAGGAGGACUAGCUG	1	0	Su(Ste)		Su(Ste)
7346	UUUCAAAGCUCGACUUAUACAAU	1	0	no annotation		
7347	ACCACCCUGAACGCGACCCGAUCU	1	no match			
7348	AACGAACGAGACCUUAACCUGCU	1	no match			
7349	GCCCAGAGUGGCGGAAUUGGUAGAC	1	no match			
7350	AGCGAUGAAAACGACAAGAGUCUUGAC	1	0	coding gene		CG15734-RA|CG15734
7351	AUCACCCAUAACCCGGGACAAGUUUAC	1	0	transposon	LINE	Doc2
7352	GGUGCGCUUGUGACGGCCCGUGA	1	no match			
7353	CAAGCUAGACUAUGUAUCCCGGG	1	1	transposon	LINE	Doc
7354	AAAAACCAUCUCCGACCGCUAUCCGA	1	1	transposon	LTR	accord
7355	CGUACACAUAUUUCGCAACGUGA	1	0	repeat		HETRP_DM
7356	UGUUUCAUCGUUAGACGACUCGGG	1	1	AT-chX		AT-chX-B
7357	AAUUGAUGAACGUCGUGGAAGACA	1	1	AT-chX		AT-chX-A
7358	AGCGAAACUAAGUGAGCCAAGUGGA	1	1	transposon	LTR	invader4
7359	AAGUACGAAAAGUAGUGACCUUGCGAA	1	0	transposon	LTR	gypsy8
7360	UCUCAUCGUCGUAGAACAAGCCCGCGG	1	1	Su(Ste)		Su(Ste)
7361	AGAUCCCCGAAUCCGGAGUGGCG	1	no match			
7362	AAUCUCGCCUGGGAGUGCUGGACCUGCU	1	1	coding gene		CG2146-RA|didum
7363	UCUUCAACGACAACUCUUUUCCACUCCC	1	0	transposon	LTR	ninja-Dsim-like
7364	UGCUCCCUGAGCCACCACAGGCCCCGUCU	1	no match			
7365	AGCGACGUCGCUUUUUGAUCCUUC	1	0	ncRNA		rRNA
7366	UAAAUACUGAUUUUCUCGAGAUCGA	1	0	transposon	LTR	GATE
7367	CGCGGGCGGCCCGGGUUCGUUUCCCGGC	1	1	ncRNA		tRNA
7368	GAUCCACUGAAGCUUAUCCUUUGCU	1	0	ncRNA		rRNA
7369	AAAACCUGAAGAUGAACCUGAGUC	1	1	transposon	LINE	Helena
7370	CUUGAAAUGAUGAUGACUAAGGUC	1	0	no annotation		
7371	AAACCAUGCAAUGCACCCGACCGAUC	1	0	transposon	LINE	BS
7372	UUAUAUCGCGAGUGCCGUAGAGGUA	1	0	transposon	LTR	invader3
7373	GAAAAGUUUAAACUGAUUGGCAGGA	1	0	no annotation		
7374	AAGAAAAGCAAAAAUUUGACGGCAU	1	0	transposon	LTR	diver2
7375	AGAGACAAUUGACGGUACCGCACAC	1	0	transposon	LTR	Dm88
7376	UAAGGCUAAAUACCGGCACGAGACCGAU	1	no match			
7377	CCAGACGGGAUUACUCUGGAAGGACG	1	0	transposon	LTR	ninja-Dsim-like
7378	AACGCAAUAUUGACAUCAUUCAGC	1	2	transposon	Low_complexity	F-element
7379	AUGAUAUGCACCGAAAGUAUAUGAG	1	0	Su(Ste)		Su(Ste)
7380	UCGGCCGAAUCCCGGGGUCCGAGGAGCGAG	1	no match			
7381	GGACAACUGUGAGGGGACUCCUCAA	1	no match			
7382	CGGACAGAGACGGCGCAUUUUCCAU	1	1	transposon	LTR	McClintock
7383	UAGGUAAGUGGCGGCCAUAAAACGC	1	0	no annotation		
7384	CGGUCUGGUGGUUUGAGCGGUGUGCCC	1	no match			

7385	CUGCGAGAGAUUUGGAGGCAGAAG	1	0	transposon	LTR	ninja-Dsim-like
7386	UGAGCAACCCGGGGGAACUGAA	1	no match			
7387	UAGAUCUUCACCGUUGACUUGGCCC	1	0	Su(Ste)		Su(Ste)
7388	AUGGUGACCACGGGUGACGGGGA	1	no match			
7389	GCCGAGAUUCCCACCCUGCCGUGGUCA	1	1	coding gene		CG11198-RA|CG11198
7390	GGCCGCCUCAGAAUUGCCCAGCGAGUG	1	0	coding gene		CG32486-RD|CG32486
7391	ACUCGCCGAAUCCCGGGGCCGAGGGGAG	1	no match			
7392	UCAAGAAUAACAGGACCAUCCAAGAG	1	0	no annotation		
7393	AACAUUCGAGACCCAACGGCACGUGACAC	1	1	transposon	LINE	Rt1b
7394	UCUCUAUCGUCGUAGAACAGCCCGA	1	no match			
7395	UGACCUAAUGACCCAAGGAAGAUUCCCU	1	no match			
7396	CUGAGAACUUGCUGAUCGAGGCG	1	2	transposon	LTR	ninja-Dsim-like
7397	CACUAACUGGAGGACCGAACCCG	1	no match			
7398	ACCCGAGAUUGAGCAAUAACAGGUCUGU	1	no match			
7399	UUGGGAACACCGCGCGUUGUUGGCCUC	1	1	ncRNA		5SrRNA
7400	ACCAAUCUUAAUCGGGGACUCGCAU	1	0	no annotation		
7401	UCUGCGGCGAAGAUGUAACGGGGCU	1	no match			
7402	AACAUGCAGAUCGUAUGGUCUUGUACCG	1	0	ncRNA		rRNA
7403	AGAACCAUUACGGAAAAAGCUCGAA	1	0	transposon	LINE	Cr1a
7404	AAGAACCUGAAACCGUAUGCGUAC	1	no match			
7405	AUGCUAACUAGUUACGCGACCCCC	1	no match			
7406	UUCGAGAAGCGUUUGAUCGAGUGUGG	1	0	transposon	LINE	BS3
7407	GGAGUGCCCUUCGUCCUGGGAAACGGG	1	no match			
7408	GCUUCCGAUGACCAAUGUUGGCGA	1	0	transposon	LTR	GATE
7409	GCCGAACGUAGUCUCGAGAGCGAGAG	1	no match			
7410	UAUCAUGAUGUUACCCGCAGCGGA	1	0	transposon	LTR	opus
7411	ACUCGCCAAAUCCCGGGGCCGAGGGAG	1	no match			
7412	CAGCUGCUGACCACCCACACGCACGA	1	0	coding gene		CG13255-RA|CG13255
7413	UCUCAUCGUCGUGAACAAGCCCGA	1	no match			
7414	GACCGAGCAGGACCCCGGGCGACACU	1	no match			
7415	GGUAGGGAGUUUGACUGGGGCGGU	1	0	ncRNA		rRNA
7416	GCCCGGGUGGAGCCGCCGCAGGUGC	1	no match			
7417	GUAAAUAAACUCGUUGCUUUGAUUGG	1	0	transposon	LINE	Doc
7418	ACCUGAUUGAUAGACGACGACCCC	1	no match			
7419	CAACACAUGACGCGACAAUUCUGG	1	0	transposon	LINE	INE-1
7420	AGCGACCGCCACAAAAUCGACGACCGA	1	no match			
7421	CCCGCAAAAAUUAAGGCCGUAGUGA	1	0	transposon	LTR	Quasimodo
7422	UGGGACAAUAAACAUGCACUCCAGC	1	1	transposon	LINE	BS3
7423	AGCGGUUCUGACGUGCAAAUUGAUCGU	1	no match			
7424	CGGGGGACAGCCCUUUGCCGC	1	no match			
7425	GGAUCCCUCAACUGUGAGGGGGAUCCUCAA	1	no match			
7426	UCUGACGAGAACUCGAGCCGCGUAG	1	0	no annotation		
7427	ACCCUUGAUAGACGGUUGACUUUC	1	1	transposon	LINE	Rt1b
7428	GCCCGGAUAGCUCAGUCGGUAGAGCAUU	1	0	ncRNA		tRNA
7429	GAUCGGAGAUAUCCGCCCAGCUGGCC	1	1	coding gene		CG6606-RA|Rip11
7430	CGAGCGUGGCCUGAUUGCUAUGCACCGA	1	1	repeat		trf
7431	CGAGAUGACAACCUUGGACCUGACCAG	1	0	transposon	LTR	micropia
7432	GCAGAACUCCGAUGGCGAUAGGA	1	0	no annotation		
7433	AACACUCAGAACGUCUUGAUUCGA	1	0	transposon	LINE	RR49783
7434	CCCUUAGAUGUCCUGGGCUGCACGCGC	1	0	ncRNA		rRNA
7435	ACUCGGAAGCUAAGCAGGGUCGGGC	1	no match			
7436	GGGGAUGCGGCGGGACCGGGGGGG	1	no match			
7437	UGGAAUCCUGAACGUCAGCGCCA	1	0	coding gene		CG18743-RA|Hsp70Ab
7438	CUGUUCACAUGAACUUGGAAUUCCC	1	0	ncRNA		rRNA
7439	UAAAGUCUCAAGGAACAACCCUAAUAGA	1	0	transposon	LTR	Stalker2
7440	GGGAUAGGUAAGUGGCGGCCAUAAG	1	1	no annotation		
7441	AUUCCGAGGAACCCAUUGAUCCAGCUGC	1	0	no annotation		
7442	CAUCCUCUACACCCUGUUUACAGCAAA	1	0	transposon	LINE	G2
7443	AGGUUGCGUGUUCAUAUCACGUCGGGG	1	1	ncRNA		tRNA
7444	AGGUGGGGAGUUUGACUGGGGCGG	1	1	ncRNA		rRNA
7445	UCAAGAGACGCUUGAUAGCCUAGUA	1	2	transposon	LINE	Rt1b
7446	CUGAUAAUGAACCGAACUUUACCUC	1	0	no annotation		
7447	GCAUUCAAUACAGCGAGAUGGUCCA	1	1	transposon	LTR	Quasimodo
7448	CCGAGGAUGACAUGCAGUCUUUG	1	0	transposon	LTR	opus
7449	AACUAACAAGGAUUUUCUUAGUAGC	1	0	ncRNA		rRNA
7450	GAGAGAGGGGCCCGUGCCUUGGAAAGC	1	no match			
7451	GUGACUGACAGGGGAGCAAAUAU	1	0	transposon	DNA	H
7452	AAUGAAACGACAGUGUCCCGCGG	1	0	transposon	LTR	springer
7453	AUGUCUAUGAGGUGACCGUAAUGAGC	1	1	transposon	LTR	ninja-Dsim-like
7454	UGUUUCAUCGUUAGACGUCUCGGGCG	1	1	AT-chX		AT-chX-B
7455	AUGACCGGACGCCCGACGCGCACUGU	1	0	coding gene		CG4750-RA|loopin-1
7456	GUGGAGCCGUACCUGUUGGUUUGUCCC	1	0	ncRNA		rRNA
7457	UCCGGCGAUAAACUUCUUGUCCUUCGU	1	1	transposon	LTR	G4
7458	ACCGAAUUGAAAUUGGAUUACGG	1	0	no annotation		
7459	UGCCCCGAAGGUUGCCCUCUGUGACG	1	0	transposon	LINE	R1-element
7460	AUGGCCCUCUCAGUCGCUUCCCGGGA	1	1	ncRNA		rRNA
7461	AGCUUCAUCCAAGAGUACUGUGCCCGA	1	1	transposon	LINE	R1-element
7462	AAAAACCUGAGAUGAUCCUGAGUC	1	no match			
7463	CGGACUAUCAGAGAAAGCUAUUAA	1	0	transposon	LTR	McClintock
7464	CAUGCCAAAUUCGGACCUAAUGCACGA	1	0	transposon	DNA	1360
7465	CCGGAGAUAAGGUUCAAUGACGGG	1	no match			
7466	CAAGAUAUGACCAAAGCCAAAGACAC	1	0	transposon	LINE	F
7467	CGAACAAGCACCUUCCGACUCAACGU	1	0	transposon	LTR	Rt1c

7468	AGCGAUGAUACCGAAGAACGGGA	1	0	transposon	LTR	Dm88
7469	ACACUCAUAACCGAGCGACUGUUCAAUC	1	1	transposon	LTR	GATE
7470	CCGAAAGAUGGUGAUCUAUGCCUGAA	1	no match			
7471	UGUCAGUUCAAGUCCGACAGCCGGCACCA	1	no match			
7472	AGAACGAUGACGAUCCCACGGCUGGAUU	1	2	transposon	LTR	ninja-Dsim-like
7473	AUGAAAUUCAUGAAGCUGUGGCCG	1	0	transposon	DNA	RR51476
7474	AUCACAUCACGUUCUCACCGUGGAGACC	1	no match			
7475	AGAUCAACAAAGUACGAUGGGGAA	1	0	no annotation		
7476	UCCGGGUAAACCGCUGAACCACU	1	0	ncRNA		rRNA
7477	CAAUUUGUAAAGCUCGAUUGCAAGA	1	1	transposon	LTR	mdg1
7478	AGUUCGGAUCUCGCCGGGGCCUCCA	1	2	ncRNA		tRNA
7479	AACGAAGACAUACACAGAGACCUAA	1	0	transposon	LINE	Doc
7480	GGGUGAGUCGGUCCCUAAGGCGAGGC	1	no match			
7481	CCGCCUACAAUACAGAUACCCACUU	1	2	transposon	LINE	Rt1b
7482	AUGAGAGACGAUGCAGAGCGGGUAG	1	no match			
7483	CAACUGUGAGGAUCCUCAA	1	2	no annotation		
7484	AAGAAGGGCGUGAAACCGUUAAGAGGU	1	no match			
7485	GGUCCUGAGAGCUGGGCGAGCGCCGUUC	1	no match			
7486	ACGGGAAAACAUACCAGGUCCGAA	1	1	ncRNA		rRNA
7487	CCGGCUAGCUCAGUCGGUAGGGCAUG	1	1	ncRNA		tRNA
7488	GCAUCGAGAAUAUUACACAGCCAUU	1	0	transposon	LINE	Doc
7489	AAACGAAUCGAAAACCAGCAGACGU	1	1	transposon	LTR	opus
7490	CUCCCCGCCCCACCCCGCGUCGCGCCGC	1	no match			
7491	UAACACAUCGAUUUCGUGACGUCGC	1	0	transposon	LINE	X
7492	AAAACCUGAAGAUGAGUCCUGAGUCG	1	no match			
7493	AAAUAACUCCAGCAGAGACCGUACCCG	1	no match			
7494	CAUGAUGUUAAUGCCCCUGGUAGU	1	0	transposon	LTR	gypsy
7495	ACCACCCUGAACGCGCCCGAUCCCGU	1	no match			
7496	UACUCGAAGACACCACGUUGAAAGUGG	1	1	transposon	LTR	ninja-Dsim-like
7497	UUCCGAAUGAUCUAGAGCUUAAUUU	1	0	transposon	LTR	diver2
7498	UAAGGGAGCUUGACUGCGAGACCUAC	1	no match			
7499	ACAGUCUAUGAAAGGUGUGUUGCCCAU	1	2	transposon	LTR	GATE
7500	GCGGGACUUACGGGAGAUGCUCAAGA	1	no match			
7501	GAAGAAGAGAUGCCCAGUGAGUUUGCA	1	0	transposon	LTR	ninja-Dsim-like
7502	AUCACAGCCCAACUGCGGACCCAUUAA	1	0	coding gene		CG14034-RB|CG14034
7503	GCGCGAGAAGGAGCUGGGCGAGAA	1	0	coding gene		CG5119-RA|pAbp
7504	UAGACCGAUAAGCUUACUAACAUGUC	1	0	transposon	LINE	Doc
7505	GAGAGUGGACGGGAGCGGCGGGGGCGG	1	no match			
7506	ACCACCCCUGAACGCGCCCGAUCUC	1	no match			
7507	ACCUCCACCUGGACCCAGAUUUCC	1	0	coding gene		CG31206-RA|CG31206
7508	AACCAACGGGAUAUCUCAUACUCGAG	1	2	transposon	LTR	micropia
7509	CUACCAUGGUUGCAACGGGUAACGGGGA	1	0	ncRNA		rRNA
7510	UCCGAAAACAAGGCCCGUCGCUAAU	1	0	transposon	LINE	G6
7511	UCAACGGAUUGGACGACGGCGGC	1	0	no annotation		
7512	AAACGGAAACUUCAUCCCAGUCGGA	1	no match			
7513	CCAACGAAUACCGAGGGUUUUGAC	1	1	transposon	LTR	GATE
7514	AACACUAGACAAUGAUGACCGAAAC	1	0	transposon	LTR	McClintock
7515	GCUCUCGAAUGUAUGCCCGAUAUACGGA	1	0	transposon	LINE	R1-element
7516	GCCGUCAUUACAUUUUUUUGAGCUGC	1	0	transposon	LINE	TART-A
7517	UGCUACUCGAAUGCUUGCCCGAUAUA	1	no match			
7518	AGAUCAGACGUGGCGACCCGCUGAAUUU	1	no match			
7519	CUUGGACUACAUAUGGUUGAGGGGUUGUA	1	no match			
7520	CGUACCAAAAAACCCGAUGCUUCUGGC	1	0	transposon	LTR	297
7521	CGCGAGAGGUGAAAUUCUUGGACCG	1	0	ncRNA		rRNA
7522	AAGGUGUGACGACAAUUGCAUCGGUU	1	no match			
7523	UCUCCGGGAGACACCAAUGCUGGCCAAG	1	0	coding gene		pseudogene
7524	AAAAUUCAAAGACGACGGCCCACC	1	2	transposon	LTR	springer
7525	UUUGGGCUCCGGGGGAAGUAUGGU	1	0	ncRNA		rRNA
7526	GACAUUUGCGAGGUAUACCGGGAU	1	0	coding gene		CG34170-RA|CG34170
7527	UAGGAACGAAAGAGGCGAGGACAA	1	0	coding gene		CG40271-RA|CG40271
7528	UGAGACUACAGACGGAUUGCGGUACC	1	no match			
7529	UGCACCAAGAACUGAAGGCCAUU	1	0	transposon	LTR	invader1
7530	ACCUUAUGGGACAUGUGCUUUUAU	1	0	ncRNA		rRNA
7531	GGCUUUGAGAGGCGCCGUAGUUAAA	1	2	no annotation		
7532	UAUUGAUUCCGUGACGCGUCUGCU	1	0	coding gene		CG8728-RA|CG8728
7533	UUCCAGACGACUAGGUGGAGCUUUAA	1	0	transposon	LINE	G6
7534	AACUGGGAACGUCGAUGGCCUGGCC	1	1	coding gene		CG5060-RB|CG5060
7535	AGUGCCCAAAAACUAUAAGUUGACCAA	1	0	repeat		RSP
7536	GAGAGAAUGAUAAGAACCAUUACGG	1	0	transposon	LINE	Cr1a
7537	GAGUCCAUGUCCUCCUAGCCUGC	1	0	transposon	LINE	G2
7538	GCAGACCCAGAAAAUCCCAGACC	1	no match			
7539	UGUUGAGAUGUGACGAAUCUGGCCCA	1	0	transposon	LTR	Quasimodo
7540	CCCGAAAGCAGAUGAGGUAACGUAC	1	0	transposon	LINE	F
7541	UGAUGACCGAAAUUUUGAAAAACG	1	0	repeat		SAR_DM
7542	UAAUUCCGAUAACGAACGAGACU	1	0	ncRNA		rRNA
7543	GGCUUUGAGAGGCGCCGUAGUUACCUAC	1	2	no annotation		
7544	UCAUCGAGAAUCUCACGCCCACAGGC	1	1	transposon	LINE	Doc3
7545	UGUUGACAAAAAUGAGGCUGACCA	1	0	transposon	LTR	roo
7546	UCCAAUAUCACCGCGGAGACAGUCGC	1	1	transposon	LINE	Doc4
7547	UUCAACCAGAUGGGCCUGGAGUACUUC	1	0	Su(Ste)		Su(Ste)
7548	GAGUGCUUGAUAUGACCAGCUUGG	1	0	Su(Ste)		Su(Ste)
7549	UAAAAGGGAAACAACCCAGACCGUC	1	no match			
7550	UGACAUAUGAGACUACACCAGAAA	1	0	transposon	LTR	Idefix

7551	CCCGAAAGCAGAUGAGGUAGCGUAC	1	0	transposon	LINE	FW
7552	UCAACGAGAAUCAUCCAUCAUCCAGA	1	0	transposon	LINE	G2
7553	UAACGACCCAUUUCUGUUACUUUGUGU	1	0	repeat		trf
7554	UCCAAACCCGAAUCCCAACAGUCCCGC	1	0	coding gene		CG7417-RA|Tab2
7555	CCCCAAUCAAUGAUUACAGCUGAGGCCC	1	0	transposon	LINE	Doc3
7556	AGGAGCUGGGCCGACUGCUGCUGGAGA	1	0	coding gene		CG7061-RA|rab3-GAP
7557	GUCUCAGAGAGCAAAUCGGCAAGAC	1	0	transposon	LTR	springer
7558	GCUGCACGAAAUGAACCCGAAAGC	1	0	transposon	LINE	Doc3
7559	UUCGAGAACAAACGAUACUGUUCAAUC	1	2	transposon	LINE	G2
7560	GAGAAUCGCAUCUUAGAGUAAAUGGCC	1	0	transposon	LTR	gypsy8
7561	AAUUCCAUGAAAAAGACAUUGGCCAU	1	0	coding gene		CG41322-RA|CG41322
7562	AAGUAUUGGCAUCUACCUAGUUU	1	0	transposon	LTR	Idefix
7563	GCAAGACGGACAAACGGGUGGGGU	1	no match			
7564	AAGGCCGAGAGCUGAUGACGAGUGUU	1	no match			
7565	UGCUAACGAAACAGUAACACUAGGUC	1	2	transposon	LTR	Idefix
7566	ACGGAAUGGAAUCGAGCGCUGAGUAC	1	1	transposon	LINE	X
7567	UGUAUCCAAAAAUCAUCUGUGCGAGU	1	0	no annotation		
7568	CACUAUCAGACAGCAACAUUGACCC	1	no match			
7569	AGAGCUCCGAUUUACCCAUGCAAC	1	0	coding gene		CG3620-RA|norpA
7570	CUGGCGAAUAAAAGUAAACGAGC	1	0	transposon	Low_complexity	F-element
7571	AAUGGCUGCAACGGGGUAACGGGGGAAUC	1	no match			
7572	AAGUUAACAACCGAAACUUAUGGCAC	1	0	transposon	LTR	rooA
7573	UGAGACUACAAGCGGAUUGCGGACC	1	1	no annotation		
7574	GCAUCCGGUGCGUCCUUAACAGUCCUU	1	no match			
7575	AUGGCUCUACUCAGUCGCUUCCCGGGA	1	no match			
7576	ACAUUCUUAACCGAGCGACUGUUCAAU	1	1	transposon	LTR	GATE
7577	CUCCCGGGGGAAACAUGGUCAAGUU	1	0	transposon	Low_complexity	F-element
7578	ACCCACCCCUGAACGCGCCCGAUCUCGUC	1	no match			
7579	UCCGAAAACGUAAUGCUGCUUUAUCCA	1	0	transposon	DNA	S
7580	UCCUGGCAGAGACGGCGUCGCUCGAG	1	0	transposon	LTR	Quasimodo
7581	UUGGAAAUUAACCAAAAACAGCAUGG	1	0	Su(Ste)		Su(Ste)
7582	UACUGAGAUCAUACGAGAGCUGAAA	1	0	transposon	LTR	flea
7583	AAAACCUGAAGAUGAUCCUGAGUCA	1	1	transposon	LINE	Helena
7584	ACCACUUUCAUGCUUGGGAUUGUGAA	1	0	ncRNA		rRNA
7585	AUGGCUCUCUCAGUUCGCUUCCCGGGAA	1	no match			
7586	ACGGAAUGGAACCGAGCUCUGAGUA	1	0	transposon	LINE	X
7587	GUCAUGUGCAUCCUGGUGGACCCCAGCA	1	0	coding gene		CG18810-RA|CG18810
7588	CAUCGAAGGAACGAUCGCAAGGCAGCA	1	0	transposon	LTR	flea
7589	CAAUGCAUCUUGAACAUCGUCCC	1	no match			
7590	AUGCCGGAGCGAGGAGGAGGACUAGU	1	1	Su(Ste)		Su(Ste)
7591	UCCCCUGGACCUGUACUGACGCUCAU	1	no match			
7592	UGGCGGAGGCGGUCGGCGGGGGG	1	no match			
7593	CCAAAGUUGAUCCGAUCCAACAA	1	0	transposon	LTR	3S18
7594	AAGAACAAAAAGAGGACUUCGAGUA	1	0	no annotation		
7595	CUUAGUUGGUGGAGCGAUUUGCCUGGU	1	no match			
7596	UAUCUCAAGACGUGUUUGACCCCAGC	1	0	transposon	LTR	1731
7597	GACAACCAGAGAUCUAGACGGCCCAA	1	1	no annotation		
7598	UCUGUUUCAUCGUUAGACGGCUCGGG	1	0	AT-chX		AT-chX-B
7599	AACCAAUCCCAACCAAACCCGUCAA	1	0	transposon	LINE	G2
7600	CCGAAUACGAGGAACAGGCAGAAUCCAA	1	1	transposon	LTR	gypsy4
7601	AAAUAACUCCAGCAGAGACCGUACCC	1	no match			
7602	AACACUGCCGAGGAUUAUUGAAGUC	1	0	coding gene		CG1503-RA|CG1503
7603	GGUCCUGAGAGGAUGGGCGAGCGCCGUUCC	1	no match			
7604	UGGCAAACGAGAGCUCUUCCGGGU	1	0	transposon	LINE	R1-element
7605	CGCUCUUCACCACAGAGUCUCCGAGGAGU	1	0	transposon	LTR	GATE
7606	AAGAGAUGGAAAGCACGCAUCGA	1	0	transposon	LTR	springer
7607	ACAAACAGAAUUCCCGAAUGAGAC	1	1	transposon	LINE	Cr1a
7608	AUUCACCGAGCGUUGGAUUGUUCACCC	1	1	ncRNA		rRNA
7609	UGAGCGAACAGACCCAAUAUAACGCA	1	0	transposon	LTR	McClintock
7610	UCAACUUUGAGGGGAUCCUCAA	1	no match			
7611	UUUGCAAAACGGGAGCGAUAGAGC	1	no match			
7612	ACCCGGUUCAUCACUCUAGGACGAA	1	0	transposon	LTR	297
7613	UUCAACCGGACGGGUCUAGAGUUU	1	1	no annotation		
7614	UGGUUGCACGGGUAACGGGGAAUC	1	no match			
7615	UACGGUGCCCAAAUUAACAACUCA	1	0	ncRNA		rRNA
7616	GGGUAGUACAAUAGAUCCGACUGG	1	0	transposon	LINE	baggins
7617	CUUAAAGGAAUUGACGAAAGGG	1	1	ncRNA		rRNA
7618	AAAACCAAGCUUUUGGGCUCCGGGG	1	2	ncRNA		rRNA
7619	CCGAUGAACCUGAAUAUCCGUUAUGG	1	0	ncRNA		rRNA
7620	CAAUGUGAUUUCUGCCCAGUGCUC	1	0	ncRNA		rRNA
7621	GGCAGCGACGACCCAUUCGAACGUC	1	no match			
7622	UCGAAUUCAACGAGGUAUGGCAUU	1	1	repeat		SAR_DM
7623	AGAGUGCGAAGAGCGAUGAUACCGA	1	0	transposon	LTR	Dm88
7624	AAACCAUUCAUAGACGACCUGCUU	1	no match			
7625	GAAUUGACGGAAGGGCACCACCAGGAG	1	0	ncRNA		rRNA
7626	GGAGACAAGGUUCACCCGCUUCCAA	1	1	transposon	LINE	Rt1b
7627	GCGGAAAGAGAGCGAUUUUGGC	1	0	transposon	DNA	1360
7628	UGAAAACGAAGGUGACCUUUUUGGGU	1	0	transposon	LTR	flea
7629	CAAGCCUGAGAGGUGAUGCGUAGCC	1	no match			
7630	UCAAAUAAUAACGGAGGUGUCCCAA	1	0	ncRNA		rRNA
7631	GUGAUCGAUACGCCCCCUCAGAG	1	0	transposon	LTR	Max
7632	AAACCUUCGUAGACGACCUGCUUC	1	no match			
7633	ACGUACGCGAAACGUGAAACCUGC	1	0	coding gene		CG1763-RA|nod

7634	UGAUUCCGCCUCCAGUUUCCUGGGCCC	1	0	coding gene		CG16817-RA|CG16817
7635	UUACAAAGGAUGGACUGAGAACC	1	0	transposon	LINE	Cr1a
7636	AUUGCAACGAACUGACGGACUACU	1	0	transposon	LTR	HMS-Beagle2265
7637	ACUACGCCGAAUCCCGGGGCCGAGGGAGC	1	no match			
7638	AUAACUAGUAGCGUACCAUGUGAC	1	0	transposon	LTR	gypsy6
7639	GUGGAGCGGGCGUGGAAUGCGAGUG	1	no match			
7640	GCGGAGCGAGCGCACGGGGUCGGCGGC	1	no match			
7641	AGGGAGCGAGAGAGCGGAGAGCGC	1	0	transposon	DNA	1360
7642	UGUCAGAAUUGAGUAUAGGGGCGA	1	0	ncRNA		rRNA
7643	GGAUAACUGUGAGGGGAUCCUCAA	1	no match			
7644	CAUAUCCGAGGCCCUGUAAUUGG	1	0	ncRNA		rRNA
7645	UGCUCUCGAAUGCUUUCCCGAUAUA	1	1	transposon	LINE	R1-element
7646	AUUCCGAUAACGAACGAGACUCAAAU	1	0	ncRNA		rRNA
7647	ACCCUACUGAUGAUGUGUUGUUGCCAU	1	no match			
7648	AGGUUAGUUUUACCCUACUGAUGAUGU	1	no match			
7649	CGUAACUCAAACGUCGAGCAUGAAG	1	0	transposon	LINE	R1
7650	CUAACUACCGUGACAGUGAUCGUG	1	0	no annotation		
7651	GCCCAACCUCGAGACUUAUAGACU	1	0	transposon	LTR	gypsy4
7652	AAACGUAUCCACAUUCGCUGAUGACAC	1	no match			
7653	CAAUAACAGGUCUGUGAUGCCCUUAGA	1	0	ncRNA		rRNA
7654	UGGCCACACUGUCUCCUCCUGAGUA	1	no match			
7655	GUUUCCAUGAGCGAGCGGGACGU	1	0	coding gene		CG14690-RA|tomboy20
7656	GAUCGUCCUGAAGCAAUCCCGUCUCU	1	0	coding gene		CG13394-RA|CG13394
7657	AAGAGCUAUAAAGAGUUUUUUGCCG	1	0	no annotation		
7658	CACGGGAAAAACUUACCAGGUCCGAA	1	no match			
7659	AAAACGAGAACUUUGAAGGCCGAAGU	1	no match			
7660	CCGAACUUGAGACCGCCCCUGGACU	1	1	Su(Ste)		Su(Ste)
7661	UCCUACGCCCUCUUCGCCAUCGCCACCA	1	0	coding gene		CG5554-RA|CG5554
7662	CGGAACGAAACGGCCUUUUUCUAAU	1	no match			
7663	CAGGAUUGUAACUCGCGCGAUGGGC	1	0	ncRNA		pseudogene
7664	AAGACAAUUAUCCGAUACCGCUUGU	1	1	transposon	LTR	flea
7665	CGCCUCGGACGAACACCAUUGGGGGGUAG	1	no match			
7666	CAAGCUAGACUAUGUGUCCCCGGGU	1	1	transposon	LINE	Doc
7667	UUUGCAAAAAACGAGAGCGAUAGA	1	no match			
7668	ACGGACCCAGCUGAGCGAACAGUAUG	1	0	coding gene		CG10859-RA|CG10859
7669	UACCAAUCCCAAUUAGGCGAACCGA	1	0	transposon	LINE	I
7670	GUCGUCAACGAGAGAUGCUGUCCUACCA	1	0	coding gene		CG6995-RA|CG6995
7671	AGCGCCGAGAACUUGAACCCGAACGCAC	1	1	coding gene		CG4528-RA|snf
7672	AUUGGGUGUAGCUACUUUUAUGGC	1	0	ncRNA		rRNA
7673	GUACGGCAUGCCUGAGGUGGAUGA	1	0	coding gene		CG6259-RA|CG6259
7674	AUUGAUCCAGCUGCUUUUGUUGU	1	0	no annotation		
7675	AAAACCUGAAGAUGAUCCUGAGUCGCCU	1	2	transposon	LINE	Helena
7676	GCCGAAAAUACGACGACCCCAAGAUC	1	0	coding gene		CG18743-RA|Hsp70Ab
7677	AAUCAUGCCCGACGGGGCGCUGACCCCCC	1	no match			
7678	CGCUUUGAAACCGAAGAGACGCCGG	1	0	transposon	LTR	G4
7679	CCCAGAUGCCUCAGGUUGCACCCAU	1	0	transposon	LINE	R1
7680	GAAUCUAUGGGUGGUGGUGCAUGGCCG	1	0	ncRNA		rRNA
7681	GUGGCUAACACAAAUCUCGUACCAGGG	1	0	transposon	LTR	G4
7682	UGCCUCAACAUUUGGUGACCUCGA	1	0	no annotation		
7683	UUAACAACGACAGGGGUACGUCAU	1	0	transposon	LINE	Rt1b
7684	GAAUGUCCGGACAGCUGGGCCCGA	1	1	Su(Ste)		Su(Ste)
7685	AUGGUUGCAACGGUAACGGGGA	1	2	no annotation		
7686	AGAGCCCAGCGCCGAAUCCCCGCCCCGC	1	no match			
7687	AACCCAGACCGUCAGCUAAGGUCCCAA	1	no match			
7688	UUGUAAAUUUGACCCCCUUCUUAGACU	1	0	transposon	DNA	1360
7689	ACCCUGAACUCGCCCGAUCUCGUC	1	no match			
7690	AGAGACGAGAACGCGUAUGUGUGCGUGA	1	1	transposon	DNA	FB
7691	GCCGAACGUAGUCUCUCGGAGAGCGA	1	no match			
7692	UUUGCAAAAACGAGGGCGAUAGAGC	1	1	transposon	LTR	micropia
7693	UUCGAGAAGCGAACCCGGCGAACU	1	no match			
7694	AAGCCACUCGAGCAGAAGGCCUCCGA	1	2	coding gene		CG32236-RA|CG32236
7695	GUGGCUUUGAGAGGCGCCGUAGUUAC	1	0	no annotation		
7696	GAUUCGAAAUCAGAUUCCCUUUGGG	1	2	ncRNA		tRNA
7697	GAUCACGAGUUAGUCGGUCCUAAGUU	1	0	ncRNA		rRNA
7698	AGACAACCAGAGAUCUAGAGGGCCCAA	1	0	no annotation		
7699	GUUAGACGAAGAGACCCCGUGCA	1	no match			
7700	AGUUUCAUCGUUAGACGGCUCGGG	1	1	AT-chX		AT-chX-B
7701	UCGUUAUAUUGGUUGAACUCUAGAU	1	0	ncRNA		rRNA
7702	AAAAAGAUGAAGAAGACGUCGACG	1	0	no annotation		
7703	UCGGGAUAAGGAUUGGCUCUGAAA	1	1	ncRNA		rRNA
7704	UAAGGAUGAUAAGGAUGAUGAUACC	1	no match			
7705	ACAUUCAUAACCGAGCGACUGUACAAU	1	1	transposon	LTR	GATE
7706	UGUUUCAUCGUUAGUCGGCUCGGG	1	1	AT-chX		AT-chX-B
7707	UAUUCUGUGGAAGGGGGACCGGGAC	1	no match			
7708	CGAAAAUGCAGACCUAUUGUCAUCCGG	1	0	transposon	LTR	flea
7709	ACUCGCCGAAUACCGGGGCCGGGGGA	1	no match			
7710	UGACAUCAUUCAGCGAGCACAAUCAAU	1	1	transposon	Low_complexity	F-element
7711	AAAUAAUACAAACAGAGUUCCCGAAUGA	1	0	transposon	LINE	Cr1a
7712	GAAGAGAGAUGCCCAGUGAGUUU	1	2	transposon	LTR	ninja-Dsim-like
7713	UUGCACGAAAUGUCGUUACCGAGAA	1	0	transposon	LTR	copia
7714	AGAGAGCAUGUUGUUGCGGGAUCCCC	1	0	coding gene		CG8712-RA|CG8712
7715	UGCGAGAUGAACCUUUAGAGCACGC	1	1	transposon	LTR	GATE
7716	ACACGGGAAACUUACCAGGUCCGAAC	1	no match			

7717	UUGAUUUCCAGGGACUUUCUCCAA	1	0	transposon	LTR	Rt1c
7718	ACCCUUGAUAAACGGUUGGCUUUC	1	1	transposon	LINE	Rt1b
7719	AGAAGCCGUCGAGAGAUAUCGGA	1	0	ncRNA		rRNA
7720	GAGAGGCGCCGUAGUUACCACCCCA	1	0	no annotation		
7721	AAAAUCCCCCGGCUGCGACCUUAUAA	1	0	transposon	Low_complexity	F-element
7722	GCGGAAAGACGGAAAGACCCCAUGAACC	1	no match			
7723	GGUGCGAGAGGUCCCGGGUUCAAAUCC	1	0	ncRNA		tRNA
7724	CGGGAUAAGGAUUGGCUCUGAAA	1	1	ncRNA		rRNA
7725	AAGAUUAUAAAGCCGAGUGUGUCG	1	0	transposon	LTR	diver2
7726	UCUCAUCGUCGUAGAUCAAGCCCGA	1	1	Su(Ste)		Su(Ste)
7727	AGCGAAUGGCAUCCGGAUGAGAAAU	1	1	transposon	LTR	flea
7728	ACAACACAUCGAGCGACCGCGUAGUC	1	0	transposon	LINE	Doc3
7729	CAAACUAUCAUGGAAUUGCGCCGGU	1	0	transposon	LINE	Doc4
7730	CAACGGGAUACCUCAUACUCGAGUA	1	0	transposon	LTR	micropia
7731	GGGUGUUGACGCGAUGUGAUUU	1	2	ncRNA		rRNA
7732	UACCCCGGAAGUGAUCAUCGAG	1	0	transposon	LINE	F
7733	GAUUCCGGAGAGGGGAGCCUGAGA	1	no match			
7734	CCGUACGCUCGAUGAUCGUCCCAGACU	1	no match			
7735	CAACGUGCGAGAGGUGCAGCCAAU	1	0	transposon	LTR	ninja-Dsim-like
7736	CAACCUAUCCAUCAACCCAGACGAG	1	0	coding gene		CG18743-RA|Hsp70Ab
7737	CUUGUACCGACGACAGAUCUUUC	1	0	ncRNA		rRNA
7738	UCAACGUGUGAGGGGAUCCUCAA	1	no match			
7739	GAUACUCGAAGACACCACGGUGAAAGU	1	0	transposon	LTR	ninja-Dsim-like
7740	CGUCGAAUUUGGGUAUAGGGGCGAA	1	no match			
7741	CGCGACCUCAGAUCAGACGUGGCGACCC	1	no match			
7742	CAAGAGCGUACCCAACUCUUAUGGG	1	0	transposon	LINE	accord2
7743	UAAACCUCAUCCGCGUGGCAUCU	1	0	no annotation		
7744	AACCGAUAUGGUUGGACCACAAUCCUU	1	1	ncRNA		7SLRNA-RA
7745	AGAAACCUCCUGUGGAGCAGAAGGGC	1	no match			
7746	AUAAGACCUCUGUUCUGCUUUCAUUG	1	0	ncRNA		rRNA
7747	GGCUUUUGAGAGGCGCCGUAGUUAA	1	no match			
7748	UCGGCCAGCGUGGAGAUAUUGGGU	1	0	coding gene		CG7762-RA|Rpn1
7749	GCAUCAGUUAAAGGCGAAUCGCCCGA	1	0	transposon	LINE	RR49783
7750	UGCCGCGUGAGUGAUGAAGGCCCUAGG	1	no match			
7751	UGCUCCAGCUGCUCCUCUUGUCCGC	1	0	coding gene		CG30460-RA|CG30460
7752	AGGUAACUAGGAACCGUACCCCAAACCGA	1	no match			
7753	UUCCGGAGAGGGAGCCUGAGUAAAC	1	no match			
7754	CAGCCCGUGGACGGUGUGAGGCCGGU	1	no match			
7755	UAACCAAGGACGUUCACACGAGGG	1	0	transposon	LTR	opus
7756	UCGGGAUAAGGAUUGGCUCUGG	1	1	ncRNA		rRNA
7757	CCCCAGACACCGACCCGUGAAUACCCC	1	0	no annotation		
7758	UGAGCUCCAAACGGACAUGACCGACC	1	0	coding gene		pseudogene
7759	CGAACAAUGAGGCCGUCACCCAAA	1	0	transposon	LINE	Rt1a
7760	UAGUACUAAUAGCCUGAUUGAUU	1	no match			
7761	UAAUGGUAUCCGAGUCGUACUCUG	1	1	transposon	LTR	springer
7762	UCCGAAGAUAACGAGGAGCAGAGU	1	1	transposon	LTR	297
7763	GAUACUGAAACGAGUUGGCAUUCUU	1	0	transposon	DNA	RR51476
7764	UGUUUCAUCGUUAGACGGCGCGGG	1	1	AT-chX		AT-chX-B
7765	CACUUACUGACGAGAUCGGCGCGA	1	0	transposon	DNA	Ivk
7766	UUGAAGGCCGAAGUGGAGAAGGGUUCC	1	no match			
7767	GAAGGCCCGCGGCGGGUGUUGACGC	1	no match			
7768	GGAUCUCAACUGUGAGGGGAUCCUCAG	1	no match			
7769	GACCGGCUCCGGGACGGCUGGGAAGGCCCG	1	no match			
7770	AGUCGCGAAAUUGGUAUCCCCGAU	1	0	transposon	LTR	diver2
7771	AGCGAGAUAAAGCGAAAACCGUCAAA	1	1	transposon	LINE	G6
7772	CCAGUCGUAGUCCGGAUCGCAGUCUG	1	no match			
7773	AAAGCAAGACAGAGUUCACCAUG	1	2	no annotation		
7774	AUAUUCCCCAUGAACGAGGAAUUCCCAGU	1	no match			
7775	CGACACAAAGCCGCUCCAUCAGCCAGC	1	0	coding gene		CG4027-RA|Act5C
7776	UACGAGAAUAUACCGCAAACCACAGUC	1	0	transposon	LTR	gypsy12
7777	AAGGGAAAUAAAGAGAAACUCGAAUA	1	0	coding gene		CG40124-RA|CG40124
7778	CUGUAACAACUCACCUGCCGAAGC	1	1	ncRNA		rRNA
7779	UCAACUACGAAACUCAAACUCCCGGACA	1	0	no annotation		
7780	CCACCCGAAAUGCGCGCUUUUUA	1	0	transposon	DNA	1360
7781	UGAACCGAGAAGUUAAGCCGACGAAUCA	1	0	transposon	LINE	Cr1a
7782	CACCGAGAUGACGCAACACGAAC	1	no match			
7783	ACCACCCUGAACGCGUCCGAUCUCGUC	1	no match			
7784	GUAUUCGAGACCCAACGGCACGUGAC	1	1	transposon	LINE	INE-1
7785	UCUGGGUCGGGGUUUCGUACGUAG	1	no match			
7786	CCCUGGGUUUUUUCCAGCUCGCCCCCCA	1	no match			
7787	AACAACUAACUGGAGGACCGAACCGACU	1	no match			
7788	CAGUCGCGAAAUUGGUAUCCCCGG	1	1	transposon	LTR	diver2
7789	ACCCUACUGAUAGAUGUGUUGUUGCC	1	no match			
7790	AAAAACCGAUGCGAUCCGGAUU	1	no match			
7791	UUGAUAUGGCCAGCUACCACCCAC	1	1	Su(Ste)		Su(Ste)
7792	AAAACCAAUAUGGACGUAUGGAACGGU	1	2	transposon	LINE	Porto1
7793	AACUACUUCAGGGACCGUGUGUUA	1	0	transposon	LINE	Rt1a
7794	UCUGCGAGAAAACGUACGCAUCGG	1	0	transposon	LTR	3S18
7795	CGACUCUUAGCGGUGGAUCACUCGGC	1	no match			
7796	AGCAAUAUGACAGCGUGAUCCUGG	1	1	transposon	LTR	Rt1c
7797	UAGUCCCGAAGAAACCUAAACCA	1	0	transposon	LTR	opus
7798	CGAGGGUGCAAGCUAGACAUUAAAA	1	1	transposon	LINE	R1-element
7799	GCUCUCGAAUGUAUGCCCGAUA	1	0	transposon	LINE	R1-element

7800	CCCAGCUCGGGGGAGGUAGUGACGA	1	no match			
7801	GUGGAGCCGGGCUGGAAUGCGAGUG	1	no match			
7802	UGAUUCAUCGUUAGACGGCUCGGG	1	1	AT-chX		AT-chX-B
7803	ACGGACGGACGGACGGCGAGGCC	1	0	transposon	LTR	micropia
7804	GAGGUAACGAUGGAGUUUUGACGGUC	1	no match			
7805	CACAACAACAUCCCGGGAAAUGAGC	1	0	transposon	LINE	Fw2
7806	GAAGAUAUGAAACCGACUCAGGUAG	1	0	transposon	LTR	Rt1c
7807	GGAUCCCAACUGUGAGGUAGGUUGUAU	1	no match			
7808	CCGCGUCCCAUGCUAAACUGUGGCCAG	1	0	ncRNA		snoRNA
7809	UUCGGAAUGCCGGGGCGAGGAGGA	1	0	Su(Ste)		Su(Ste)
7810	GGUGCGCCGCGACCGGCUCCGGGACG	1	no match			
7811	AGCACAAGACCGAAGCUGAUGACUGAUG	1	no match			
7812	AUAAAUCAAAACCAGUAUUCACCGCGUU	1	0	transposon	LTR	Idefix
7813	AUUCGUGAUGGGGAUCGGGGAUU	1	no match			
7814	GUGUGCUAUACCGGUAAUUACCGCUGG	1	0	no annotation		
7815	CAACCCCGCCAGUGACGUAUGCAA	1	0	transposon	LINE	Doc
7816	UGCGAUUCCGGCGCAGGUGACUCC	1	0	ncRNA		pseudogene
7817	UCAAAUGAAAUGGACUCAGAGUC	1	no match			
7818	GGCAUAAAGAGACAAUUGACUGUACC	1	1	transposon	LTR	Dm88
7819	AUGCAAGGGAACGUGAGCUGGGU	1	0	ncRNA		rRNA
7820	UAGAUGUGGAAACCAGGGGCAUAUA	1	0	transposon	LTR	invader6
7821	UAACAAGAUGCAUACGAUGUGGCUCAA	1	1	transposon	DNA	1360
7822	CAGUUCAUCGGAGUGCUCGGAGG	1	0	coding gene		CG33957-RB|cp309
7823	AUGAUUAGAGGUUCCGGGGUCGAAAU	1	no match			
7824	UGCUAGAAGACGAUUUUCGGGCGGA	1	0	transposon	DNA	1360
7825	UGAGACUACAAACGGAUUGCGGACA	1	1	no annotation		
7826	GGUUCCGGGGGGAUAUGGUUGC	1	no match			
7827	GCACUGCGAGCGUUCGAGUCGUUAACGA	1	0	coding gene		CG2727-RA|emp
7828	CACAUCAACAGGAAGCGUAUCCAGGCC	1	0	transposon	LINE	G2
7829	AGUCCCCAGGAUUUUGACGCGACU	1	0	transposon	LINE	BS3
7830	UACAGAGAAAUGGCAGAACACGCUGU	1	0	transposon	LTR	micropia
7831	GCAGUUAUUACGACGAGAAGAUGGCC	1	1	transposon	Unknown	RR50960
7832	CCUACUAAUACACCUACAUACACCUGCC	1	0	Su(Ste)		Su(Ste)
7833	UCCGGACAAACACACAAGACCCGAUA	1	1	transposon	LTR	opus
7834	AAAGAGAGUAAUUUUGGCCGUCAAC	1	0	transposon	DNA	1360
7835	GAACUGGGAAGGUGGGACUUGAGUU	1	0	transposon	LINE	G2
7836	CGAAUAGUGAUUGCGAACCCCCGAAUC	1	0	transposon	DNA	S
7837	GAUCAAGAAUCCCGUGCCAAGA	1	0	transposon	LINE	Circe
7838	UGGCCGAAGUAUUCAACCGCAUUUGCG	1	1	coding gene		CG5757-RA|CG5757
7839	GCUGAACCACUUUCAUGCUUGGGAUU	1	0	ncRNA		rRNA
7840	GUUCGAGCCCAAGUGCGUCCACC	1	no match			
7841	UCCGGUCCCGGGACAUGGCGUGGUGGGUA	1	no match			
7842	UUUCGAACCAGAGGCUGGGUACAUCGC	1	0	no annotation		
7843	ACCGAUAUCCCCCAGGCAAAUGAU	1	0	transposon	LINE	INE-1
7844	CACUGCAAUACUAAGUCGACCCAGG	1	2	transposon	LINE	Doc3
7845	AGCAGUCGUGGCCGAGCGGUUAAGG	1	0	ncRNA		tRNA
7846	GUCAUGCGAGGCGAACCGGGUGA	1	no match			
7847	CUCAACUGUGAGGAGGUAGUAGGUUGUAU	1	no match			
7848	GACCGGCUCCGGGGACGGCUGGGAAGGCCCC	1	no match			
7849	CUGGGAAUACCGGGUGCUGUAGGCUU	1	no match			
7850	UCCUCGAUAAUAUAGUGGUGAGUAUCCC	1	2	ncRNA		tRNA
7851	AAAGUUCAAUAUCCCAAAUUCGGCGUGA	1	1	no annotation		
7852	UCAAAACACACAACACACUCCCAUC	1	0	transposon	LTR	Dm88
7853	AGCGGCGGCAUUGGCGUUGAACCCGGC	1	0	coding gene		CG3788-RA|CG3788
7854	ACUCGCCGAAUCCCGGGGCCGUGGGA	1	no match			
7855	ACGAGUAAUAUGAGCGCCCCAGAGU	1	no match			
7856	AAAACCUGAAGAUCGAUCCUGUAGUCGC	1	no match			
7857	GGCAAAUGCUGACUUGAUCUCGGU	1	0	repeat		LSU-rRNA_Dme
7858	GAUACUUUUAUCAGAGGCCUUCGAG	1	0	transposon	LTR	HMS-Beagle
7859	CAGUACUAUGUGUCCUGCGUGU	1	1	coding gene		CG4033-RA|RpI135
7860	ACUCGCCGAAAUACCCGGGGCCCGAGGGA	1	no match			
7861	AGGACCAUAUCCGAAGACGCGGCGC	1	0	transposon	LINE	Rt1b
7862	AAGCACGGGAGACACACGGCGGGUGCU	1	no match			
7863	CGCCUUGCGUGUUACGGUUGUUUC	1	0	ncRNA		rRNA
7864	UAAAGAAUCCUGACAAGAUCCCGA	1	1	transposon	LTR	invader1
7865	AACGAGGAUGGGAAUGUGAAUGUGG	1	1	no annotation		
7866	GCUGGCGACGCUCGAAGUACUAUCG	1	1	coding gene		CG12142-RA|Tsp42Eg
7867	AUCCUCAACUGUAAGGGUAUCCUCAA	1	no match			
7868	CCAGCUCGGGGAGGUAGUGACGAAA	1	0	ncRNA		rRNA
7869	AUCCGAAGAUAAGUGCAGUUGGAAA	1	no match			
7870	AACGUAACAAAAAAGCUGGGUCGA	1	no match			
7871	UAAUAAACUUGACCAUGUUUCCCCCGGG	1	0	transposon	Low_complexity	F-element
7872	CUUGAUUGCGACGGACUUAGGGAAA	1	0	transposon	LTR	HMS-Beagle
7873	UUUCAUCGUCGUAGAACAAGCCCGA	1	0	no annotation		
7874	AAGGAUUGACGGAAGGGCACCACC	1	2	ncRNA		rRNA
7875	GCCGAACGUAGUCUCGAGAGCGAG	1	no match			
7876	GUUGCAAGAAGUGCGACCGAUCCCAC	1	0	transposon	LTR	GATE
7877	AAACGCACCAGCGCCCAUGGUACUGU	1	no match			
7878	UAUUAUUAAGAUGGGCUUGUAGG	1	0	transposon	LINE	baggins
7879	UGAUCUUCCAGAGAGCGUUUACCGU	1	1	transposon	LTR	Idefix
7880	CCGAAGAUGACCAAGUUGGCUAC	1	0	transposon	LTR	gypsy6
7881	CAAACAGUUAAUCUUGACAUUCGAGA	1	0	transposon	LTR	invader3
7882	AGCUUCAUCCAAGCGAACUGUGCCCGAC	1	1	transposon	LINE	R1-element

7883	UUCAACGGAUUGGACGACGUCGG	1	1	no annotation		
7884	CUCUGAGAAAGCGGACGCGUGAA	1	0	transposon	LTR	invader6
7885	CCGGGAUGGACAUACCUCUGGUGUACCUA	1	no match			
7886	UGUUUCGAAACACGGAGUUAACGAUC	1	0	no annotation		
7887	GAACUUGAGACCCAACAACGCAACCGA	1	no match			
7888	ACCUGGGAAUUGCAUCUGUGACU	1	no match			
7889	AUUAAAACAUCCCUACCCGAAAGCACAC	1	0	transposon	Low_complexity	F-element
7890	CAACCAGAGAGCCUACCAACGUGGUGA	1	0	transposon	LTR	ninja-Dsim-like
7891	AAAAUACUUCAAACCAGCAUCUAGCC	1	1	transposon	LINE	Cr1a
7892	GGCUUUGAGAGGCGCCGUAGUUACCAA	1	1	no annotation		
7893	GGCUGCCAAGAAGGUCCUCGACGGUUU	1	1	coding gene		CG13168-RB|CG13168
7894	GAGAGGCGCCGUAGUUACCACCCCAC	1	0	no annotation		
7895	UUCACAAAGUCGGAGACACCCAUAAGG	1	1	transposon	LINE	accord2
7896	CCGAGAAUAACGGAACUACAGAAGAGCAA	1	0	transposon	LTR	HMS-Beagle
7897	AGCCCGUGGACGGUGUGUGGCCGGU	1	no match			
7898	CGAGAUAGUACGAAUCGAUUACGUA	1	1	transposon	LINE	I
7899	UUCAGAGAUUCUCGAGCUUCAGUCG	1	0	transposon	LTR	ninja-Dsim-like
7900	UAGUACGAGAGGAACCGCAGGUACG	1	0	ncRNA		rRNA
7901	UGCUCUCGAAUGUAAGCCCGAUA	1	1	transposon	LINE	R1-element
7902	AGCCAAGAGUUCCCGUGGUAGGCAGUAG	1	0	ncRNA		7SLRNA-RA
7903	GAGGUCUCCGGACGUGAUCACUGUG	1	0	ncRNA		rRNA
7904	UGCCCGGGAUAGUCCUUUCGUUUAGC	1	0	transposon	LTR	springer
7905	UAGACUUAAGAUGAUGAACCCAUA	1	no match			
7906	GCAAACCGUAAGACCCAGGCUGUGG	1	1	transposon	LTR	ninja-Dsim-like
7907	AGCCAGUUACCGAAGCUGCGGAUUU	1	no match			
7908	AGAGGAGAAGGUGGCUUUGAGA	1	0	no annotation		
7909	UGCGAAAUAAUUGCCAGAGUCACCGGA	1	0	transposon	LTR	Quasimodo
7910	UGAGGAGUACCCGUUUCCCGACCUGCGUA	1	no match			
7911	AGGAUCCUCAACUGUGAGGGGGAUCCUCAA	1	no match			
7912	UCCAAUAUCACCGCGGAGACUGUCGUC	1	1	transposon	DNA	1360
7913	UAACAAGAUGCAUACGAUGUGGCUCCA	1	0	transposon	DNA	1360
7914	ACCGACAAGAACUGCUCAUUCCGAACA	1	0	transposon	LINE	Doc
7915	UUCAAGGUUACCCCAGGAUGCUGUGUC	1	1	repeat		trf
7916	UGGCGAGCGAAACGGGAAGAGCCCAAA	1	no match			
7917	AUCGAUUCUGAUCGGACGGAGAGGG	1	0	no annotation		
7918	AGGCCCGUAUAACGUUAAUGAUUA	1	0	ncRNA		rRNA
7919	AGGCCGUGAGUUCAGCCCUCACAGGA	1	1	ncRNA		tRNA
7920	UGUGAUAGAAGACGAUUUUCGGGCCGA	1	0	transposon	DNA	1360
7921	CGAGGAGAUGUACCAGCUGACCGUG	1	0	coding gene		CG9124-RA|eIF-3p40
7922	AACGAACGAGACCUUAACCUACUAA	1	no match			
7923	CUCAACUGUGAGGGGAUCUUCAA	1	no match			
7924	CUCCGGGGGAAGUAUGGUUGCAAA	1	0	ncRNA		rRNA
7925	GUGGUGGGGUGUGGGGGCGGCUGGAC	1	no match			
7926	ACUCGCCGAAUCCCGGGGCCAGGGAG	1	no match			
7927	CAAAGGAUACUCGAAGACACCACGG	1	0	transposon	LTR	ninja-Dsim-like
7928	CAGCACACGUAUUCCUUCCACUUCUCCUC	1	no match			
7929	AAGAUCAAGAAUCCCGUGCCAAGACUGC	1	1	transposon	LINE	Circe
7930	ACCACCCUGAACGCGCCGAUCUCGUU	1	no match			
7931	UUCAACCAGAUGGGCCUGGAGUA	1	0	Su(Ste)		Su(Ste)
7932	CAGUAUACGAGUCAUGUGCAGCAU	1	1	transposon	LTR	Rt1c
7933	ACGGAUGGAAUCGAGCGCUGAGUAG	1	no match			
7934	UAUUUCAUCGUUAGACGGCUCGGGCG	1	1	AT-chX		AT-chX-B
7935	CACAUUCUGAUUCUGAUUCGAGUUC	1	0	transposon	DNA	transib2
7936	GGGCCUCGUGCUCAUCCUGGCAACAG	1	0	ncRNA		rRNA
7937	GGGUUCGACUCCCGCUCGGUGUACCA	1	no match			
7938	AACCCUCUAAUUCGGCCCGAACCC	1	0	transposon	LINE	baggins
7939	CAUGUACCGGAGGUACGGUUCCAGA	1	0	transposon	LINE	R1
7940	CAAGAUAUGACCAAAGCAAAGAAAA	1	1	transposon	DNA	1360
7941	AUCCGAGGCCCUGUAAUUGGAAUGAG	1	0	ncRNA		rRNA
7942	UCCGUACUGCACGUGGAGUUGUUAAA	1	0	transposon	LTR	GATE
7943	ACGCAUCCUAGAAUCGACCGACC	1	0	transposon	LTR	flea
7944	ACCUUGAUAAACGGUUGACUUUC	1	1	transposon	LINE	Rt1b
7945	AACCGCACUGAGGAGCUAACCGACAA	1	1	coding gene		CG8977-RA|Cctgamma
7946	UCGAGUUGCAGAAUUGAGGACCGC	1	0	transposon	LTR	Quasimodo
7947	GUAAGUGGAGAUGAGCCAGGGCAA	1	0	coding gene		CG7143-RA|DNApol-eta
7948	CAACGCCGCCAGAUCAUCCUCAACAUC	1	no match			
7949	CGGGAUAAGGAUUGGCUCUGAAGG	1	1	ncRNA		rRNA
7950	AUAGAACAUAGAACCAGAUGUCCUAU	1	1	repeat		trf
7951	GAGGACUUGGGCGAUUUAAGUGGC	1	0	Su(Ste)		Su(Ste)
7952	AAUACCGUGCUGAAACCCCUCUG	1	0	transposon	LINE	Doc2
7953	AAGGAAAUGAUAGAACAAGGGAUUGU	1	1	transposon	LTR	Idefix
7954	CAAGAUAUGACCAAAACAAAGCAAC	1	0	Su(Ste)		Su(Ste)
7955	GCCAAUCACAUUAGACGAGACGGCCC	1	1	transposon	LTR	opus
7956	GGCGUUCGAGAGAAUCAUGGUUAAGGAA	1	no match			
7957	GCCGAUCACAGGUCCGCACCUUCGC	1	0	transposon	LINE	R1-element
7958	CAGUACUGUAAACUGACCUAAGG	1	0	transposon	LINE	Doc3
7959	GAUUCCGGAGAGGAAGCCUGAGAAAC	1	1	ncRNA		rRNA
7960	AAUCACAAAACAAGACAUGACACCA	1	0	transposon	LTR	297
7961	CCAGAACGAGGUGGUCCGACAAUU	1	0	transposon	LTR	gypsy3
7962	AUACCACCCUGAACGCGCCCGAC	1	no match			
7963	AGCAAUUGACGGUACCGCACACC	1	2	transposon	LTR	Dm88
7964	GAAUCUAUGGGUGGUGGUGCAUG	1	0	ncRNA		rRNA
7965	GGUUCGGGCCCUCCUUCUAGCGCCA	1	no match			

7966	CUCGUGUUUACCCAGAAUGAUGUUGA	1	0	transposon	Simple_repeat	I
7967	AAACACAGCCCAACUGCGGACCCAUUAA	1	2	coding gene		CG14034-RB|CG14034
7968	GAAGAUCAAAAGUUGGCGCCUGU	1	0	transposon	LTR	flea
7969	UGAGACUCCAAACGGAUUGCGGACC	1	1	no annotation		
7970	ACCCUGAACGCGCCCGAUCUCUUC	1	no match			
7971	AAUAGAUCCUGAAACCGCGUGCU	1	no match			
7972	UCCGAAGAUAACCAGGAGCAGAGCUGAU	1	2	transposon	LTR	297
7973	GGAAAACCCAGAAAAAACCCGAUCAGAA	1	no match			
7974	UAAUACGUUAUCCCGGCACCAUAUU	1	0	transposon	LINE	Cr1a
7975	GUGUGAAGGCCAAAAACGAAACAACU	1	0	coding gene		CG4836-RA|CG4836
7976	AAAGCAAAAAUAACUCCGAAUUAUGGCAG	1	2	transposon	LINE	Cr1a
7977	CUCGGACCUUGAAAAUUUAUGGUGGGG	1	0	ncRNA		rRNA
7978	UGUAUACCUAUACAUUACCGCUAAA	1	0	ncRNA		rRNA
7979	GACGUCGGCUACCCACCCGACCCGUC	1	no match			
7980	UCAACCUGAGAGUUUGGUCCUGGCU	1	no match			
7981	CACCAAGAUCAUCAACCCGUGUUUC	1	2	transposon	LINE	Cr1a
7982	CCAAGGGCUGUGGUCCUGCUCGGUUC	1	0	coding gene		CG2127-RA|CG2127
7983	GCCAAUGAAAACUACACCCAGACG	1	2	transposon	LTR	McClintock
7984	UGAGAGUAGACUUAUUGGUCUGUAAGA	1	0	transposon	Low_complexity	F-element
7985	ACGGGCCUAGAGUUAUUCCCCCAGA	1	0	coding gene		CG40635-RA|CG40635
7986	CACGGUAAAACUUACCAGGUCCGAACA	1	1	ncRNA		rRNA
7987	UCUGCAUGAAGACGAUUCCGAAGA	1	0	transposon	LINE	Circe
7988	GAGACAAACAUGUUCGAGCUACAGCAA	1	0	transposon	Unknown	RR50960
7989	UUCGAGAACAAACGAUACUGUUCGGU	1	0	transposon	LINE	G2
7990	CGCAAACCUAAAUGACAAUGCACGA	1	no match			
7991	UCCAAGGAGUGCGAUUGCUGGGA	1	0	coding gene		CG2127-RA|CG2127
7992	UGCUCUCGAAUGUAUGCCCCGAUAUC	1	no match			
7993	AGUUCGCCGAAUCCCGGGGCCGAGGG	1	no match			
7994	UGAUAUAUAAUGCCAUACUCAGGG	1	0	transposon	LINE	Rt1b
7995	UAGGAACCUGCGGAAGGAUCAUUA	1	no match			
7996	GCCUGAUGACUAUAGCAAGUUGGUA	1	no match			
7997	AAGGAAAUUAGCUGGACGCCCGAA	1	0	transposon	DNA	hopper
7998	UCCAAGCGAACUGUGCCCGAGGCCGA	1	0	transposon	LINE	R1-element
7999	AGAAUCUCGACUAUUGCUACUAUCUGCC	1	0	coding gene		CG17349-RA|CG17349
8000	GGCGCAAAGAGAUGGAUGCCGUGAG	1	0	transposon	LTR	ninja-Dsim-like
8001	UGGUUAAUUCCGAUAACGAACGAGACUC	1	0	ncRNA		rRNA
8002	CAAUCCGAUACAUUUCACGCCCGACA	1	no match			
8003	AGACCAAACACGAUACUACCGGA	1	0	transposon	LTR	RR50207
8004	CACUAUUACACGAUAUACCUUGAGGUA	1	1	transposon	LINE	Cr1a
8005	GCAAAACAUAGACAAUGGGGUUGC	1	1	transposon	LINE	G2
8006	GCUUCCGGGAAGAAUGGCAGACCU	1	0	coding gene		CG5405-RA|KrT95D
8007	UCGAGCGAAGAAGACGAGGACGACCA	1	0	transposon	LINE	G6
8008	AGCUUCAUCCAAUCGAACUGUGCCCG	1	1	transposon	LINE	R1-element
8009	ACACAGCAUUUGUUACCCGAGAAG	1	0	transposon	LTR	flea
8010	CUCAACUGUGAGGGGUAUCCUCAA	1	no match			
8011	CCGCGUCUUGUGGCCAGUGCGUCG	1	0	transposon	LINE	TART-A
8012	GGCCAAGUGAUAUCGUCGGUGGAU	1	0	coding gene		pseudogene
8013	UGCGAAAGUAUAAGAUGUCCAGCGAA	1	0	transposon	LINE	Rt1b
8014	AUGGUUGCAACGGGUAACGGGA	1	1	ncRNA		rRNA
8015	GUAACAACGAAGAUGAUUCUGCCCC	1	1	transposon	LTR	invader4
8016	CCCAGCUCGGGAGGUAGUGACGAA	1	no match			
8017	GUAACUUCGGGAUAAGGAUUGGCUCUA	1	1	ncRNA		rRNA
8018	GGAUCAGCUGUGAGGGGGAUCCUCAA	1	no match			
8019	GAUCCUUCUGACCUUUUGGGUUUU	1	no match			
8020	AUGCUGUGGACUUCGAGAUGACC	1	0	coding gene		CG6527-RA|CG6527
8021	AAAAACCCAUGACCUAAACACGAUUCU	1	0	transposon	LTR	Idefix
8022	CAACAAAGCACCAGGACAUGACG	1	0	transposon	LINE	X
8023	CCCCUGAUGAAGAAGAGAUGCCCAGU	1	0	transposon	LTR	ninja-Dsim-like
8024	AAUACAUGCCGACGGGCGCUGUACCCC	1	no match			
8025	ACGAUCAAUAAAACAUACGCCCGGU	1	0	transposon	LTR	GATE
8026	ACUGUCCCUACCUACUAGUCCAGCCA	1	no match			
8027	GGCUCCGGGGAAGUAUGGUUGC	1	no match			
8028	AAUAGCGAAAAUUGGCCCAAUCUGAUU	1	2	repeat		SAR_DM
8029	UAGUACAAUAGAUCCGACUGGGU	1	0	transposon	LINE	baggins
8030	CAAGCUAGACUAUGUAUCCCCGGGUCC	1	2	transposon	LINE	Doc
8031	AAGGGACGGUCGGGGGCGAUCAGUAUUC	1	no match			
8032	UGGCUCUCUCAGUCGCUUCCCGGGUA	1	1	ncRNA		rRNA
8033	UGUGCCACCAGGUGUUCAAGUUGUCCA	1	0	transposon	LTR	mdg3
8034	GGUUCCUUCCGAAGUUUCCCUCAG	1	0	ncRNA		rRNA
8035	GAUUCCGGAGAGGGAGCCCGAGAAA	1	1	ncRNA		rRNA
8036	ACAUUCGCGAAAAAAGCUCAAGUGCA	1	0	transposon	LINE	Doc2
8037	CAGUCCGAAACACACCCCCCGAUUA	1	0	no annotation		
8038	UGUUUCAUCGUUAGGCGACUCGGG	1	2	AT-chX		AT-chX-B
8039	GGGCGGAGGCGGGUGGCGGGGG	1	2	no annotation		
8040	GAACGUGAGCUGGGUUUAGACCGU	1	0	ncRNA		rRNA
8041	UCAUUGAGCGAACCCGGGGAACUGAA	1	no match			
8042	UGCCGAGCCAUCUGCCAAUCCAGGGACC	1	0	coding gene		CG9029-RA|CG9029
8043	GCACCAGGCAGUGAUUAGAGAUU	1	0	transposon	DNA	HB
8044	GACACGUUCAACCAGAUGGGC	1	0	Su(Ste)		Su(Ste)
8045	AUACCGUAAACGAAAAACUAAGA	1	1	transposon	LTR	Idefix
8046	UCGUGACGAAGCUCCAAGCGGUA	1	1	coding gene		CG2759-RA|w
8047	GGCAGCCGAAUCGAUUUGUGAAACC	1	0	coding gene		CG17945-RA|Mst84Dc
8048	GCUCCCUGGAUUCCACUCGGCGU	1	0	transposon	LTR	ninja-Dsim-like

8049	AGUGGAGCCGGGCGUGGAAUCGCGAGU	1	no match			
8050	GGGCUCCGGGGGAAGUAUGGUUG	1	0	ncRNA		rRNA
8051	UUCCCGGACAUCUUCUUCUCGCUGCUG	1	0	repeat		trf
8052	AACCCGACCCCACGAAGCUGGAUUCGCU	1	no match			
8053	CCACCCUGAGCGCGCCCGAUCUCGUC	1	no match			
8054	GGGGCAUUAGUAUUACGACGCGA	1	0	ncRNA		rRNA
8055	GCCGAGAAAAUGAACACGAAACGA	1	no match			
8056	AAAAACCCUGAAGAUGAUCCUGAGUCGC	1	no match			
8057	AACCUUGACUCCGGCGAUCCCGA	1	no match			
8058	AUAACCAUGAGACCGAUAGUAAACA	1	0	ncRNA		rRNA
8059	UCCAAUAUCACCGCGGAGACUGUC	1	0	transposon	LINE	Doc4
8060	CGCGGUAAUUCCAGCUCCAAUAGCGUA	1	0	ncRNA		rRNA
8061	GACUGAUUGCCCAUGCUGGCUCGCU	1	0	coding gene		CG10206-RA|nop5
8062	ACUCGAAUGAGUGCUUGAUAUGACC	1	0	Su(Ste)		Su(Ste)
8063	CAAGAAUGUCUAGCCGAAACGAG	1	no match			
8064	GCUUUGAUCCAGGAUUUCCGAAU	1	no match			
8065	AAUUUUGUUACGAAUUCAACGAGGU	1	0	repeat		SAR_DM
8066	GCUGAUUGGCUCAUUAAUGUAUCUG	1	0	transposon	LTR	DM88
8067	CCCGCAAUAUGACAAUCGCUCGAUG	1	1	coding gene		CG40609-RA|CG40609
8068	AAAUACCAUAGCAAUCUUCACACGC	1	1	transposon	Low_complexity	F-element
8069	GUAGCUUCGGGAUAAGGAUUGGCUCU	1	1	ncRNA		rRNA
8070	ACACAACAACACAUACCCGCUGGC	1	2	transposon	LTR	GATE
8071	GAGAGUGGACGGGAGCAGCGGGGGCG	1	no match			
8072	AACACUCGUAGAUACCCAACAUGUCCC	1	0	coding gene		CG17567-RA|CG17567
8073	GCGGAGGCGGUGGCGGCGGAGGCG	1	0	repeat		(CGG)n
8074	CACGAUCGAGUCCACGGACAAACCC	1	0	transposon	LTR	blood
8075	GACAGAAACGAGGCCACCGCCGCCC	1	no match			
8076	UGUAUAGAUAUUAGGGAAUAUGUAGA	1	1	transposon	LTR	invader4
8077	AACCGAAAACAUAUCACACGCAACUGAC	1	0	coding gene		CG17618-RB|CG17618
8078	GGAACAAUGUCCGAGUCGGCUCC	1	0	coding gene		CG32575-RA|hang
8079	ACCGAAUUGAUCGCGAGGCUGCAAGA	1	1	transposon	LTR	mdg3
8080	AACUACAAAAUUGCACCCAAAUAUA	1	0	transposon	LINE	TART-A
8081	CUCGGUGUUCAGUACACACAGGG	1	0	repeat		LSU-rRNA_Dme
8082	UUGUUCCCCGGAUAGUUUAGUUACGGU	1	1	ncRNA		rRNA
8083	CCAAAGCUUUGGGCUCCGGGGGAAGU	1	no match			
8084	ACCCAAGAUGAUCAGCCUCUAUCCAC	1	0	transposon	LTR	gypsy7
8085	UGCACCGAAAAUAUAUGCGAGGAGAUU	1	0	repeat		trf
8086	ACGGUACCCGGAGAAGAAGCCCCGG	1	no match			
8087	UUCAACCGGACGGGCCUAGAGUUAUUCC	1	0	coding gene		CG40635-RA|CG40635
8088	CCAACGGGAUACCUCAUACUCGAGUA	1	0	transposon	LTR	micropia
8089	AGGGAGUUUGACUGGGGCGGUACAUCU	1	0	ncRNA		rRNA
8090	AACCAAGCGCGGGUAAACGGCGGGAGUA	1	no match			
8091	CGCGCGCCAAGGAGCACCCCGUCUUUUCUG	1	no match			
8092	ACACCAAUCGCUGAUGAGACACCGGG	1	1	transposon	LTR	accord
8093	CGAGCUACGAAGGGAUCUUGGAGU	1	0	transposon	LTR	ninja-Dsim-like
8094	GAACUGGCGCUGCGGGAUGUAA	1	no match			
8095	CAGACAAAUACCCUAUGCCAGAU	1	0	transposon	LTR	gypsy3
8096	CAGGUAGGGAGUUUGACUGGGGCGGUAC	1	0	ncRNA		rRNA
8097	GUGAUCGAUACGCCCCCUCAGAGUCC	1	0	transposon	LTR	Max
8098	GGAGCAAUCCUGGUGGCGAGUGGCGA	1	no match			
8099	UCUCAUCUUCGUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
8100	CUAACACGUGCGCGAGUCGGGGCU	1	no match			
8101	CGGAGGUGUCCCAAGGCCAGCUCAGU	1	0	ncRNA		rRNA
8102	UGAACGUGUCCUGAAAGUAGUUGG	1	0	Su(Ste)		Su(Ste)
8103	CCCUUACAAUUCCCCUAUCUGGAUA	1	0	transposon	LTR	opus
8104	GGACCCGUCGGACGCCCCCCCAACCCCGA	1	no match			
8105	UAAUGCCGAGAGCUGAUGACGAGC	1	no match			
8106	CUGGCAUUUAUACAGUGCAUUGGACG	1	1	transposon	LINE	baggins
8107	CAGACACGAACGAAAUUUUGAGUAC	1	2	transposon	LTR	Idefix
8108	ACCAACAUGAUGUUUUCGCCGAUGAA	1	1	repeat		trf
8109	GACAAUGAGAGGUACUUUGACCAA	1	0	transposon	DNA	1360
8110	AUAACAACAACUGGAUGGCCCAUGA	1	0	transposon	LINE	Rt1b
8111	UACAUCGAAAAGCUAGUUGACGUAU	1	1	transposon	LTR	invader4
8112	GACGAGAUAAACCAGCACUCAAGC	1	0	transposon	LINE	X
8113	AAAACCUGAAGAUAAUCCUGAGUCGCA	1	1	transposon	LINE	Helena
8114	UCCACAAGGGAGGCGAGCUGUUUGAC	1	0	coding gene		CG3492-RA|CG3492
8115	ACCCGAAGAUAACUCGCAGUCGCUACA	1	1	transposon	LTR	GATE
8116	UGUUUCAUCGUUAGACAGCUCGGGC	1	1	AT-chX		AT-chX-B
8117	AGAUCAGUGAAAUCAUCUUCCGUGA	1	1	transposon	LTR	micropia
8118	CUGCGAGGUAGCCGAAGUCAUUGU	1	0	transposon	Simple_repeat	I
8119	AACCGGACGAAGGUGGGGAUGAC	1	no match			
8120	GAACGAGUGAACUGGCAACAUUUCG	1	1	Su(Ste)		Su(Ste)
8121	UCAACUGUGAGGGAUCCUCAACUGUGAG	1	no match			
8122	ACCGAAACUAGACUAGACAACAAU	1	0	transposon	LINE	G2
8123	ACGGAAUGGAAUCGAGCGCUGAGUUU	1	2	transposon	LINE	X
8124	UGUUUCAUCGUUAGACGGCUCGGGUG	1	1	AT-chX		AT-chX-B
8125	UCUGCCCGAUGAGAAGAAUGGCCCG	1	0	transposon	LINE	RR49783
8126	AGGAAUAAUGGAAUAGGACCGCGGU	1	no match			
8127	CCGCGUGAGUGAUGAAGGCCUUAGG	1	no match			
8128	AUGCAGUGACCUUGGAUGAUGCUCUGG	1	0	coding gene		CG4750-RA|loopin-1
8129	AAAAACCUGAAGAUGAUCCUGAG	1	1	transposon	LINE	Helena
8130	GCCGUUACGAAAACACCGUUUAAG	1	0	no annotation		
8131	AACCCAAGCUUUUGGGCUCCGGGGGAAG	1	1	ncRNA		rRNA

8132	CGCUCAACUACGAAACUCAAACUCCCGG	1	0	no annotation		
8133	UGUAAACGGUAGCUAAUUCGAGCGUC	1	1	transposon	LINE	Cr1a
8134	UCAUCACCAGGAGUUCGUCUUUA	1	1	no annotation		
8135	UGCUCUCGAAUGUAUGCCGAUAUU	1	no match			
8136	CCUGACUUUAUGCCCCAGUUUUUCCAG	1	0	transposon	LINE	G2
8137	GUGGCGGCGACGACCCAUUCGAAC	1	no match			
8138	AGUCCUGAGAAAUCUUUAGAAAAUGAU	1	1	transposon	LTR	mdg3
8139	GGAUCCUCAACUGUGAGGGGAUCUCA	1	no match			
8140	AGUUGAGAUUAUCGCCAAUGACCAGG	1	1	coding gene		CG31359-RA|Hsp70Bb
8141	UGCACUAAUAUAAGCCCUGACGUCCGUA	1	1	transposon	LTR	G4
8142	UCAUGACCGAAAUUCGGAAAAACGAAUU	1	0	repeat		SAR_DM
8143	CCGGUCCCGGGACAUGGCGUGGUGGGUAG	1	no match			
8144	ACCCUUGAUAUACGGUUGACUUUC	1	1	transposon	LINE	Rt1b
8145	CAUGUCAUAACAACUAAACCCGAAGA	1	0	transposon	LTR	Quasimodo
8146	GACGUUAAUGAAGAUGCCGUGGUGC	1	0	coding gene		pseudogene
8147	CUUAGAAGCAUUGGGCCAAACCGA	1	0	transposon	LTR	opus
8148	UCCGGGAGACACCAAUGCUGGCCAAGU	1	0	coding gene		pseudogene
8149	AGGUAGGGAGUUUGACUGGGCGUAC	1	no match			
8150	AGGACAGAGACGCCUAAGCUAUUCGGC	1	0	transposon	LINE	X
8151	UAGAUCAGUCACAAUAAAUACUUGA	1	0	transposon	LTR	DM88
8152	UACUGAGACAUAGUGACAUCGGCC	1	1	no annotation		
8153	UCCGAAGUGACCGACCGAAACAGUC	1	0	transposon	LTR	diver2
8154	CGCUACAUAAGAAAGCCGACAUCCCGA	1	0	transposon	LTR	GATE
8155	UGUUUCAUCGUUAGACGGCUCUGGC	1	1	AT-chX		AT-chX-B
8156	UCCCCAUACGGUACAUGGCCGCGAGGA	1	0	transposon	LINE	F
8157	CAGUUCAUCAAAGCGCCUUACUGGAGGA	1	0	transposon	LTR	flea
8158	GCACACAGAGGAGAGCAGAUGAGAACA	1	no match			
8159	UGACCACCCAUACAAAUCCCCCUCGC	1	no match			
8160	UCCGAGGAACCAAUUGAUCCAGCUGCUU	1	0	no annotation		
8161	UCAUCAUCAAUGACCGGCCAGCCAG	1	0	transposon	LTR	springer
8162	GAACCAUCAGACGCCUCCAGAGGAU	1	2	transposon	LINE	G
8163	GGCUAAAACCAAGCGAUCGCAAGA	1	0	ncRNA		rRNA
8164	GUAGUACGAGAGGAACCGCAGGUA	1	0	ncRNA		rRNA
8165	CCCAGAAACACAGUGUUUGACGCCUG	1	2	coding gene		CG18743-RA|Hsp70Ab
8166	UGCUCGAUCAAAAGGCCCUGGCAU	1	0	no annotation		
8167	CGCUCACCCAGACCAUGAAAACUUCG	1	2	transposon	LINE	G
8168	GGAGACAAGAUGCCAGAUGUACCAAU	1	0	no annotation		
8169	UGGGCACGAGGGCUAUCCGGGCGAGCU	1	0	coding gene		CG32445-RA|CG32445
8170	UGCUCUCGAAUGUAGUGCCCGAUA	1	no match			
8171	UUCAAAGUUACCCCAGGAUGCUGU	1	0	no annotation		
8172	CCGCCGAAUCCCGGGGCCGAGGGAGC	1	no match			
8173	UUCGCCGGAGACGGGAACGGCUGCUGA	1	0	transposon	Low_complexity	F-element
8174	UUUGUACGAGAGGCUUUAAGAGA	1	1	transposon	LTR	invader1
8175	GAGGAACUGAGGAACGCAUUUCAAA	1	0	transposon	LTR	diver2
8176	UUCCCCACCUGCGGUCCGGCGGCAUUCC	1	0	coding gene		CG17377-RA|CG17377
8177	CGUUGACCAAUUUGCGAACAUACGCC	1	1	transposon	LTR	DM88
8178	AACCUAUCGGACGACGCAGCGCAU	1	2	transposon	LINE	Quasimodo
8179	GCCGAAUUUGAGACCCCAGCCGCUCC	1	0	coding gene		CG40635-RA|CG40635
8180	GCCGAACGUAGUCUCUCGAGAGCGAUA	1	1	transposon	LTR	invader1
8181	AGGAACCAUCACUGGGGCACCGUGG	1	1	transposon	LINE	F
8182	ACCACCCUGGACGCGCCCGAUCUCGUC	1	no match			
8183	CAACUGUGAGGGGAUCCUCUAA	1	no match			
8184	AAGAGAUGGGGAAGCUCCGUUUC	1	no match			
8185	GCAGCAUAUCUCGGCAGCCCGCGGU	1	1	transposon	Simple_repeat	I
8186	CAAGCUAGACCAUGUGUCCCCGGGUAA	1	2	transposon	LINE	Doc
8187	UCGCGGGACAACUGGGCCGAGGC	1	0	coding gene		CG5265-RA|CG5265
8188	CUAGCCGGGCUGAAUGACCCAUAUUA	1	0	ncRNA		tRNA
8189	CUGGGGUUGCCACUUAGGACGAGGC	1	0	transposon	LTR	ninja-Dsim-like
8190	CGAAAAUGCAGACCUAUUGUCAUCCGU	1	1	transposon	LTR	flea
8191	AAGGAUCUAACACGUGCGCGAG	1	no match			
8192	GCGAAGGUGGCGGAAUUGGGAUUCG	1	no match			
8193	CCCGUUAUAUCUGGCUCGUCAAAU	1	1	transposon	LTR	Quasimodo
8194	UCCUCAAACUGUGAGGGGAUCCUCAA	1	no match			
8195	UCUCAUCGUCUUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
8196	AUCAGCGAUGUGUGGGGCAAGACA	1	2	Su(Ste)		Su(Ste)
8197	UCCCAGCUCGGGGAGGUGGUGACGUAAAA	1	2	ncRNA		rRNA
8198	GGAGCCUGACGACCACGACCUCUUAG	1	1	transposon	LTR	diver2
8199	AGUAUUACGACGCGAGAGGUGA	1	0	ncRNA		rRNA
8200	UUCCGAGAAAGCCCGCAGAAUUUGCAUC	1	0	transposon	LTR	flea
8201	ACCAAGAUCAUCAACCCGUGUUAUUCU	1	0	transposon	LINE	Cr1a
8202	AAGAACUGGCGCUGCGGGAUGAAC	1	2	ncRNA		rRNA
8203	AAGACAGACAUGCCUCGCUCCCGAA	1	0	transposon	LTR	invader1
8204	CGAGCGCACGUGUUAGGACCCGAAA	1	no match			
8205	UGGCCGUGAGGGGGAUGCCCGG	1	no match			
8206	AAAACAACGACCCGCUACCGGAG	1	0	transposon	LTR	opus
8207	CACCAAGCGUUGGAUUGUUCACCCA	1	0	ncRNA		rRNA
8208	UACUCGAAUGAGUGCUUGAUAUGACCA	1	0	Su(Ste)		Su(Ste)
8209	CCACCCUGAACUCGCCCGAUCUCGUC	1	no match			
8210	UUGUAACAACUCACCUGCCGAAGCA	1	1	ncRNA		rRNA
8211	CAGAUCCGAAAUCGCACGAGCUGCGUU	1	0	coding gene		CG7578-RA|sec71
8212	GCAUUCGCGUCGAGGCCCAGGAUUU	1	1	transposon	LINE	R1-element
8213	GCAUUCAAUACAGCGAGAUGGCCCC	1	1	transposon	LTR	Quasimodo
8214	GCCUGGGCGACGGGCGGUCCCUCGACCAC	1	0	coding gene		CG10377-RA|Hrb27C

8215	ACAAGAUGCAUGCCCGCGCCCGA	1	no match			
8216	CUGCACCCGCUCACGCUCUGACUGA	1	0	transposon	LINE	F
8217	AACACUGAGAGAUGAUGACGAGGCUCU	1	no match			
8218	UGCCUUGACAUCGAAUGGAGUGCCAGA	1	0	transposon	LINE	R1-element
8219	ACUGAAUAAACGAAAUGGAUGACUA	1	2	transposon	LTR	invader4
8220	CCCUGGGUUCAUUUCCCAGCUCGCCCC	1	no match			
8221	UCGAACGAUAAACGGUGCUAACCAGCC	1	0	transposon	LTR	GATE
8222	AGAGGAGGCGAGAAUUUUACAUUCA	1	1	AT-chX		AT-chX-B
8223	AUUGCUAACAUCGUCCCGGAUAAC	1	0	coding gene		CG31988-RA|CG31988
8224	UGCUCCCGCAAUCUCCCUGUGGAUCCG	1	0	coding gene		CG41423-RA|CG41423
8225	CUGUAACUCGGGUCAUAAAAAAAAA	1	no match			
8226	ACUUCAUAACCGAGCGACUGUUCAUA	1	no match			
8227	GUGGAAAGAGAGCAAUUUUGGCCGU	1	0	transposon	DNA	1360
8228	AGAUCAAGAAUUCCGUGCCAAGA	1	0	transposon	LINE	Circe
8229	AAGAGCAUAAAGACGACGAAAGUGAC	1	1	transposon	LINE	Doc3
8230	UUAAGUCCCGUUACUUGUGCCCAG	1	0	transposon	LINE	G6
8231	CGUUUCAUGAGGAAUCGGCCCAGUG	1	1	transposon	LINE	G2
8232	GUAAGUGGAGCCGUACCUGUUGGUUU	1	0	ncRNA		rRNA
8233	GGCUCGAGAUCGAAGAGUAUGAGU	1	0	transposon	Simple_repeat	I
8234	ACCCCGUGCACCUUUA	1	1	no annotation		
8235	AAUUGACAAGGGCAUGAGCGACAC	1	0	transposon	LTR	springer
8236	AACCUGAGAGUUUGAUCCUGGCU	1	no match			
8237	AACUGAACAUGGUCAUCCGUAAGCU	1	no match			
8238	CCAGAUGAUAUUCCCGAAAACAGCCGU	1	no match			
8239	CGCCGUAUACGCCCUGGACUGCGA	1	0	coding gene		CG14801-RA|CG14801
8240	CACCCUGAACGCUCCCGAUCUCGU	1	no match			
8241	UGAACCACUUUCAUGCUUGGGGU	1	1	ncRNA		rRNA
8242	UUCUACCACAACACGCACACAUACAC	1	0	transposon	DNA	1360
8243	GCGGGCGGCCCCGGGGUUCGUUUCCCC	1	no match			
8244	UGCUCUCGAACGUACGCCCGAUAU	1	1	transposon	LINE	R1-element
8245	UUCCGAGGAACCAAUAGAUCCAGCUG	1	1	Su(Ste)		Su(Ste)
8246	CCGGCAUUAAUUUCUUUAAUUAUGACA	1	0	no annotation		
8247	UGGAGUUCGAGGAGUGGCCCGUUA	1	no match			
8248	AUACCACCUGAACGCGCCGAUCUCG	1	no match			
8249	GAAGAGAGAAGAGAGAAGAGAGA	1	0	repeat		(GA)n
8250	CCCUGAACGCGCCCGAUCCCGU	1	no match			
8251	AAAUUAUGCACCGAAUAUAGCGAUA	1	0	transposon	LINE	RR49783
8252	CGAAUCCCGGGGCCGAGGGAGCGAGAC	1	no match			
8253	UUCGAGAGAGCCUGGAGCCACC	1	0	transposon	LINE	Cr1a
8254	UGCUUUCGAAUGUAUGCCCGAUAU	1	1	transposon	LINE	R1-element
8255	UGUCAUCGUCGUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
8256	GUAGUACGAGAGGAACCGCAGGUACGGA	1	0	ncRNA		rRNA
8257	CUGACGAUAUUGCCCUGUUGUACAGC	1	0	transposon	LINE	BS
8258	UCAGUCUUGAAAAUGUUAAUACCGAAU	1	1	transposon	LTR	invader6
8259	CUCGGGAUCAAGGGCAACGAGUCC	1	1	coding gene		CG40635-RA|CG40635
8260	CAGACAGAGACGGAGCUUUCUCCAGU	1	0	transposon	LTR	297
8261	CACCGACCGAUCUUAUUGGCGCACCCGC	1	1	no annotation		
8262	ACAACGACUCCUGGAAUCUCUGAA	1	1	coding gene		CG4230-RA|CG4230
8263	CAACUGUGAGGUGUAUCCCUCC	1	no match			
8264	CACGACCCCCAGCUACGUGGCCUUCACC	1	no match			
8265	UCGAGUAAUAAAAUGCCAUUUCAUG	1	no match			
8266	GUCCCGCCGGUUGGUACCUACAGUGAAU	1	no match			
8267	AUCCUCAACUGUGAGGAA	1	2	coding gene		CG13004-RB|CG13004
8268	AACCUGAACGCGCCCGAUCUCGUC	1	no match			
8269	AGCAGAACUCCGAUGGCGAUAGCAA	1	0	no annotation		
8270	UACCCAGAAACAAUGCUGAAACACUA	1	0	no annotation		
8271	ACGUUGUGGCCGAUGAAACCCUGGUUC	1	no match			
8272	UUGGUCAACGAAAUGGGUUUUGGUGAU	1	1	transposon	LTR	DM88
8273	UCAAAUAGAAAUGGACUCAGAGUCU	1	1	transposon	LTR	rover
8274	CAGAACGUCAUCGAACCCUAGAGG	1	0	transposon	LTR	GATE
8275	CACGAUGAUCGAUGACGAACUACGA	1	0	transposon	LTR	ninja-Dsim-like
8276	AACACAUCGAGCGACCGCGUAGUC	1	0	transposon	LINE	Doc3
8277	ACGAGUAAUCGAUCGUUCAAGUAUUGC	1	1	transposon	LTR	micropia
8278	CGAGAACAAACGAUACUGUUCGGC	1	0	transposon	LINE	G2
8279	UGCCGUUCGUCCUGGAGUCCCGA	1	0	coding gene		CG10252-RA|CG10252
8280	AGACUUCAAAGUCUACGUGCUAAAAG	1	no match			
8281	AAUACAUGCCGACAGGCGCUGACCCC	1	no match			
8282	GAGUUCGAAUCUCUUAGCAACCACCA	1	no match			
8283	UCAAGAGUUAUUUAAUGCAAUACAAG	1	0	transposon	DNA	HB
8284	GCAACUGAACCCAAGAAAAUCGCCU	1	0	transposon	LINE	Doc
8285	GGAUCCUCAACUGUGAGGGGUAUCCUCAA	1	no match			
8286	GCGAGGACGAGGGACGCGGACGCA	1	0	coding gene		CG7564-RA|CG7564
8287	AGGGCAAAUGCUGACUUGAUCUCG	1	0	repeat		LSU-rRNA_Dme
8288	CUGGAUCGAUUGGUUCCUCGGGAUCAA	1	0	repeat		trf
8289	CAGACAUUAAAGUUACGAUAUUU	1	2	transposon	LTR	McClintock
8290	AUGGCUCUCAGUCGCUUCCCGGGUA	1	no match			
8291	CUGAACCCGCAGAACACCGUGUUCGACG	1	no match			
8292	CUCAAUAAGCUGCCAGACCUGAUCGAC	1	0	transposon	Low_complexity	F-element
8293	CACGGGAAAACUUACAAGGUCCGAAC	1	1	ncRNA		rRNA
8294	AGGGUGCGAGCGUUGUCCGGAAUU	1	no match			
8295	ACCACAGAGUACGGGGCUUACUAAAGU	1	0	transposon	LINE	baggins
8296	AAACGCUUUCAACACCGCCAGAUGGCCCA	1	2	transposon	LINE	F
8297	CGGUCCGACACCAUGUCCCGUUGC	1	0	transposon	Simple_repeat	I

8298	GUGGCUUGAGAUGUGCCGUAGUUA	1	no match			
8299	AAGGCCGGCGAGUUGAUCGAAA	1	no match			
8300	GAAACAGAGAUCUUCAGCAGUGGGA	1	0	AT-chX		AT-chX-A
8301	GUGGAAUUGGUCUUGCUGGCCACAG	1	0	transposon	LTR	blood
8302	ACCUACGAAAUAAGCUGUCUGAGUU	1	0	coding gene		CG40084-RA|CG40084
8303	AAAAAGUCCAAGCUGCGAGAGAAAC	1	0	transposon	LINE	G2
8304	CAGAAAAUGACGACCCAGAUGAGAA	1	0	transposon	LTR	roo
8305	CCAGAUGAUAUUCCGAAAACAGCCGUCAC	1	0	transposon	LINE	Doc4
8306	UAGAUCUUCACCGUUGACUUGCCCCG	1	0	Su(Ste)		Su(Ste)
8307	UAAACGAGUGUUAUUGUGGGCCGGU	1	0	ncRNA		rRNA
8308	UCGAGGUGGGCCUUCUGACGAGAA	1	1	no annotation		
8309	UUGAUACGAAUGAGACCCGCAGUAAAG	1	0	repeat		trf
8310	UGGGCUCCGGGGAAGUAUGGUCGCA	1	no match			
8311	CAGCCUAGGAGACGUGGACACGGA	1	1	transposon	LINE	R1-element
8312	UCUCAUCGUCGUAGAACAAGCCCCG	1	2	Su(Ste)		Su(Ste)
8313	AUACCACCCUGAACGCGCCCGAUC	1	no match			
8314	AACCGAUAACGACCUUACCACCGCUCC	1	0	coding gene		CG3051-RB|SNF1A
8315	AGUGGAGCCGGGCGUGGAAUGCGAGUG	1	no match			
8316	GCCACCGAGAUAUGAGGACGAAGUG	1	0	transposon	LTR	ninja-Dsim-like
8317	AAGGGGGACCUGAGCUGUCGUGCACCCG	1	no match			
8318	ACCGGACGGGCCUAGAGUUAUUCCCC	1	0	coding gene		CG40635-RA|CG40635
8319	CGGCAAGACCAGUUCCACUCAGUUG	1	0	coding gene		CG8838-RA|CG8838
8320	CACGCAAUAAAAACUUUACGAGUGC	1	no match			
8321	CACCAGAGAUCUAGAGGGCCCAAGG	1	2	no annotation		
8322	CAAGAUAUGACCAAAGCAAAGACUAG	1	1	transposon	DNA	1360
8323	UGUUUCAUCGUUAGCCGGCUCGGGCG	1	1	AT-chX		AT-chX-B
8324	CAACACAUGAACAGAGUCAUUUUCAA	1	0	transposon	LTR	McClintock
8325	AGCGGAUGAGCUGUGGAUAGGGGU	1	no match			
8326	UACUAUCUCGGGGAGCUUAUGCGAU	1	0	transposon	LINE	I
8327	GUGUCAAGUUCUUUAGCCACCAGUACA	1	0	transposon	LINE	Doc3
8328	GGGGAGUUUGACUGGGGCGGUAC	1	1	ncRNA		rRNA
8329	UGUUUCAUCGUAGACGGCUCGGGCG	1	no match			
8330	AACUGACGGCGCCACUGUAAAUGUGA	1	0	transposon	LINE	R1-element
8331	UUAGACCUCGGAGAAUUUGGUAG	1	no match			
8332	ACUCGCCGAAUCCCGGGGCCGAGGGGAGC	1	no match			
8333	GGACGAGGCGUUCCGAGCUGCCCAG	1	0	coding gene		CG4836-RA|CG4836
8334	AGCUUGAUCUAGGAACAGUCAGCUC	1	no match			
8335	AUUGGCGAGAGACCGAUAGCGAACAAGU	1	no match			
8336	GAUCCUCAACUGUGAGGGGAUCCUAA	1	no match			
8337	AAUAGGGACGGUCGGGGGCAUCAGUA	1	no match			
8338	GCGAGAGACCGAUAGCGAACAAGUACA	1	no match			
8339	GAAAACCGACGACACUAUAGCCCAUU	1	0	transposon	LTR	INE-1
8340	UUGACAUAUCAAGAUAUUCCGAGCUC	1	no match			
8341	CCCCUGAUGAAGAAGAGAUGCCCAG	1	0	transposon	LTR	ninja-Dsim-like
8342	UAGUGUUCAAUGCCUCCAGCCCU	1	0	transposon	LTR	GATE
8343	CACGUCCAGUGCUGGCUGAAGUACU	1	0	Su(Ste)		Su(Ste)
8344	GUCCCUGCCCUUUGUACACACCGCC	1	0	ncRNA		rRNA
8345	GCAUCGGUGGUUCAGUGGUAGAAUGCUU	1	1	ncRNA		tRNA
8346	CGCACAAGCGGUGGAUGAUGUGGUU	1	no match			
8347	CCGGAGCGAGGAGGAGGACUAGCUGGC	1	0	Su(Ste)		Su(Ste)
8348	CAGACCUAUACAGAUUGGAACAUUU	1	0	transposon	LTR	McClintock
8349	UGCUGACUAAGAGGGCUGCACAG	1	0	transposon	LTR	HMS-Beagle
8350	AAGUGAAAUAACCCAAUACGUAGGC	1	0	no annotation		
8351	GUCCCUACCUACUAUCCAGCGAAACC	1	2	ncRNA		rRNA
8352	AAACCCGAAGAAGACUCUACUCGAG	1	1	transposon	LINE	Circe
8353	AAUUUUGAACGAAAGGGAAUACGGUUC	1	0	ncRNA		rRNA
8354	GAGGAACUAGCAGGAUUCUUUGCC	1	0	transposon	LTR	3S18
8355	AAGGAGUGUGUAACAACUCACCUGC	1	0	ncRNA		rRNA
8356	GACGAGAUAAACCAGCACUCGAGUA	1	0	transposon	LINE	X
8357	CGGCUCCGAAAUCGAAAACACAACACA	1	0	coding gene		CG8287-RA|Rab8
8358	GACAAAACAACAUCCCGGGAAAUGAG	1	0	transposon	LINE	baggins
8359	ACCAUCUGUGGGAUUAUGACUGAA	1	no match			
8360	AAUGUGAGGGGAUCCUCAA	1	no match			
8361	CCGCAAGGCGUAGGCGAUGGAUA	1	no match			
8362	CGGUCCUGAGAGAUGGGCGAGCGCCGUU	1	no match			
8363	UCCUGGGAAACGGGGCGCGGCCGGAAA	1	no match			
8364	GGGGAACUGAGGAACGCAUUUCAA	1	0	transposon	LTR	invader3
8365	UGCCCCACAUCUAGUGUUUCAGCAUUG	1	0	no annotation		
8366	AUUGAGAGGCGCCGUAGUUACCAC	1	1	no annotation		
8367	GAGCGCACGUGUUAGGACCCGAUAU	1	no match			
8368	AAAACCUGAAGAUGAUCCUGAGUUCGC	1	no match			
8369	AUGAGGCCAUUUACCCAUAGAGGG	1	0	ncRNA		rRNA
8370	CCCUGAUGACACGCAGUACCAGAGG	1	0	transposon	LINE	INE-1
8371	GUGCUACUCGAACUGUAUGCCCGAUAU	1	no match			
8372	GGAUCCUAAACUGUGAGGGGAUCCUCAA	1	no match			
8373	ACAACAAUUACAACCGAGGCAGCAU	1	1	transposon	LTR	gypsy8
8374	CUCAGCGAUGUGUGGGGCAAGACAA	1	1	Su(Ste)		Su(Ste)
8375	UUACAUAAGAUAUGAACGGAGCCC	1	0	transposon	LINE	Cr1a
8376	UGUCGCUGAGCGAGGGAAUCCGGG	1	0	coding gene		CG3704-RA|CG3704
8377	GCUAAACCAUUCGUAGACGACCUUCU	1	no match			
8378	UGCCCCCACUGGACGACGUUUGCACCUU	1	0	coding gene		CG3722-RA|shg
8379	ACGAACGAGACUCUGGCAUGU	1	no match			
8380	UUCUGACGAGAACUCGAGCCGCGUA	1	0	no annotation		

8381	UGUUUCAUCGUUAUACGGCUCGGGC	1	1	AT-chX		AT-chX-B
8382	AGCGACCCACUCUUAUGCUCGGCAA	1	1	coding gene		CG4439-RA|CG4439
8383	GCAAAACUUAGGCUGGGACGAUAA	1	0	transposon	LTR	Rt1c
8384	CCGGCAUUAAUUUCUUUAAAUAUGA	1	0	AT-chX		AT-chX-B
8385	CCCUACUGAUGAUGUGUUGUUGCCA	1	no match			
8386	CGGGCCGAAAUCAAUGUUGAUCGAAU	1	1	transposon	LINE	G
8387	GCCCAAGCGAUCGAUGCGAUAUUAUC	1	1	coding gene		CG33933-RA|Indy-2
8388	AUAUAGCAGACAUCCUGAAACGUU	1	0	no annotation		
8389	AACCAUAUCGGACGACGCAGCGC	1	0	transposon	LINE	Rt1a
8390	UGCACCGAAAAUAUAUGCGAGGAGAUA	1	1	repeat		trf
8391	GAGGCAAAAAGGACAUUCACCCAGG	1	0	transposon	LTR	HMS-Beagle
8392	ACCUUAACGAAACCAGAGACAUAAUCC	1	1	transposon	LTR	gypsy4
8393	CCUGACCAGAAGAACCGUAAGGCAU	1	0	transposon	LTR	micropia
8394	CAAGAUAUGACCAAAGCAAAGACUUUU	1	2	transposon	DNA	1360
8395	UGAUCUCGGAAGCUAAGCAGGGU	1	no match			
8396	ACGUCCGCACUAAGCUUGCCAUCA	1	0	ncRNA		7SLRNA-RA
8397	GGCAUUGAGAGGCGCCGUAGUUACCAC	1	1	no annotation		
8398	UUCCCAACAAGACAUCCGAUUAGUUC	1	0	transposon	LINE	Circe
8399	AGCCGGUGAUCGAUAGUUCCUCU	1	0	repeat		trf
8400	AUGCGAGUGCGAUCUCACCGAUGUC	1	0	coding gene		CG5458-RA|CG5458
8401	CCUACAUGGAGACAUCACCGUUGGU	1	1	transposon	LINE	G2
8402	AGGAGAUAUACUUUCAGCCGCGUA	1	1	coding gene		CG16781-RA|CG16781
8403	UCCGAUUUCAAGUACCAGGCGAACA	1	0	repeat		RSP
8404	UAGGAAACUGGAUGAAGACAUGCUA	1	0	transposon	LINE	Rt1a
8405	UGCAAGAUAAAGACCCUUCUACCUGCGUC	1	1	transposon	LINE	Helena
8406	CUGGUUGAUCCUGCCAGUAGUUAUAUGC	1	0	ncRNA		rRNA
8407	AUGGAGCGCAUGUUCACCCAGAUGACAC	1	0	transposon	LINE	G6
8408	AGGUGAACCGUGAGAUUAUCGAG	1	0	coding gene		CG6671-RA|AGO1
8409	UUGCAACGGGUAACGGGGAAUCAGG	1	0	ncRNA		rRNA
8410	CAGGCAAGGUGUUGUAAGUGGUUGAG	1	0	ncRNA		rRNA
8411	GCCAGCAUGACGUUCUUCUAUCUGAUU	1	0	coding gene		CG4692-RA|CG4692
8412	UAGUACAAUAGAUCCGACUGGGUGA	1	1	transposon	LINE	baggins
8413	AAAUACAUGCCGACGGGCGCUGACCCCCC	1	no match			
8414	UAACGAUGACAGUGCCUGGGACAA	1	0	coding gene		CG5621-RA|CG5621
8415	UCCUGACGAAGAGCGCAAGAAGA	1	0	transposon	LTR	opus
8416	UGUUUCAUCGUUAGACGGCUCAG	1	1	AT-chX		AT-chX-B
8417	UCCAGAGUUAUGUUAUUUCCAGUAGGUC	1	0	transposon	LINE	accord2
8418	AAGACAUUAAAGAAAUAGUAGAUCACGA	1	1	no annotation		
8419	UGCAACUCGAGCGACGCCGUCUCUGCCUA	1	2	transposon	LTR	Quasimodo
8420	GCAUUCAAUACAGCGAGAUUGCC	1	1	transposon	LTR	Quasimodo
8421	CAGUCUAUGAAAGGUGUGUUGCCC	1	0	transposon	LTR	GATE
8422	GGCAAUACCAUCACCGUGGGCGUGAAGA	1	0	coding gene		CG4362-RA|CG4362
8423	UGCUUGGACUACAUAUGGUUGAGG	1	0	ncRNA		rRNA
8424	ACAACGAGAAGACAGUAAUCGACAGCAC	1	0	no annotation		
8425	AGUAUCUAGGACUACCAUGGUCGCA	1	1	ncRNA		rRNA
8426	UGACGUGACCCCAGUGUUGAAUAGC	1	1	coding gene		CG32026-RA|CG32026
8427	CGGUACUAUUACUUUGAACAAAUUAGA	1	0	ncRNA		rRNA
8428	AUCGAAGGAACGAUCGCAAGGCAGC	1	0	transposon	LTR	flea
8429	AAACCUGAGGGGAUCCUCAA	1	no match			
8430	ACCAGACGGAGGUCUGCCUUUCGCAGUC	1	no match			
8431	GUUGCUUUGAGAGGCGCCGUAGUUA	1	1	no annotation		
8432	GCAACUGUGAGGGGAUCCUCAA	1	no match			
8433	UCCAAGUCCAUCUAGGUCCCGCAGA	1	0	transposon	LINE	R1-element
8434	GGGACGUUAAUGAAGAUGCCGUGG	1	0	coding gene		pseudogene
8435	AUGGAUUGAAACGUUUCAGUCAGA	1	0	no annotation		
8436	GUGGGCCGGUACUAUUACUUUGAA	1	0	ncRNA		rRNA
8437	CACCCACUGGAGGUCCGAACCGAUUAA	1	no match			
8438	GCCCAAGACCCCCGAGAAACCAAAGCC	1	0	transposon	LINE	G6
8439	AACGAAAGUUAGAGGUUCGAAGG	1	0	ncRNA		rRNA
8440	CACCUCCGCCCAGUUGGAAUAUCCAAA	1	2	coding gene		CG3748-RA|CG3748
8441	UUCAAGGUUACCCCAGGAUGCUGUGA	1	1	repeat		trf
8442	AUAAACGCGAACGACUUGAAAAA	1	no match			
8443	AGUACCUCGACCAGACGGUGGCCCCCC	1	1	coding gene		CG11591-RA|CG11591
8444	AACGAGUUCCUCUGCCACAUGCCCAU	1	2	Su(Ste)		Su(Ste)
8445	CCGGGAAGACGUUGAAUUGAUCGG	1	no match			
8446	UUCAGAGAAAACCAAGGAGAAGU	1	0	no annotation		
8447	GCAUUCGAGACCCAACGGCACGUGU	1	1	transposon	LINE	INE-1
8448	AGGAUCAAUAGCCGAGUCGCUCUAGC	1	0	transposon	LINE	INE-1
8449	CGGUUCAUUCCCGGGUUUCGGCACCA	1	no match			
8450	AAAACCUGAAGAUGAUCCUGAUUC	1	1	transposon	LINE	Helena
8451	CGCAUGUGAAUGUUGGCCCCAGAA	1	0	no annotation		
8452	UCCUGGGAAACGGGGCGCGGCCGGAA	1	no match			
8453	GCCUCGGGACUUCCUAUAUAAGCGGAA	1	0	ncRNA		snoRNA
8454	UACUCAUAGACACCCUAGGCUGAGACC	1	0	transposon	LTR	G4
8455	ACUGCAGAUCGUAUGGUCUUGUACC	1	2	ncRNA		rRNA
8456	UACUCAUCGUCGUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
8457	UGCCGAGAGCUGAUGACGAGCUU	1	no match			
8458	ACAUAAGAAAGCCGACAUCCCGACA	1	0	transposon	LTR	GATE
8459	UUGGGAACACCGCGUGUUGUUGGCU	1	1	ncRNA		5SrRNA
8460	AAGAGGGCGUGAAACCGUCAAGAGGU	1	no match			
8461	AAACACUCAGAACGUCUUGAUUCG	1	0	transposon	LINE	RR49783
8462	CAAACCAAUCCCAGCCAAACCCGUCAA	1	0	transposon	LINE	G2
8463	ACUUCCAUCCUAAAACUGAUAGACAGC	1	1	Su(Ste)		Su(Ste)

8464	UAGAGAGCGAGAGAGCGGGGAGC	1	0	transposon	DNA	1360
8465	GAUCUCCCGUGUGUGUUGUUCGC	1	0	coding gene		CG6963-RB|gish
8466	GAAGAAGAGGUGCCCAGUGAGUU	1	0	transposon	LTR	ninja-Dsim-like
8467	AAGUAAAAGACCAAGCCAAGAGAC	1	no match			
8468	GCGGAGGUUGCGGUGCUGUACACUG	1	0	ncRNA		snoRNA
8469	UCUCAUCAUCGUAGAACAAGCCCGUA	1	2	Su(Ste)		Su(Ste)
8470	ACAUUCAUAACCGAGCGACUGUUCACAC	1	2	transposon	LTR	GATE
8471	UCCGAGCUAAUCCGAGCACCAAGCUGA	1	no match			
8472	UGAUUUGAUAUCCGUGGACCAGUACA	1	1	transposon	DNA	transib2
8473	AGCUUCAUCCAAGAGAACUGUGGCCG	1	1	transposon	LINE	R1
8474	UUGAAUGAAAACCCGGCCUCUGUCAU	1	1	transposon	LTR	micropia
8475	CCCCAGUGGCCUAAUGGAUAAGGCAUC	1	0	ncRNA		tRNA
8476	UUCACCAAGCGUUGGAUUGUUCACCCA	1	0	ncRNA		rRNA
8477	GGCUCUGAAGAUUGAGAUAGUCGG	1	0	ncRNA		rRNA
8478	UCGGCUCAUCUCAUCCUGGGGCUG	1	no match			
8479	UACAAUAUCAGAAGUGGGCCGAUCA	1	0	transposon	LTR	rooA
8480	GGCUCGUCGCAUCCUGGGGCUGG	1	no match			
8481	ACUUUGAGGGGAUCCUCAA	1	2	coding gene		CG2675-RA|Csat
8482	CCCGAAUUGAACUCAUUAAGUGUAC	1	0	transposon	LTR	rooA
8483	UGCUAGAGGAGAGUGUGCGCUUGGC	1	0	no annotation		
8484	UCAGCAGCGGGAUGAUCAGUAGCACC	1	0	transposon	LINE	R1-element
8485	ACACAAGUUAAGGAUGGAGGAUCUCG	1	0	no annotation		
8486	GCGGUGAAUACGUUCCCGGGCCUUGU	1	no match			
8487	AGUACGAUCCGUCUGCAAACUUCCUC	1	0	transposon	LTR	opus
8488	GCCUUACGAAACGAAACCCGAUCAGA	1	0	transposon	LTR	gypsy
8489	UAACUACAUGACGCGACAAGUUCUGG	1	no match			
8490	CCGUACUUUACUUUUGCACAAAAC	1	2	no annotation		
8491	CACCAUGAUAUCGCGUACCCUUAAA	1	0	transposon	LINE	accord2
8492	GGUCCUGAGAGAUGGGCGAGCGCUGUU	1	no match			
8493	CGGUCUAUUCGCCAGGUCCCGAUG	1	0	transposon	LINE	Cr1a
8494	CCAUCAUUCAAAGCCAGCUGGAUGUCC	1	1	transposon	LINE	G
8495	AAACUACCAUAGCAAUCUUCACACGC	1	2	transposon	Low_complexity	F-element
8496	GUGUUCAAGUCCCGUCCUCGUUACCA	1	no match			
8497	GCGUACCGGAGUGGACUGCCGUU	1	0	ncRNA		7SLRNA-RA
8498	CCAACUCCGAUUGCAGCAACAUGC	1	0	transposon	LTR	297
8499	UGUUUCAUCGUUAGACGGCUCGCGC	1	1	AT-chX		AT-chX-B
8500	ACCCCUCCCACGGCGCGACCGCUCUC	1	no match			
8501	GCGGGUGUUGACGCGAUGUGAUU	1	2	ncRNA		rRNA
8502	CAACAAAGCACCUGGACAUGACGUCA	1	0	transposon	Simple_repeat	I
8503	UGGGUCGGGGUUUCGUACGUAG	1	no match			
8504	GGUUCGAGCCCCAUCAGCCACCCCA	1	no match			
8505	AGAACCAUCACUGGGGCACCGUGGUA	1	0	transposon	Low_complexity	F-element
8506	GACGGGCCUGAUGUUCGUUGAUCGCCA	1	no match			
8507	CUGUAGCCUUCAUUCAUGUUGGC	1	1	ncRNA		rRNA
8508	GGUGCGCUUGUGACGGCCCGUGUA	1	no match			
8509	GGGAAACCCUGAUGGAAGACCGAAA	1	0	ncRNA		rRNA
8510	AAGUUAACUAUCCCAAAGGCAUCACGC	1	0	no annotation		
8511	UGAAAACACAUCAGAACCCAUGGAG	1	0	transposon	LTR	springer
8512	UGCUCUCGAAUGUAUUCCCGAUAUACG	1	1	transposon	LINE	R1-element
8513	GCUGGGUCGGUCGGGCUGGGGCGA	1	no match			
8514	AUAGUAAUGAUUCCCAAGCCAGGCAAU	1	1	transposon	LINE	Doc2
8515	AUCGCCAAUUCGAGGACCCCAUGGAAC	1	0	coding gene		CG6691-RA|ttm3
8516	AAGACAAGAACAGCUCGAUACGAA	1	0	transposon	LTR	invader6
8517	CAAACCAAUCCCAACCGAACCCGUC	1	1	transposon	LINE	G2
8518	GGCUUUGAGAGGCGCCGUAGUUACCACCCC	1	0	no annotation		
8519	AAUAGAAGCAGUUUGGGGGCAUUAG	1	0	ncRNA		rRNA
8520	CACAUCAACAAUAAGCGUAUCCAGU	1	1	transposon	LINE	G2
8521	GCGCUCUCCGCUCUCUCGCUCCCUAA	1	0	transposon	DNA	1360
8522	GGUCCGAGGAUCGCUGUAACUCGAU	1	no match			
8523	UACAAGCAAGACCACCACACCCGAC	1	0	transposon	LTR	gypsy12
8524	UGUGGGCCGGUACUAUUACUUUG	1	0	ncRNA		rRNA
8525	UCCGGGAGGGUGUGAUUGGAAGGUC	1	0	transposon	LINE	G
8526	GCAUUCAAUACAGCGAGAUGGCCCAAU	1	0	transposon	LTR	Quasimodo
8527	GAUAACGAACGAGACUCAAAUAUAUUAA	1	0	ncRNA		rRNA
8528	ACGAUCAAUAAAACAUACGCCCGGUUU	1	2	transposon	LTR	GATE
8529	UACCCCAUCCCAGAUAUAAACGCUACGC	1	0	transposon	LTR	opus
8530	UGUUUCGCAAAAGUUGACCGAACUUG	1	0	ncRNA		rRNA
8531	GACAACAACAACUGGAUGGUCCAUG	1	0	transposon	LINE	F
8532	AAACUGAAUAAACGAAAUGGAUGACAC	1	0	transposon	LTR	invader4
8533	ACGUCUAUAAUUGAUACCUACGGCAC	1	0	transposon	DNA	1360
8534	GUUGCAAUUAAACAGAGUUCCGCAGUGC	1	no match			
8535	CAAGCCAAUCUAGACGUACGGAAUAC	1	0	transposon	LINE	G2
8536	UGGCUCUCUCAGUCGCUUCCCGGUA	1	1	ncRNA		rRNA
8537	GAACCCUGACCUACGAGUCCGUCCAUCA	1	0	coding gene		CG6255-RA|CG6255
8538	UCAACGAGCAUCAGCCAUCAUCCA	1	0	transposon	LINE	G2
8539	AGAGCUGAGAGCCCGCAGACUGAUUUU	1	0	ncRNA		alphagamma-element
8540	GGCUACCCACCCGACCCGUCUUGUAAC	1	no match			
8541	AGACGCAGGACGGUGGCCAUGGAAGU	1	no match			
8542	CCACCAGGUGAUCCUGAGCAAGGG	1	no match			
8543	CCAUGGUUGCAACGGGUAACGGGG	1	0	ncRNA		rRNA
8544	CCACAUCUAAGGAAGGCAGCAGGCGCGU	1	0	ncRNA		rRNA
8545	CCGAGGCCCUGUAACUUGGAAUGAGUA	1	no match			
8546	AACUGAAUAACGAAAUGGAUGAC	1	no match			

8547	CGUAACUUCGGGAUAACGGAUUGGCUCU	1	no match			
8548	AUGGCUCUCCCAGUCGCUUCCCGGGA	1	1	ncRNA		rRNA
8549	CCGAUACAUCAGGUCAGAGCGUG	1	0	repeat		trf
8550	GCCUGAUGACCAUAGCGAGUUGGUACCAC	1	no match			
8551	CUACGUCCGUUGGAUUAUGCCUGAAC	1	0	ncRNA		rRNA
8552	UAGUUUCAUCGUUAGACGGCCUCGGGCGU	1	no match			
8553	GCAGAAGUUCGAAGGCAGUUUUCA	1	0	transposon	LTR	flea
8554	UUCGAGAAAAAUUGACGAAUUGCUGA	1	0	transposon	LTR	invader2
8555	GGCGGUAAUCCCAAGCCGGAGCUUUCU	1	0	coding gene		CG11444-RA|CG11444
8556	CCAGAGAAUAUGAGCAGCUAGAGA	1	0	transposon	LTR	3S18
8557	AUUGAGCGAACCCGGGGAACUGAAA	1	no match			
8558	UCCAUCUCGAAUACGUCACCAAAGC	1	0	transposon	LTR	ninja-Dsim-like
8559	AGGUACGGACCAAUGGCACAAUA	1	0	repeat		LSU-rRNA_Dme
8560	CAACACCCAAAAUAACAAAUAAUAAA	1	1	transposon	Low_complexity	F-element
8561	UCAAGAAUAACAGAACCAUCCAAGA	1	1	no annotation		
8562	AAAGAGAGCAAUUUUGGCCGUCA	1	0	transposon	LINE	Cr1a
8563	GGGUCGAAACGAUCUUAACCUAUUCUCAA	1	0	ncRNA		rRNA
8564	CGGAUGUGACCAAUAUCAAGGCUC	1	0	transposon	LINE	G6
8565	ACCUGAAGAUGAUCCUGAGUC	1	0	transposon	LINE	Helena
8566	CAGUCGCGAUAUUGGAAUCCCCG	1	0	transposon	LTR	diver2
8567	GUCUGGUUAAUUCCGAUAACGAACGU	1	1	ncRNA		rRNA
8568	UGGGAACACCGCGUGUCGUUGGCCU	1	1	ncRNA		5SrRNA
8569	UGAAACUGCUUAGAGGUUAAGCCCGA	1	0	ncRNA		rRNA
8570	GGGUGAAUCCUUUGUAGACGACUU	1	no match			
8571	AUUGAAUUAUUUAGUGAGGUCUCCGGA	1	0	ncRNA		rRNA
8572	UAGAGGUGAACCCGAUUGAAGUAUCA	1	no match			
8573	AAAAAAACCUGAAGAUGAUCCUGAGUCGC	1	no match			
8574	UCCAGACGUAAUACUCCACUACCCGU	1	1	transposon	LTR	HMS-Beagle
8575	AUCCUCAACUGUGAGGGGAUCCUAA	1	no match			
8576	CAGAUCAACAAAGUCCGAUGGGG	1	1	no annotation		
8577	GGGGGAAUGUCCCGAGCGGCAAAGGGGG	1	no match			
8578	GCACACAGAGGAGAGCAGAUGAGAG	1	no match			
8579	UAGUACAUUAAUUACUCGAUGACGC	1	0	transposon	LTR	invader4
8580	ACCGAGAGACGCCGGCCGCUAC	1	no match			
8581	CUCAUCGUCGUAGAACAAGCCCGA	1	0	Su(Ste)		Su(Ste)
8582	UGCUCCUGAUGGCGAGUGGCGAACGGG	1	no match			
8583	GUCCUGAAGAUCUAUCCCAGCCUAAG	1	1	transposon	LTR	Max
8584	GCGUCAUCGUGUGAACUGCAAGACACA	1	1	ncRNA		rRNA
8585	GCCGGGGAUUGUGGGUUCGGGUCCCAUCU	1	1	ncRNA		tRNA
8586	ACAGAGGGCAACCUACGGGGCAUUA	1	0	transposon	LINE	R1-element
8587	AACGCAUGGACGACUCAUUGGCUGUAA	1	0	coding gene		CG11719-RA|Mst98Ca
8588	AACGCAUAAUGUGGGUUGGGCCA	1	no match			
8589	CAGAUAUGACCAAAGCAAAGACA	1	1	transposon	LINE	Cr1a
8590	CUUGAUUGCGACGGACUUAGGGAA	1	0	transposon	LTR	HMS-Beagle
8591	AUCGCAAAGAAGCUGUUUUUGAAGGAC	1	no match			
8592	AAUAGUGAAAUCCUGGUACAGAACCGAC	1	0	transposon	LINE	Circe
8593	ACGGAAGCGAUCCCACUGCUGAAGA	1	1	AT-chX		AT-chX-A
8594	UGGUACGAUACGGAAGAUGGCCCA	1	0	transposon	LINE	F
8595	AGCGUGCUGGAGGUAUCAGAACUGC	1	no match			
8596	GUCGCUACCGGGUCUUCGUCUGCAC	1	no match			
8597	GGCGCUAAUCCAGGACGACGCAGU	1	1	transposon	LTR	blood
8598	CACACAAGAUGACGACAUUAAUGCAACC	1	1	transposon	LINE	G2
8599	GACACCGAAUUAGUCGACAAUGAGAA	1	0	transposon	LTR	micropia
8600	AGAGCGUCCGACGAAUACCCUGGGGGG	1	no match			
8601	UCUGUUUCAUCGUUAGACGGCUCGGGC	1	0	AT-chX		AT-chX-B
8602	CUCACAAUUAGGUGGGUCCCGGGACAA	1	0	transposon	LINE	baggins
8603	CGCGGAAGACGGAAAGACCCCAUG	1	no match			
8604	UGAUUUGUCUUGAGUAUAUUCCUAC	1	0	transposon	LINE	Cr1a
8605	CACUCAACUCUUCGCCCACCGACU	1	0	transposon	Low_complexity	F-element
8606	UCGUCCGGUAUGACAUAAACCAGUU	1	0	no annotation		
8607	AAGGCCUUGAUGCCACUCAAAUCGUC	1	0	Su(Ste)		Su(Ste)
8608	UUUCCCGAAAUCCGAGUGAAUAUGGC	1	0	no annotation		
8609	CCGUGAUCGUCUAGUGGUUAGGACCCC	1	0	ncRNA		tRNA
8610	UAUUAACUACUGGUAUGACUGAGGA	1	0	transposon	DNA	transib2
8611	AUGUAGUGACGAAAAAUAACGAUACG	1	no match			
8612	AGCACUCGCCGAAUCCCGGGGCCGAGG	1	no match			
8613	GGGCUAAUGACGGUACCGUAAG	1	no match			
8614	GAAACCGAAAAUCACAAUUUGCC	1	no match			
8615	CUUGAAAUGAUGAUGACUAAGGUCGCA	1	0	no annotation		
8616	ACCACCCUGACCGCGCCCGAUCUCGU	1	no match			
8617	CAGACAUCUACUUCCCCCCCGCUCCUAU	1	1	transposon	LTR	invader1
8618	CCAGGGAGUUCUUCGAUGUGGCAGCGUU	1	no match			
8619	GUAACUUCGGGAUAAGGAUUGG	1	0	ncRNA		rRNA
8620	CAGACAAAGACUCCGCGUUUAUAAGCAC	1	0	transposon	LTR	Idefix
8621	GCGAGAAUAAACGAAAUACAGUGGGAAA	1	1	transposon	LTR	diver2
8622	UUCCGAGGAACCAUUGAUCCAGCUGC	1	no match			
8623	GACCGGCAACGACGUUCCACACGCA	1	0	no annotation		
8624	UCCAAAAGAAACGCACGCGAAAAGUAA	1	1	no annotation		
8625	GGUGAUGUCAACGUUUGAUCCCCUGGA	1	0	transposon	LTR	ninja-Dsim-like
8626	ACAAUCUUAAACGUAAGGUGAUGGA	1	1	no annotation		
8627	UAGAAAUAUAAAAUGUAUCCGGAGCG	1	1	transposon	LTR	297
8628	AAACUGAAUAAACGAAAUGGAUGACACAC	1	0	transposon	LTR	invader4
8629	UGGAUUAGGACUCAGAAUUUGAAGA	1	0	no annotation		

8630	AGGUCUCCGGACGUGAUCACUGUGAC	1	0	ncRNA		rRNA
8631	GAGAUAUUGAGCGACUUUCCCCGUAG	1	0	transposon	LTR	3S18
8632	CCCGAAAAAUCCCCGGCUGCGAC	1	no match			
8633	UGCGUCGGGUCUUCGAGCGGUGCGGC	1	0	coding gene		CG5442-RA|SC35
8634	AGCCGUCGAGAGAUAUCGGAAGAGU	1	0	ncRNA		rRNA
8635	GUCGGAGGCUGGCGCGGGGGGGU	1	no match			
8636	GAGGCAAUCACCGAGAGCAUCAACG	1	0	transposon	LINE	F
8637	CGGGGUUGCGAGCAAGUUGACCA	1	0	no annotation		
8638	AAGAUUUGAUGUGUAAACUGAAC	1	0	coding gene		CG9603-RA|CG9603
8639	AAACUUCGGGAUAAGGAGUUGGCUCU	1	no match			
8640	CGGGUAAAGAAAACCACGUAGCCGA	1	0	transposon	LTR	FB
8641	ACGGAAUGGAACCGAGCUCUGAA	1	1	transposon	LINE	X
8642	UGCACAGAGACCUGAAACUCGCAAC	1	0	transposon	LINE	Doc3
8643	CCUGCUCUCCCGCUCUAUUGGCUCCAA	1	0	transposon	LINE	F
8644	ACCCUGAACGCGCCCGAUCUCGUAAA	1	no match			
8645	GAAGCUUAUCCUUUGCUUGAUGAUUCG	1	0	ncRNA		rRNA
8646	UCGGAUACACGAAGCUGCACCACCACC	1	0	coding gene		CG8416-RA|Rho1
8647	AUGGAACACAUAGAGGCUUCGAG	1	0	transposon	LTR	HMS-Beagle
8648	UCUCAUCGUCGUAGAAGAAGCCCGAGG	1	1	Su(Ste)		Su(Ste)
8649	AAGAAUCCGGAUGGUAUUCGACCCUUUA	1	0	coding gene		CG32240-RA|CG32240
8650	CAAGAUAUGACCAAAACAAAGCAACGU	1	0	Su(Ste)		Su(Ste)
8651	GCCGGGCCGUACCCAUAUCCGCAGC	1	no match			
8652	GCCUCCAAGACGAGCGUUGCUUCAC	1	no match			
8653	AUUACGACGCGAGAGGUGAAAUUCUU	1	0	ncRNA		rRNA
8654	GCGAGAACCGGGAUAUUAAUUUGG	1	0	coding gene		CG6815-RA|bor
8655	ACUGCCCGACGAGCAAACCCGCGUA	1	2	coding gene		CG5977-RA|spas
8656	GUAUGACCCCUGGUACUGAAAUGC	1	0	transposon	DNA	Bari1
8657	GCCCCCGUACGUGAAAUGACCUGUGUG	1	no match			
8658	ACUGAGGUGCGAUUAUUGCUAGUUGAA	1	0	ncRNA		snRNA
8659	AAAAACUAAACUAUAAAGCCGGGG	1	0	transposon	LINE	G2
8660	ACACGGAAAACUACCAGGUCCGAACA	1	no match			
8661	CCAGUCAGUAGAGCACUACCGCCGCC	1	1	repeat		trf
8662	GUGGAGCCGGGCGUGGAAUGCGAG	1	no match			
8663	ACACAGAGUACGGGGCUUACUGAAGUCA	1	1	transposon	LINE	baggins
8664	UCCGAGGAACCAAUUGAUCCAGCUGCU	1	0	Su(Ste)		Su(Ste)
8665	CCUGGCGGCGAGUGGCGAACGGGUGA	1	no match			
8666	UAGAAUAUUAACUUCUGUAAACGGCGGC	1	0	transposon	LTR	mdg3
8667	AACUGAAUAAACGAAUGGAUGAC	1	no match			
8668	GCCUGAUGACCAUAGCGAGUUGG	1	no match			
8669	AGAGGGACGGCCGGGGGCAUUCGUAU	1	no match			
8670	UGCUCUCGGAAUGUAUGCCCGAUAUAC	1	no match			
8671	GAGCCAGUCCCAGAAGAAGCAGC	1	0	coding gene		CG10869-RA|CG10869
8672	UAUACUCGAAUGCCAUUUGGUUU	1	0	transposon	LTR	Quasimodo
8673	ACGUGCAACAAACCCCGACUUCCGG	1	no match			
8674	ACUGGACGAGGCUACGACAGCAGAG	1	0	transposon	LINE	Rt1b
8675	UGAUGACCGAAAUUUGGAAGACGGA	1	1	repeat		SAR_DM
8676	AGUGAGGACAGCCCUAGAUGAACAA	1	1	transposon	LINE	baggins
8677	UCGGCUCAUCACAUCCUGGGGCUGG	1	no match			
8678	CUAGACUCGAAAAUUCGUUUUAUG	1	0	Su(Ste)		Su(Ste)
8679	CGGCGGGGAAGGUGGCUCGGGGGGC	1	no match			
8680	CUUCUCGUUGUUGAAGUGAUUGUUG	1	0	no annotation		
8681	CCCCAAAAGCAGACGAGGUAACGUACC	1	2	transposon	Low_complexity	F-element
8682	GAUUCCGAAGAGGGAGCCUGAGAA	1	1	ncRNA		rRNA
8683	UGAAAAAUGAUCCCGUAUGCGAGAGU	1	0	repeat		HETRP_DM
8684	UCCCUUCUGAACGAGUUCGAUAUAA	1	0	transposon	DNA	H
8685	GCAUCCUAGAAUCGACCGACCAGCUA	1	0	transposon	LTR	flea
8686	UCCAAUAUCACCGCGGAGACUGUCGUA	1	1	transposon	DNA	1360
8687	AAGACGUCCAGAGGAGGCGAGAAUU	1	0	AT-chX		AT-chX-B
8688	CUGCUUUACAUACCCUUUGGUGCCC	1	0	transposon	LINE	Circe
8689	GUGGCAAGAUGCGAUGCCGCCGCCCCAG	1	1	coding gene		CG4994-RA|Mpcp
8690	UCCGAAUCAAAUGCUACGUAACCAAG	1	0	transposon	LTR	diver2
8691	UUGGGUGCGAGAGGCCCUGGGUUC	1	2	ncRNA		tRNA
8692	CAAGCUAGACUAUGUAUCCCCGGGCA	1	1	transposon	LINE	Doc
8693	UGAUUGCUAACAAGAACUCGGAGGA	1	1	transposon	LTR	flea
8694	GUCAACUGUGAGGGGAUCCUCAA	1	no match			
8695	CACGUACUCAGACAAGAUAUCCCUA	1	0	transposon	LTR	gypsy
8696	ACCACCCUGAACGCGACCGAUCU	1	no match			
8697	CAACCCAAUACACACGACUGACAC	1	0	transposon	LINE	Cr1a
8698	CGAAUCGAGACCGGGCCAGAAA	1	0	transposon	LTR	G4
8699	ACCAAGCUUUUGGGCUCCGGGGG	1	2	ncRNA		rRNA
8700	UCAUUCGAGACCCAACGGCACGUGA	1	1	transposon	LINE	INE-1
8701	GAAACUCCAUAAAAAUCAUUGCCGCC	1	0	transposon	LTR	accord
8702	UGCUCUCGAACGUAUGCCCGAUAU	1	1	transposon	LINE	R1-element
8703	AUGAUCUCGGAAGCUAAGCAGGGUC	1	no match			
8704	GGAAUAAUGGAAUAGGACCGCGGUU	1	no match			
8705	AAAAGACAGGAGAGGCUUCGUACGCAG	1	0	transposon	LINE	Rt1b
8706	AAGACGACGUAUAGGGCCUGACGCCU	1	no match			
8707	GACAAGCGCAAGGGCGGUCCCGGU	1	0	coding gene		CG3481-RC|Adh
8708	UGGCCGAUGGAGUGGGCGGUUGGCGA	1	0	coding gene		CG15035-RA|CG15035
8709	CUCGGAGUUAGAGUACAUGAGGUACA	1	1	transposon	LINE	R1-element
8710	GAUACUUUUAUCAGAGGCCUUCGAGGA	1	0	transposon	LTR	HMS-Beagle
8711	UCGAGCGAACAGUGGUAUGACGCUACGU	1	0	ncRNA		rRNA
8712	AGAGACAGGAUCAUAUGCCGCCGCC	1	0	transposon	LTR	springer

8713	AGGGUCGGGUAGUGAGGGCCUU	1	no match			
8714	GCAACAUCCAGACCAAACGAGUCCAA	1	0	transposon	LINE	Rt1b
8715	GGCUCCGGGGGGAAGUAUGGUUGCA	1	no match			
8716	AGAGAGUGGACGGGAGCGGCGGGGGCG	1	no match			
8717	CACGAAAUGGACCCGAACGCACUGAG	1	0	transposon	LINE	Doc3
8718	CAUCUGUGGGAUUAUGACUGAACGC	1	no match			
8719	CAGGCCGAGAUUGAUCGCAUGGUGA	1	0	coding gene		CG18743-RA|Hsp70Ab
8720	GAUUCCGGAGAGGGAGCCUGAGUA	1	1	ncRNA		rRNA
8721	UGACUCUCGAAUGUAUGCCCGAUAUUU	1	no match			
8722	UCGAACGAUAAACGGUGCUAACCAGU	1	1	transposon	LTR	GATE
8723	UCCGCCCCCUGAACGGAACUCAGC	1	0	no annotation		
8724	GACUCUUAAUCUCAGGGUCGUGGGUU	1	0	ncRNA		tRNA
8725	UCGAGGGAGACGUCAGCGGGUACAUA	1	1	coding gene		CG13898-RA|CG13898
8726	AAGCACGGUACCGAAGCGGCGGAC	1	no match			
8727	ACUCGCCGUAUCCCGGGGCCGAGGGA	1	no match			
8728	AAGAGGGACGGCCGGGGGCAUUCGUAU	1	no match			
8729	CCUCAACUGUGAGGGGAUCCUCUAA	1	no match			
8730	UGCCUCAACAUUUGGUGACCUCGAUA	1	2	no annotation		
8731	AAGGCUGGAAUGAAUGGUUGGACGAA	1	0	ncRNA		mt
8732	GCCUUACGAAACGAAACCCGAUCAACA	1	1	transposon	LTR	gypsy
8733	CUUCGACCCCGCCAGUGAAACCAA	1	0	transposon	LINE	Doc
8734	CAGCCAAGAGUUCCCGUGGUAGGCAGUA	1	0	ncRNA		7SLRNA-RA
8735	GAAUGGAGCGAGGCUGCCCAGGAGU	1	0	coding gene		CG31501-RA|nxf4
8736	GAAGGUGGUUUUCCCAGGGCGAGGC	1	no match			
8737	ACUCCGGAUGUGCUGACCCCUGCGAU	1	no match			
8738	AACUUCGGGAGUAAGGAUUGGCGUCU	1	no match			
8739	ACCUGAUGACUCUUUGACAUAUA	1	2	transposon	LTR	Dm88
8740	UGGACAAAAAAAAACGAAUGACACAG	1	no match			
8741	CAACCAGAGAUCUAGAGGGCCCAAGAG	1	1	no annotation		
8742	AGCGAGGAUAGGCCCGUAAAAAACAGCC	1	1	coding gene		CG14472-RA|poe
8743	AAUCCGGACAAACACACAAGACC	1	0	transposon	LTR	opus
8744	AGACAGAGCAAAACCCAUGACCAGC	1	0	transposon	LTR	297
8745	UGAAUCGUGCAUGCCACCUACGGACC	1	1	transposon	LINE	Rt1b
8746	GCUGUAUACCGAAGAUGUGAGUGCU	1	no match			
8747	UACAGAGUCGAAGCGGUUACAAGA	1	0	transposon	LINE	Rt1b
8748	CCGAACUUGAGACCGCCCUGGUAC	1	no match			
8749	AGACGAAUUUGUGGGCCGGUAUU	1	2	transposon	LTR	Rt1c
8750	AAAACCUGAAGAUGAUCCUGAGUAU	1	2	transposon	LINE	Helena
8751	CAGUGACGGAGCCGGACGACCUC	1	0	transposon	LTR	Quasimodo
8752	GCCCGGCUAGCUCAGUCGGUAGAGCAUGUA	1	2	ncRNA		tRNA
8753	UCGAGAAGCGAACCCGGCGAACUGAA	1	no match			
8754	UGUGCUAGAAGACGAUUUUUCGGACCG	1	no match			
8755	AAGAAUCUCCGCAAAGCCUGAGGA	1	0	transposon	LTR	diver2
8756	AAAUUACAAAGACGACGGCCCACC	1	0	transposon	LTR	springer
8757	GAAGAAGAGAUGUCAGGUGAGUU	1	0	transposon	LTR	ninja-Dsim-like
8758	UGUUCCACAUGGCGCUGGUUUACGGG	1	0	coding gene		CG18418-RA|CG18418
8759	UCCGAGGAACCAACUGAUCCAGCUGC	1	1	Su(Ste)		Su(Ste)
8760	AGCGCAACCCUUGUCAUUAGUUGCU	1	no match			
8761	UCCGAUUUCAAGUACCAGACAAACA	1	0	repeat		RSP
8762	CGUAUAACAACUACUUCUCGGUGA	1	0	transposon	LTR	accord
8763	CUCGCCGAAUCCCGGGGCCGAGGGAGUC	1	no match			
8764	AAACUUCGGGAUAAGGAUUGGCUCUGA	1	1	ncRNA		rRNA
8765	CACGAUGAUCGAUGACGAGCUACU	1	1	transposon	LTR	ninja-Dsim-like
8766	CAAGCUAGACUAUCUAUCCCCGGGU	1	1	transposon	LINE	Doc
8767	UCAUACGGGUAGUACAACUUACA	1	0	ncRNA		rRNA
8768	CGCUCGAAAAAAGGUUAUACCUAAGUUG	1	0	transposon	LTR	opus
8769	GGCGGGUGUUGACGCGAUGUGAUUA	1	no match			
8770	GUGGAGCCGUACCUGUUGGUUUGUC	1	0	ncRNA		rRNA
8771	GUUCGAGUCCGACCCUCGGUACCA	1	no match			
8772	ACUACCAUCCCACGGAACACCAGAGCAAUGCC	1	no match			
8773	UGCAACGAACCGAACGGAGUCCAC	1	0	no annotation		
8774	CAGACAGAGAGGCUCAUUUACAACCAC	1	0	transposon	LTR	McClintock
8775	CGAACAACUAGAGCACCGCAUCCGG	1	0	transposon	LTR	Tirant
8776	CGAAAUGAUCGAGUGGAUAUACAGUA	1	no match			
8777	CAUCGCGAUGGAUCUGGAGGCCC	1	no match			
8778	UGAUGAGACAUUCCCGUUUCUCGCGAC	1	1	transposon	LTR	rover
8779	CCUACUCGUAGCCGAAACCCACUUC	1	0	transposon	LINE	G6
8780	AUGGUUUCAACGGGUAACGGGGAAU	1	1	ncRNA		rRNA
8781	UGAUAAGACAUUCCCGUUUCUCGCGAC	1	1	transposon	LTR	rover
8782	GCCGGAAGCGAUCCCUCUGCUGAAGA	1	1	AT-chX		AT-chX-A
8783	UCAUUGCUAUGUUGACGCGGAUU	1	1	transposon	LTR	DM88
8784	CAACCUAUGACAAAUCAAAACCAGUA	1	0	transposon	LTR	Idefix
8785	CAAAGUAUCCGAAUGCCCUGGCGCA	1	0	coding gene		CG41423-RA|CG41423
8786	UUCCUCGGAAUCAAGGCCAACGAGU	1	1	Su(Ste)		Su(Ste)
8787	AAAAGGACAAUUGACGGUACCGCAU	1	no match			
8788	AUCUCACGGAACACCCGAGCAAUGCCACC	1	1	transposon	LINE	G2
8789	AGAAAGAUAAAAUGUAUCCGGAGCGA	1	0	transposon	LTR	297
8790	UCGAUAUUAUCCCUGCACUACGAAGC	1	1	no annotation		
8791	CCAACGAACGGUUAAGCGCGACAUC	1	0	transposon	LTR	HMS-Beagle
8792	GCUUCACUUGACCACGAACCGGCCUUCC	1	no match			
8793	UGGUGAACGACGGCGACAGCC	1	no match			
8794	AUGGACAGAUAGAAAGAUUUCACUCCA	1	0	transposon	LTR	HMS-Beagle
8795	AACGAGCGCAACCCUCGCCAUUAGU	1	no match			

8796	GCUCCGAGGUGAUUUUUCAUAUUG	1	no match			
8797	GCGUUCGUUGACGUAGAUGGCUGC	1	1	transposon	LTR	HMS-Beagle
8798	AAGACGAGAAGACCCUAUGGAGCUU	1	no match			
8799	GCAGAACUCCGAUGGCGAUAGGAA	1	0	no annotation		
8800	CAAAGCAUUGUGAUGGCCCUAGCGGU	1	1	ncRNA		rRNA
8801	AACGAGAAUAUACACCGAGACUUACA	1	0	transposon	LINE	Doc3
8802	CCAUCAGAGACUGCGGUGGCCAAC	1	0	coding gene		CG12717-RA|CG12717
8803	GAAAGAUGGUGAACUAUGCCUGGGC	1	no match			
8804	CGGCUCCGGGACGGCUGGGAAGGCC	1	no match			
8805	CCGGGUGGCGGAAUGGUAGACGCGC	1	no match			
8806	GCUAACGAUAUCCACAUUCGCUGA	1	0	transposon	LINE	Doc3
8807	AACGAUUCUAUUUUUCCGAACCC	1	2	transposon	DNA	FB
8808	CACCCUGAACGCGCCCGAUCUCGUU	1	no match			
8809	ACGAGCGCACGUGUUAGUACCCGAAA	1	no match			
8810	AAGGCUGAGGCGUGAUGACGAGGCACU	1	no match			
8811	ACAUUUGAGAGUGUUUGGUGCAA	1	0	transposon	LINE	Cr1a
8812	UACGAGAUUUGUGUUAGCCACUGAGA	1	0	transposon	LTR	G4
8813	AACUAUAAAAGCGAGAAGGUUGA	1	0	transposon	LINE	INE-1
8814	GAGUACUUCAACCAGACACUGGAC	1	0	no annotation		
8815	UAGAAGCAGUUUGGGGGGCAUUAGUAUU	1	no match			
8816	AACCUCGGGAUAAGGAUUGGCUCU	1	1	ncRNA		rRNA
8817	CCGAGCUUGAGUUUGCCGAUGUUU	1	no match			
8818	GGAAGCCGAGAUGUGCACCACGUGCCCC	1	1	coding gene		CG3982-RA|CG3982
8819	CCGAAAGUAUAUGAGAAGAACUACC	1	no match			
8820	UCCGACGACUGUGCUACUCCAUCCAAA	1	0	transposon	LTR	GATE
8821	UGCGCAGAGUGUUGCCAGUACUGGAU	1	0	transposon	LINE	G
8822	CGUUCUGGGUAAAUAGAGUUUC	1	0	ncRNA		rRNA
8823	ACAAACUUAACUAUGUUCCCCGGGA	1	0	transposon	LINE	Doc3
8824	UGCCUCAACAUUUGGUGACCUCG	1	0	no annotation		
8825	GGUCAAGUUAAGAAGCGCAUACGGUGG	1	no match			
8826	GAGGCCCUCGUAAUUGGAAUGAGUA	1	no match			
8827	GCCAACGAUCGAGCGUACUUGUCCGUGA	1	no match			
8828	UUGGCAUUACUCGGUACUGAUUU	1	1	no annotation		
8829	GGGUGCGAGAGGCCCUGGGUUC	1	2	ncRNA		tRNA
8830	CAUCCUGGCAACAGGAACGACCAUAA	1	0	ncRNA		rRNA
8831	CGAGGCCCUGUAAUUGGAAUGAGU	1	0	ncRNA		rRNA
8832	UUGUUCCCGGCGGUGGCGUCGUCCC	1	no match			
8833	GCGGAUGUGACCUGCGUUCUGGGUAC	1	0	no annotation		
8834	CCCCCAGACACUCGAUCACCGAGAC	1	0	transposon	LINE	G2
8835	UCAACCGGACGGGCCUAGAGUUAUUCC	1	0	coding gene		CG40635-RA|CG40635
8836	AUAAGGAUGAUGAUACCGACUAAAGGC	1	0	no annotation		
8837	AAACGGACUUCCCCCGAUUCGGAUA	1	no match			
8838	CGUGACAGGACGAUUGCAGGGCGUA	1	2	no annotation		
8839	UACUGAGACAUGGUGACAUCGGCCAA	1	0	no annotation		
8840	UGUGGAUCGGAGUGAAAGGCUAAUCA	1	no match			
8841	ACAACAAACAACACAUACCCGCUGGCCA	1	0	transposon	LTR	GATE
8842	UCCCAGCUCGGGGAGGUGGUGACGAAAA	1	1	ncRNA		rRNA
8843	GAGAAGCAGCAAUGGACAUCCCGUA	1	2	coding gene		CG15530-RA|CG15530
8844	AUAGGGAACGUGAGCUGGGUUU	1	2	ncRNA		rRNA
8845	AGCCUGAGAGGUGAUGCGUAGCCG	1	no match			
8846	GCUGCGGUGUCCGAGUGGUUAAGGU	1	1	ncRNA		tRNA
8847	GCAAUAACUGUAUUUAGCCCUGAGAA	1	0	transposon	LTR	GATE
8848	UGACAUUGUGUGGGAUCCUUCGAUA	1	0	transposon	LINE	G2
8849	UCGAUGCUGACCCCCUACCCCCACC	1	0	transposon	LINE	R1-element
8850	ACCAACCAGACUGACCUCUACCAAAU	1	2	transposon	LINE	F
8851	UAACCGGGGCUAAGCAUAUUACCGA	1	no match			
8852	UGCGUCGAGGCCCAGGAUUGCG	1	0	transposon	LINE	R1-element
8853	CUCAGGACUCUGAAUCCUGCGAUCCGA	1	0	ncRNA		tRNA
8854	GUAACAACGAAGAUGAUUCUGGCCCU	1	1	transposon	LTR	invader4
8855	CAACAUAAUGGAAAAUCUCUGGU	1	1	transposon	DNA	Tc1-2
8856	GUCCUGGGCUGCACGCGCGCUACA	1	0	ncRNA		rRNA
8857	AGCCACACUGGGACUGAGACACGGCC	1	no match			
8858	GCACACAGAGGAAGAGCAGAUGAGAGCA	1	no match			
8859	AAGGAGUCUAACACGUGCGCGAGUU	1	no match			
8860	UGGCACAUCUCGACACCCGUGGCACU	1	1	coding gene		CG13340-RA|CG13340
8861	GGAUUCAACUGUGAGGGGAUCCUCAA	1	no match			
8862	CAGCUGCCUGCGACCCACGCCGCC	1	0	coding gene		CG10043-RA|rtGEF
8863	GACAACCAGAGAUCUAGAGGGCC	1	0	no annotation		
8864	CCACCCUGAACGCGCCCGAUCUC	1	no match			
8865	GAACCUGCUGAAUGAUUGCAUCG	1	0	coding gene		CG8318-RB|Nf1
8866	CGACACAUCCGGUCGAUCCAGCAAA	1	2	coding gene		CG15258-RA|CG15258
8867	UGCAAUCGAAACGGGAAACUUCAUC	1	1	coding gene		CG41521-RA|CG41521
8868	UGGGAACACCGCGUGUUGUUGGCCCUU	1	no match			
8869	CGCAAUACUUAGCCUAUCUAGAUGC	1	1	transposon	LINE	Doc2
8870	GUUAAUUCCGAUAACGAACGAGA	1	0	ncRNA		rRNA
8871	GACACUGACAAACGCGAGUAACUACGG	1	0	transposon	LINE	TART-A
8872	ACGAGAGAUAUCCGUCUAGGUGA	1	0	transposon	LTR	micropia
8873	AAGAGAUGGAUAAAAUUUCUCAUCUA	1	0	transposon	LINE	Cr1a
8874	GCCCUGAAUCGUACCGAGCGUAG	1	0	transposon	LINE	Rt1b
8875	CGUAAACGAGUGUUAUUGUGGGCCGG	1	1	ncRNA		rRNA
8876	UCCAAUAUCACCGCGGAGACUGUCGCA	1	1	transposon	LINE	Doc4
8877	GCCCGAAGCGUUUACUUUGAAAAAAUU	1	no match			
8878	GCAUUCGAGACUCAACGGCACGUC	1	0	transposon	LINE	Rt1b

8879	CGCACGCCGUUUGCUGGUGACCCAUGU	1	1	coding gene		CG11253-RA|CG11253
8880	AUCCGCAGAGGGAGCCUGAGAAACGG	1	2	ncRNA		rRNA
8881	AACUUCGGGAUAAGGAUUGGCUCUU	1	1	ncRNA		rRNA
8882	GAAUCGGAAUGAGCGGGAGUGCG	1	0	coding gene		CG13733-RA|CG13733
8883	GGCUUUGAGGGGCGCCGUAGUUACCAC	1	1	no annotation		
8884	GUUCAAUCCCCGGCAUCUCCACCA	1	no match			
8885	GAUCAGAUACCGCCCUAGUUCUAACC	1	0	ncRNA		rRNA
8886	UUCAACUUUAACCCAAGAGACGUGAC	1	0	transposon	DNA	transib4
8887	GCCGAACUUGACGACCCACGAUAA	1	0	Su(Ste)		Su(Ste)
8888	GAUGACUAUAUCAUUCUCGGGAUUAU	1	0	transposon	LTR	gypsy8
8889	AGCAGAACUCCGAUGGCGAUAGCAAC	1	0	no annotation		
8890	GUAGCCAAAUGCCUCGUCAUCUA	1	0	ncRNA		rRNA
8891	AGAGCAUAUAAUGCCCCGAAGGUAGUCCC	1	2	transposon	LINE	R1-element
8892	CAACAACGUUCGCGUGAUGACCC	1	0	transposon	LINE	G6
8893	ACGGAAUGGAAUCGAGCGCUGAGUAGG	1	0	transposon	LINE	X
8894	GCAAAGUGAACACGUCGCUAAGCGAA	1	0	ncRNA		alphagamma-element
8895	AACAUUAACCGAAUUGCCUCGUUGU	1	1	transposon	LTR	gypsy4
8896	AGACUCCGUUAGUAGUGGCGAGCGAAC	1	no match			
8897	GAAUCCAACAAUCCAUAGACCAGGCGA	1	0	transposon	LTR	G4
8898	CCAGAUAGGACGCCGAUUUUCCG	1	0	no annotation		
8899	AACGGGCGAAAGACUAAUCGAACCAUCU	1	no match			
8900	GCCACCCUGAACGCGCCCGAUCUCGUC	1	no match			
8901	UCUCAUCGUCGUAGAACAAACCCGA	1	1	Su(Ste)		Su(Ste)
8902	AAUUGCUAUAACGAACGAGACCUUAACC	1	no match			
8903	GCGGAGAGAAGACAAUUUUGGCC	1	0	transposon	DNA	1360
8904	GUGGCUUUGAGAGGCGCCGUAGUA	1	1	no annotation		
8905	AGCAAACCAAUCCCAACCAACCCCGU	1	1	transposon	LINE	G2
8906	AGAGUGUCCUGGGAGUUCGAUAUGC	1	0	no annotation		
8907	UAACGAGAGAAGCAUUUUUUGCCGC	1	no match			
8908	ACUUCGGGAUAAGGAUUGGCUCUG	1	0	ncRNA		rRNA
8909	GCCGUACCUGUUGGUUUGUCCCAUUAU	1	0	ncRNA		rRNA
8910	CUUAAACGAGUGUUAUUGUGGGCCG	1	0	ncRNA		rRNA
8911	CGGUGGGCGAAGAGUUGAGUGGGGC	1	2	transposon	Low_complexity	F-element
8912	GCCAACGACCAUACCACGCUGAA	1	0	ncRNA		5SrRNA
8913	GAACCUGAGACCGCCCCUGGACGACC	1	1	Su(Ste)		Su(Ste)
8914	GAGCUCCCUGAUCGAUGACUAUC	1	0	coding gene		CG14708-RA|CG14708
8915	UGAGGCGAAUUCAGUGAUCCUAUGCU	1	no match			
8916	UCGAAACAAUAGACAGAGCAACGCC	1	1	transposon	LTR	Rt1c
8917	ACCCUCGAGGUAGAGCCAUCGGACACC	1	0	repeat		trf
8918	GAUACAAGCAAGAGAUGUCCAGU	1	0	transposon	LTR	rooA
8919	GCAGCCAAGAGUUCCCGUGGUAGGCA	1	0	ncRNA		7SLRNA-RA
8920	CCGAAGAUGACCAAGUUGGCAACCGCU	1	no match			
8921	ACGAUAUUAUUUCUGGCCCGGUCUCGA	1	0	transposon	LTR	G4
8922	CCAGCUAAGGUCCCUAAAAUUGGCUA	1	no match			
8923	ACCCUGAACGCGCCCGUAUCUACGUC	1	no match			
8924	AGAUGUCCGGGGCUGCUGCGCGCC	1	no match			
8925	UGUUUCAUCGUCAGACGGCUCGGGC	1	1	AT-chX		AT-chX-B
8926	ACCACCCUGAGCGCGCCCUAUCU	1	no match			
8927	CGCUUUGAAACCGAAGAGACGCCGGC	1	0	transposon	LTR	G4
8928	UGAGUGUCCCGCGGGGCCCGAAGCGUU	1	no match			
8929	UCAAAGAGACGCUUGAUAGCCUAGA	1	no match			
8930	UGCGCGAAGAAGGAGAACGAGAAGG	1	0	transposon	LINE	G6
8931	UGUUUCAUCGUUAGACGGCUCGGGAG	1	1	AT-chX		AT-chX-B
8932	GAGGCAAUCACCGAGAGUACCGAGAU	1	0	no annotation		
8933	AAUACAUGCCGACGGGGCGCUCGACCCCC	1	no match			
8934	GCCACGCAGACGGAGGCACAACCGGUUCC	1	0	coding gene		CG30324-RA|CG30324
8935	GGACCUACUGUGAGGGGAUCCUCAA	1	no match			
8936	UGCCGGAGCGAGGAGGAGGACUAGCU	1	0	Su(Ste)		Su(Ste)
8937	AGUCCCGCAACGAGCGCAACCCUCGCCA	1	no match			
8938	AGGUGGGAGGCUUUGAAGCGUGGACGCC	1	no match			
8939	AACUGAAUAAACGAAAUGGAUGACUAC	1	no match			
8940	GACCAUAUCCGGAGACGCGGCGCUAG	1	1	transposon	LINE	F
8941	GACGUGGCGACCCGCUGAAUUUAAG	1	no match			
8942	ACACAACACACAUACCCGCUGG	1	no match			
8943	GAAGAAGUAAAUAAGCUCAGCCAAGG	1	0	transposon	LTR	Quasimodo
8944	GGAUACUAAGCAGGGACGUUAAUGA	1	0	coding gene		pseudogene
8945	UCGAGGAUGAACUGGGUGACCCU	1	0	transposon	LINE	G2
8946	GGCUUAAUUUGACCCAACACGGGA	1	1	ncRNA		rRNA
8947	UACUCUGGAAGGACGACCACGUUG	1	0	transposon	LTR	ninja-Dsim-like
8948	CGUACUUCGGGAUAAGGAUUGGCUCU	1	no match			
8949	AUGCCGAGUCGUCGUAUCGAGUAAC	1	0	no annotation		
8950	AGCCGGACGGUGGCCAUGGAAGUCGG	1	no match			
8951	GCCAACGACCAUACCACGCUGAAU	1	0	ncRNA		5SrRNA
8952	ACCGAGAAACAAGCCUCCGUC	1	no match			
8953	UUCGAGACAAACGAUACUGUUCGGCC	1	no match			
8954	GCACACAGAGGAGAGCAAAUGAGAGCA	1	no match			
8955	GCCCGUAAAAACGACUUGGUUGA	1	0	transposon	LTR	gypsy
8956	AUGAGACUACAAACGGAUUGCGGACC	1	1	no annotation		
8957	UACCUAAUCCCUAUACCAACGAGCAGC	1	0	coding gene		CG13958-RA|CG13958
8958	GUCUAUGAGAUGACCGUAAUGAGC	1	0	transposon	LTR	ninja-Dsim-like
8959	ACGUCGCUUUUUGAUCCUUCGAUGUCG	1	0	ncRNA		rRNA
8960	GCUUGCACCAUCACCAUCACCAU	1	2	coding gene		CG11505-RA|CG11505
8961	CUCUGAGAAAGCAGACGCGUGAGUA	1	0	transposon	LTR	Max

8962	CAGCUCGGGGAGGUAGUGACGAA	1	0	ncRNA		rRNA
8963	CUCGCAACCAUUCCUGAACUUGAUCAUC	1	0	no annotation		
8964	UCACUGUGAGGGGAUCCUCAA	1	no match			
8965	AGACAACCAGAGAUCUAGAGGGCC	1	0	no annotation		
8966	GCAUUCAAUACAGCGAGAUGGCCCAUU	1	0	transposon	LINE	Rt1a
8967	UCUACGAGAGUCUGAAUAUUGUUGA	1	0	transposon	LTR	flea
8968	GGAGAUGAAAAUGUAGUGAACGCCA	1	0	transposon	LTR	invader4
8969	AAUCCCAAACUACUAAACUACGAAA	1	1	transposon	LTR	McClintock
8970	GGCAGCGCCGCGGAGCCUCGGUUGGCC	1	no match			
8971	CCAGACUUGACCUCCUUCGACGCGUA	1	no match			
8972	GAGGUAACGAUGGAGUUUGACGGCAG	1	0	transposon	LTR	flea
8973	CAUGGCCCAAAUCGUUGCGACGGUCC	1	0	transposon	LINE	HeT-A
8974	UGAGAGGUGGCAGAGUGGUCGAAUGUA	1	no match			
8975	GACUAUGGACGAUCAAAGACCCUGU	1	0	coding gene		CG6315-RA|fl(2)d
8976	AAACAAUGGAAAAUAAUUAUGCGGUUGG	1	no match			
8977	GUGGCUUUGAGAGGCGCCGUAGUUUU	1	2	no annotation		
8978	UGACGCCUUGCGUGUUACGGUUGUU	1	0	ncRNA		rRNA
8979	UAGGGAUGAACCUUCACAUGAUGA	1	0	no annotation		
8980	CCGAAGAGAAAGAAUAUUGAACCCCC	1	1	coding gene		CG40228-RB|CG40228
8981	CCUGAAAGCAGACGAGGUAACGUACC	1	1	transposon	Low_complexity	F-element
8982	UAUUCCCCAUGAACGAGGAAUUCCCAGU	1	no match			
8983	AUGGUGACCACGGGUGACGGGGUAA	1	no match			
8984	GGUCUACUUCCCCCCAGACACUCGA	1	0	transposon	LINE	G2
8985	GCCAGGUUGGCCGAGCGGUCAAAGGCC	1	2	ncRNA		tRNA
8986	CACUACAACAAUAAAUCUAAAGGGCC	1	1	transposon	LTR	rooA
8987	AGCCUGGACGGUGUGAGGCCGGU	1	no match			
8988	CGAGUGUUAUUGUGGGCCGGUACUAUU	1	0	ncRNA		rRNA
8989	ACUCGCCGAAUCCCGGGGCCGUGGGAG	1	no match			
8990	AAUACAAACAGAAUUCCCGAAUGAGAGU	1	0	transposon	LINE	Cr1a
8991	AAACCGGGCCAGGCCAGUGCUCUUGUG	1	no match			
8992	ACAGAUGAUUUUUGGAUAGCCACA	1	1	no annotation		
8993	CAGACAUCUACUUCCCCCGCUCCU	1	0	transposon	LTR	invader1
8994	GCAGGAACUCCCGACUUUGAUGA	1	0	transposon	LTR	gypsy4
8995	UUAUACCCAGGUGAUUACCACUAU	1	0	Su(Ste)		Su(Ste)
8996	AAGAAGGUAAACGGGUGGGGGUC	1	no match			
8997	ACCAUGGUUGCAACGGGUAACGG	1	0	ncRNA		rRNA
8998	GGCGGCGGCGGCGACUCUGGACGCG	1	no match			
8999	UCCUGAACGUAGUCUCUCGAGAGCGA	1	2	transposon	LTR	invader1
9000	AAACCUAAAGAUGAUCCUGAGUC	1	1	transposon	LINE	Helena
9001	GACGAACUACCGAGCCCGAGCCUUA	1	0	transposon	LTR	Max
9002	UAAUAUGAUAUUUAUUUGAUGUGAACGA	1	0	no annotation		
9003	ACAGGCUGAUAACCCCCAAGAGCUCA	1	no match			
9004	CGGUCCCCGGGACAUGGCGUGGUGGGUA	1	no match			
9005	UCUCCCCGCGGAAAGACGGAAAGACCC	1	no match			
9006	CGUGUGAACAGUGGUUGAUCACGA	1	0	ncRNA		rRNA
9007	GUUUCGUGUGAACAGUGGUUGAUC	1	0	ncRNA		rRNA
9008	ACCCGAAACCCGGCGAUCUAGGCAU	1	no match			
9009	AUGUAGCCUUCAUUCAUGUUGGCAGUA	1	0	ncRNA		rRNA
9010	UCGGCUCGUGCGUCGAUGAAGAACG	1	2	ncRNA		rRNA
9011	UGUUUCAUCGUUAGACGGUCUCGGGC	1	no match			
9012	AACUCAAAGACAGCUAUGUACCCGA	1	0	transposon	LINE	G2
9013	GUGGAAGUCAAAAGAGUUGAGAUCGC	1	2	transposon	LTR	Quasimodo
9014	GGCCCCGAUACCAACGACAGACCCAUA	1	0	transposon	LTR	accord
9015	UGUUUCAUCGUUAGACGGCCCGGGCG	1	1	AT-chX		AT-chX-B
9016	ACAUUCAUAACCGAGCGACUGUUCCAUC	1	0	transposon	LTR	GATE
9017	GCUCAGAGCGAACGCUGGCGGCAGGC	1	no match			
9018	GCCCGAAUUGACAAAUCCCGGAACU	1	0	coding gene		CG8330-RA|tomboy40
9019	AACCAGGUGGCGCUGAACCCGCAGA	1	no match			
9020	AACCCGAAACCCGACGAAAUUGC	1	0	coding gene		CG42266-RA|CG42266
9021	CGAGCGAACAGUGGUAUGACGCUAC	1	0	ncRNA		rRNA
9022	UCCAGAAGGCAAAUGAGUGUAGAU	1	1	no annotation		
9023	CCCCUGAACGCGCCCGAUCUCGUC	1	no match			
9024	UAAACUACUACAACAACGCCGACUGG	1	0	transposon	LTR	Stalker2
9025	UAAUGCCGAGAACUGAUGACGAUCC	1	no match			
9026	AACUCUAGAUAACAUGCAGAUCGUA	1	0	ncRNA		rRNA
9027	AUCAAGGGCAACGAGUUCCUCUGC	1	0	Su(Ste)		Su(Ste)
9028	UGUCCCAAGAGAGAAAAGAGUUCAA	1	1	transposon	LTR	invader4
9029	AACACGGGAAAACUUACCAGGUCCGAAA	1	1	ncRNA		rRNA
9030	ACAAGCUUAGUGACGACAGCAAGGAU	1	0	coding gene		CG33523-RA|CG33523
9031	AAACUGAAUAAACGAAUGGAUGAC	1	no match			
9032	AUAUCCGUGAACUACAUCCCAACGACCGA	1	0	no annotation		
9033	CAGCUAGACUAUGUAUCCCCGGGUUAA	1	no match			
9034	GUGGAUCCGGGCGUGGAAUGCGAGUG	1	no match			
9035	GUGCUCGUUCUGGGUAAAUAGAGUUU	1	0	ncRNA		rRNA
9036	CGACUCAUUAUCCCGCUCAGCUGA	1	0	transposon	LINE	G2
9037	GGCGAAGGCGGCUCACUGGACUGGU	1	no match			
9038	CAGAGGUAGACGACCGGGCGAUGG	1	0	transposon	LINE	R1-element
9039	AAGAAAUACAGAGUCGUGAUUGG	1	1	transposon	LTR	HMS-Beagle2265
9040	GCUCACCAAGGCGACGAUCAGUAGCU	1	no match			
9041	AGCAAGAAGGCGCCGGUGGAAACC	1	0	coding gene		CG17764-RA|CG17764
9042	GCGACCUCACGAUCAGGACGUGGCGACC	1	no match			
9043	UCCUGAGAUACCCUGAUUACAAAAAAGC	1	0	transposon	LTR	blood
9044	AUCUCGAUCAUGUGACUCCGGGUAG	1	1	transposon	LINE	Doc2

9045	CCAUUUUGGACGAACUGCACGAGC	1	1	coding gene		CG9755-RA|pum
9046	ACUCGCCGAAUCCCCGGGGCGAGGGA	1	no match			
9047	UCUGACGAGAACUCGAGCCGCGUAGC	1	0	no annotation		
9048	AGCGGCGAGCGAAGUGGGACCACGCCC	1	no match			
9049	ACGGGAAACCUCACCCGGCCCGGACACG	1	no match			
9050	UUGGGAACGCCGCGUGUUGUUGGCCU	1	1	ncRNA		5SrRNA
9051	AAAAAGUGAAGAAGACGUCGACG	1	2	no annotation		
9052	UAUUUAGAGGAAGUAAAAGUCGUAAC	1	0	ncRNA		rRNA
9053	AAGCUACCAUCUGUGGGAUUAUGACUG	1	no match			
9054	GGCUUUGAGAGGCGCCGUAGUUACACCC	1	2	no annotation		
9055	CUAUCUAGGACAAUGAUGCGACCCGUC	1	0	transposon	LTR	ninja-Dsim-like
9056	GGAGGAAAAGAAACCAACCGGGAUUGC	1	no match			
9057	ACCCCACGGAACACCCGAGCAAUGCCAC	1	1	transposon	LINE	G2
9058	UGCGGGAUGAACCGAACGUAGAGUU	1	no match			
9059	AGUGACGAAAAAUAACAAUACAGGAC	1	0	ncRNA		rRNA
9060	CCCGGGCGUCGCAUCCCCCCCCGCUAGC	1	no match			
9061	UACCAACCCAGAAUAUGACCCAGCAUU	1	0	transposon	LTR	McClintock
9062	AUCUCGUCGAGAAUGAACUUCG	1	no match			
9063	CACGAAACUGGAACGGGAUUAAGAU	1	0	no annotation		
9064	AACUUAACUAUGUUUCCCCGGUAGCGC	1	1	transposon	LINE	Doc3
9065	AAAGUGGUACGCGAGCUGGGUU	1	no match			
9066	GAUGUCUAUGAGAUGACCGUAAU	1	0	transposon	LTR	ninja-Dsim-like
9067	CAAGAUAUGACCAAAGCAAGACAC	1	no match			
9068	UGAGAUAUGCUAGCAGGCUGGAGA	1	1	transposon	LINE	G2
9069	GACAUAAACAUCCGCAGCCGCAUUC	1	0	transposon	LTR	Quasimodo
9070	AAACGAGUGUUAUUGUGGCCGGUA	1	no match			
9071	GCCUGCUCCAAGUGUUUCUCGCGCC	1	0	coding gene		CG4452-RA|CG4452
9072	CGGCCAAGCGUUCAUAGCGACGUCGC	1	0	ncRNA		rRNA
9073	AAACGACAAUGGAUGUGAUGCC	1	0	ncRNA		rRNA
9074	CGGACCUAAAGACGGUUCCAAGCCCA	1	1	transposon	LINE	G6
9075	GCCCCGGGUAGCUCAGUCGGUAGAGCAUUG	1	2	ncRNA		tRNA
9076	AGUGGAGCCGUACCUGUUGGUUUGUC	1	0	ncRNA		rRNA
9077	CACAGAGUACGGGGCUUACUAAAA	1	1	transposon	LINE	baggins
9078	CUCGCAAGAAAACUAAUCCGAGUA	1	0	transposon	LINE	Doc
9079	UUUGAUAAUAACCCCAGGAAUUAAAA	1	0	transposon	LTR	opus
9080	AUGCUCUCGAAUGUAUGCCCCGAUAU	1	no match			
9081	GCAACUAGACGACGCUGCUAUACC	1	no match			
9082	GGUCUACGAGCGGGCUUGGACGCGU	1	no match			
9083	AAAGCCGAAAUCAUUGCCAAGGUCGC	1	2	coding gene		CG2113-RA|CG2113
9084	GACAGUCCAAACUCGAACCCCGUAAC	1	0	transposon	LTR	GATE
9085	AAACCAAAAGCUUUUGGGCUCCGGGGGA	1	no match			
9086	AGCGCGAGAAGGAGCUGGGCGAGAA	1	0	coding gene		CG5119-RA|pAbp
9087	AACAGAGAGAAGACUCACCCGACGAGA	1	0	no annotation		
9088	UCCUUAGUUCAGUCCUGAGUGCGAGCUCC	1	no match			
9089	UUCAAGGUUACCCCAGGAUGCUGUGU	1	1	repeat		trf
9090	GCCCAGCUUCCCGGACAUCUUCUUUC	1	1	Su(Ste)		Su(Ste)
9091	ACCCUUUCAUGCUUGGGAUUGUG	1	2	ncRNA		rRNA
9092	GCGACGAAAACGAUCUUUGCGAC	1	0	transposon	LTR	diver2
9093	UGCGAUGAACCUGUCACCUAGUGUG	1	0	transposon	LINE	Doc
9094	GAGCAGAGAACCCUCUAAUUCGGCC	1	0	transposon	LINE	baggins
9095	UUGGUGGUGCAUGGCCGCUCCUAGUU	1	no match			
9096	CAAGCUAGACUAUGUAUCCCCGA	1	1	transposon	LINE	Doc
9097	GGAUCGAGGCGACGCCGUACCGUUAUG	1	0	coding gene		CG31178-RA|CG31178
9098	ACUCGCCGAAUCCCGGGCCGAGGGAGU	1	no match			
9099	CCCGUUAUAUCUAUGUGGAUGGUGUGC	1	0	transposon	LTR	micropia
9100	CGGGAAAUGAAAAAAUAUUGGCGA	1	0	transposon	DNA	1360
9101	UGCUCUCGAAUGUAUGCCCGAUACUAC	1	no match			
9102	ACGAGUAAUAUGAGCGCCCGGAGU	1	1	transposon	LTR	opus
9103	UGGGCUCCGGGGGAAGUAUGGUCGCUA	1	2	ncRNA		rRNA
9104	ACGUAAACAAUAAGUGACCACCAUGC	1	0	transposon	LTR	17.6
9105	GGAUCCUCAACUGUGGGCUCAA	1	no match			
9106	CCGCCCUAGUUCUAACCAUAAACG	1	0	ncRNA		rRNA
9107	AUGGAUGACUCGAUGACUCCGAUCUGCA	1	0	coding gene		CG7045-RA|CG7045
9108	GCGAUGCGGUGGUGCAGUCCGACAUGA	1	no match			
9109	AAGACCGUAAACGAAAAGCUAAGAA	1	0	transposon	LTR	Idefix
9110	AAAACCUGAAGAUGAUCCUGAGGC	1	1	transposon	LINE	Helena
9111	ACCCGGAGAAGAAGCCCCGGCUAACUUCGU	1	no match			
9112	AUGCCAAUCACGCCGUUUGAACCCU	1	0	transposon	Simple_repeat	I
9113	GUUCGAUCCCGUCAACCUCCACCA	1	no match			
9114	CAAACGUAAUGUUACGGUGCCCAAAUUA	1	0	ncRNA		rRNA
9115	CGCAUUGAUUACGUCCCUGCCCUUUG	1	0	ncRNA		rRNA
9116	CAGACCAUAUCUUUAGCUGUGCGCU	1	no match			
9117	GUCACCUACAGGCCCCGAUGCGAGCAC	1	1	transposon	LINE	BS3
9118	AGAGAACAUACGAAACACCACCCA	1	0	transposon	LTR	flea
9119	GAGCAACUGUGAGGGGGAUCCUCAA	1	no match			
9120	ACAUACUUAGGUAUUCAUCUGGACA	1	0	transposon	LINE	Doc3
9121	GAAGCCGUCGAGAGAUAUCGGA	1	0	ncRNA		rRNA
9122	CGUGGGUGGUGGUGCAUGGCCGUUC	1	2	ncRNA		rRNA
9123	UCUCAUCGUCGUAGAACAAGCCCGAUC	1	2	Su(Ste)		Su(Ste)
9124	AACAACGUUCGCGUGAUGACCCCGGA	1	0	transposon	LINE	G6
9125	UGAGUUGAAAAAUUCCGAGAAUUGG	1	0	transposon	LINE	Doc2
9126	ACUAAUACGAUUGCGAGUUAAGCACC	1	0	transposon	LINE	Doc3
9127	GGCACAAUUACCAGAAGAUGUUGUCU	1	2	transposon	LTR	roo

9128	UCCUGAGUUUCUUCAACCUGUUGCAGGCU	1	0	transposon	Simple_repeat	I
9129	UGAUCAGUACCCAUGAGUGAUCCACU	1	0	transposon	LTR	Quasimodo
9130	GGCUGACAUGAAGAUCACCUCUGCA	1	1	coding gene		CG5016-RA|Mst57Db
9131	AGGACGUUCACACGAGGGUUAC	1	0	transposon	LTR	opus
9132	ACCCUGAACGCGCCCGAUCUAGUC	1	no match			
9133	AUGAUUCCCGGCUGGUUGGCCCUCGU	1	0	coding gene		CG33233-RC|CG33233
9134	CGACAGAGACGGCGCAUUUUCCA	1	1	transposon	LTR	McClintock
9135	UUGUACACACCGCCCGUCGCUACUACCG	1	0	ncRNA		rRNA
9136	ACCCUUGAUAAACGGUUGACUUUCAG	1	0	transposon	LINE	BS
9137	CACGAUGAUCGAUGACGAACUACG	1	0	transposon	LTR	ninja-Dsim-like
9138	CUGAAGUGGAGAAGGGUUUCGUGUGAA	1	0	ncRNA		rRNA
9139	CACACAGUCAGGGGUUCCAUCGG	1	0	transposon	LINE	Cr1a
9140	AUGCUGACUAAGAGGGCUGCACAGU	1	0	transposon	LTR	HMS-Beagle
9141	UUCAAUCCCCGGCAUCUCCACCA	1	no match			
9142	UCUGGUUAAUUCCGAUAACGAACGAGA	1	0	ncRNA		rRNA
9143	CUGGACGGUCAGCAACAUCACAGC	1	0	coding gene		CG4799-RA|Pen
9144	AGAAAGCACCGUUUCCGUCCGAUC	1	no match			
9145	AAGCUUCAUCCAAGAGAACUGUGCCCGA	1	1	transposon	LINE	R1-element
9146	CAUCGCUGGAGGACGCCCGGGCUG	1	0	transposon	LINE	R1-element
9147	CUGAAGUGGAGAAGGGUUUCGUGUGA	1	0	ncRNA		rRNA
9148	AAGGUGGAAAUAAAGAAAAGGAUCCUCUAA	1	no match			
9149	AUCUCGAGGAAUACGGCCUUACCGCAG	1	0	transposon	LTR	ninja-Dsim-like
9150	UGACGAUGGGGUGAGGCUUAAGCAA	1	0	transposon	LTR	Idefix
9151	UAUAGGGUCUGACGCCUGCCCGGUG	1	no match			
9152	CCCAGCUCGGGGAGGUAGUGACGUAA	1	1	ncRNA		rRNA
9153	UUCGCUAUCAUACCCGAACACCAGUUA	1	1	transposon	LINE	G4
9154	GGCUUUGAGAGGCGCCGUAGUUACC	1	0	no annotation		
9155	ACCGGGACCAGGCCAGUGCUCUUGUGUA	1	no match			
9156	UACAUCGAGAAUCUCACGCCCACAGGCA	1	0	transposon	LINE	Doc3
9157	UAUCAAGAAAUCCUUUAUCACCCGAAC	1	0	transposon	LTR	micropia
9158	CUGUCUCAGUCGCUUCCCGGGAAA	1	1	ncRNA		rRNA
9159	CUGUAACUCGGGUCAACUGUAACUCGG	1	no match			
9160	GGCUCAACUGUGAGGGGAUCCUCUAA	1	no match			
9161	AACGCGACCGAUCCCGGAGAAGC	1	no match			
9162	UCCGAGGAACCAAUUGAUCCAGCUG	1	0	Su(Ste)		Su(Ste)
9163	ACUACGCCGAAUCCCGGGGCCGAGGGAG	1	no match			
9164	CGGAGCGAUGUUCGGGCCCAGCUU	1	0	repeat		trf
9165	ACAUAGGGACGGUCGGGGGCAUC	1	no match			
9166	ACGAGUAAUAUGAGCGCCCAGAC	1	1	transposon	LTR	opus
9167	AGGUUCGAAGGCGAUCAGAUACCG	1	0	ncRNA		rRNA
9168	UAAUGCCCCGAAGGUUGCCCCUCUG	1	no match			
9169	UGUCUCAUCGUUAGACGGCUCGGGCG	1	1	AT-chX		AT-chX-B
9170	CACAGAGUACGGGGCUUACUAAAGUCUA	1	1	transposon	LINE	baggins
9171	CUGUGAGCGGAUCCUCAA	1	2	coding gene		CG10431-RA|CG10431
9172	AUACAUGCCGACGGGCGCUGACCC	1	no match			
9173	UUACUGAUAACGAUAUCUGCUGUGG	1	1	no annotation		
9174	AACUGAAUGUGGAGACGUCGGC	1	no match			
9175	GCUCUCGAGAGACUACGUUCGGACAGUA	1	no match			
9176	CAAGAUAUGACCAAAGCAAAGCAACG	1	0	Su(Ste)		Su(Ste)
9177	CGACGUGGCGACCCGCUGAAUU	1	no match			
9178	GGAUCCUCAACUGUGAGGGAUCCUCUAA	1	no match			
9179	UCCGAAGACAAGUAUACCGUGGC	1	0	transposon	LTR	micropia
9180	CAUCAUAUUCAUCCUGUUGGUCU	1	no match			
9181	GAAGAUAAAAACUGGAAAGCCCGGGCGC	1	1	transposon	LTR	Dm88
9182	CUUGGUUAGGUCGAUUCUUUUGAAU	1	no match			
9183	GAGAGGUGUAGAAUAAGUGGGAGGCC	1	no match			
9184	CCCCGCGGAAAGACGAAAAGACCCCA	1	no match			
9185	ACAUUCAUAACCGAGCGACUGUUCU	1	1	transposon	LTR	GATE
9186	UGUCCCUACCUACUAUCCAGCGAA	1	2	ncRNA		rRNA
9187	CGAAGCCGAGCAGGAACCGCGGGA	1	0	coding gene		CG8531-RA|CG8531
9188	UUCACAGUAAAAUUUUGUAGCCGA	1	0	no annotation		
9189	CAUAUCCGAGGCCCUGUAAUUGGA	1	0	ncRNA		rRNA
9190	UCCGUUCAAAAAUGUUCAUGCAGGAGA	1	0	transposon	LTR	Rt1c
9191	ACCGAAGAAAACGAAGCCUCCGCCAA	1	1	transposon	Low_complexity	F-element
9192	AGACUGGCGAUGCGGGAUGAACCUAAC	1	no match			
9193	UCGGCCCGAACCCUUCAGGUGUGUUG	1	1	transposon	LINE	baggins
9194	UGCGCUCGAAUGUAUGCCCGAUAU	1	1	transposon	LINE	R1-element
9195	UGCCCUGAAUGUAUGCCCGAUA	1	no match			
9196	GGAUCCUCACUGUGAGGGGAUCCUCAA	1	no match			
9197	UUUGCCAAUCUCGAGCCGGCCAAGA	1	0	transposon	LINE	G6
9198	ACCACCCUGAACGCGCCCUAUCUCGUC	1	no match			
9199	UGAGCUAAUAGAGAACAGGAAGGAUC	1	0	transposon	LTR	rooA
9200	AACAUCGUCCCGGAUAACCAUGGUGCAU	1	0	coding gene		CG31988-RA|CG31988
9201	GUCCUGAGUAUUCUCCUGUUGUCC	1	0	transposon	LTR	Dm88
9202	UUCCGAGGAACCAAUUGAUCCAGCU	1	0	Su(Ste)		Su(Ste)
9203	CGGCCCGAACCCUUCUGGUGUGU	1	0	transposon	LINE	baggins
9204	ACCGAUGGGCCUGGAGUACAUCAUC	1	2	coding gene		CG40820-RA|CG40820
9205	GAUUCCGGAGAGGGAGCCUGAAA	1	1	ncRNA		rRNA
9206	UAGAGAAAAACAUGGAACCAUCG	1	0	transposon	LINE	Doc
9207	GUCCCUGUGGAGGUUGCUGUAGUCC	1	0	coding gene		CG31740-RA|CG31740
9208	CCAGGCCCGUAUAACGUUAAUGAUU	1	0	ncRNA		rRNA
9209	AACAGGCUGAUAACCCCAAGAGCUCA	1	no match			
9210	UCCCCCUGAGAGAGGGGCCCGCGCCCU	1	no match			

9211	CAACCUGAGAGUUUGAUCCUGGCUUC	1	no match			
9212	UGCAGCCAAUAAGGACUUACGUUGUG	1	0	transposon	LTR	flea
9213	AACUUGAGACCCAACAACGCAACCGAA	1	no match			
9214	UAAAGGAUUACGUGUCUAAAGGAUA	1	0	transposon	LTR	ninja-Dsim-like
9215	UAUGAAUCGAGUUAAAACUGCAGG	1	1	transposon	LTR	diver2
9216	GAACUUGAGACCGCCCCUGGACGAC	1	0	Su(Ste)		Su(Ste)
9217	GCAUUCGAGACCCAACGGCACGUGAAAA	1	2	transposon	LINE	INE-1
9218	UUAAUGCCGGAAGCGAUCCCACU	1	0	AT-chX		AT-chX-B
9219	UCUUAAACGAGUGUUAUUGUGGGCCGGU	1	0	ncRNA		rRNA
9220	UGAGAACGAUGACGAUCCCACGG	1	0	transposon	LTR	ninja-Dsim-like
9221	AAGGAUUGGAUGGUCAUUAGAUGG	1	2	transposon	LTR	frogger
9222	CUGCCACACGUAACCCCGUUUUUGU	1	no match			
9223	UCCGAACGUACCACGGUGCCCCAGU	1	0	transposon	Low_complexity	F-element
9224	UGUUUCAUCGUUAGACGCCUCGGGC	1	1	AT-chX		AT-chX-B
9225	GGCACAUCUCGACACCCGUGGCACU	1	0	coding gene		CG13340-RA|CG13340
9226	ACCACCCUGAACGCGCGCUAC	1	no match			
9227	UCAGCUCAUCUGCGCCGGGAGAUCGA	1	0	transposon	LTR	gypsy12
9228	CAAGGGAACGUGGGCUGGGAUUAG	1	2	ncRNA		rRNA
9229	CCCGAAUUAAUCGCGAGGCUGCAAGA	1	1	transposon	LTR	mdg3
9230	ACCACCCUGACGCGCCCGAUCUCG	1	no match			
9231	CACCAGGGGUUGCGGUUGGUCGACC	1	0	transposon	LTR	opus
9232	ACAUUCGAGAAGCGUUUGAUCGAGU	1	0	transposon	LINE	X
9233	ACCAGAGAUCUAGAGGGCCCGA	1	1	no annotation		
9234	CGAGACUUUGAACUAAAAACAGUUACC	1	0	transposon	LTR	Rt1c
9235	AACUCGUAAUCGAUGGCCAUGCGUUCC	1	0	coding gene		CG8561-RA|AIs
9236	CACGAAACUGGAACGGGAUUAAGAUC	1	0	no annotation		
9237	ACACGACAACAUUCCGGGAAAUGAGC	1	1	transposon	LINE	baggins
9238	GCUAACACAAAAGUCGACUUCUA	1	0	transposon	LINE	Doc2
9239	CGCCGCAGGGCAGUUGGGGUAAGGAC	1	0	coding gene		CG32688-RA|Hk
9240	ACGGGGAAACUCACCAGGUCCAGA	1	no match			
9241	GCCCCAAUGAGAACGAUGACGAUCCC	1	0	transposon	LTR	ninja-Dsim-like
9242	ACCCCUGAACGCGCCCCGAUCUCGUC	1	no match			
9243	UUAUCGACUCGUGUACCUGGUUAA	1	0	transposon	LTR	micropia
9244	CAGUACAGUGGCCAGCACGAAAACCGA	1	0	transposon	LINE	BS3
9245	ACCGACGUUUUCAAAUUAUGUAGGCAA	1	0	transposon	LINE	RR49783
9246	UAAACCUGAAGAGAGCCCUGAAUCG	1	0	transposon	LINE	Rt1b
9247	ACGAGUAAUAUGAGCUCCCAGAGU	1	1	transposon	LTR	opus
9248	GUCGUCCGAGGCGAAGACUGCGAGGGGU	1	0	no annotation		
9249	AACCUCAUCAAGUUACCAUUCCG	1	0	transposon	LTR	Rt1c
9250	GCCGAACUUGAGACCGCCCCUGGACGACC	1	0	Su(Ste)		Su(Ste)
9251	AUACCUUCGGGACCCGUCUUGAAACAC	1	0	ncRNA		rRNA
9252	UCGGGCGGCCCGGGUUCGUUUCCC	1	2	ncRNA		tRNA
9253	ACAGUGCCAAGGCUAAAGAUGAUUCGU	1	1	coding gene		CG2658-RA|CG2658
9254	GAGGAGGAGGACUAGCUGGCCGAGUA	1	0	Su(Ste)		Su(Ste)
9255	GGCUUUGAGAGGCGCCGUAGUUACCUACCC	1	2	no annotation		
9256	CCGAACUUGAGACCCCAACCGCUCC	1	1	no annotation		
9257	ACGAGCAACGUCAACAGCCACCACAGC	1	0	coding gene		CG18811-RA|CG18811
9258	UAUUACAGGAGAUAUUUGGACACGAU	1	0	Su(Ste)		Su(Ste)
9259	UGAUAGUUCAAAGACCCCCAUAUG	1	no match			
9260	CAACGAGCGCAACCCUCGCCAUUAGUUG	1	no match			
9261	GUCGAGGGAUAUCGGAAGAGUUU	1	1	ncRNA		rRNA
9262	CAACACCAAACAACAAAGAGAACAACGC	1	0	transposon	LINE	Cr1a
9263	GGUUAAUUCCGAUAACAAACGAGACUCU	1	2	ncRNA		rRNA
9264	CUCCCACGGAACACCAGAGCAAUC	1	0	transposon	LINE	G
9265	UUAUUAACGAAACCUGAGACCGACA	1	1	transposon	LTR	297
9266	CCCGAAAAAAUCCCCCCGGCUACGACC	1	no match			
9267	AACCCUUGUCCUUAGUUACCAGCACGU	1	no match			
9268	AGUAUACCUAUACAUUACCGCUAAA	1	1	ncRNA		rRNA
9269	UGCACGAACAAAGCCAGUAAUCAU	1	0	transposon	LTR	flea
9270	AAAACCUGAAGAUGAUCCCUAGUCG	1	2	transposon	LINE	Helena
9271	UCUAGAGACAAGGACGCAGCGCAAU	1	1	transposon	LINE	INE-1
9272	AAAAUUACAAAGACGACGGCCCAU	1	1	transposon	LTR	springer
9273	UCUCAUCGUCGUAGAACUAGCCCAA	1	1	Su(Ste)		Su(Ste)
9274	CACACAACAAAACUAUAACCAGCUGC	1	no match			
9275	CUGACAAGGAAUCAUUACUCAAUGA	1	0	transposon	LTR	springer
9276	UAGUACAAUAGAUCCGACUGGGCCAGU	1	0	transposon	LINE	baggins
9277	AGGAACGAAAGUGGACAGCGAGCGAAA	1	0	transposon	LTR	diver2
9278	AACGUAAAUAAAAAAGCUGGGUCG	1	1	transposon	Low_complexity	F-element
9279	UGCCCGACAAACUCAAGGAGGUGGCAG	1	0	coding gene		CG5215-RA|Zn72D
9280	UCUGGGAUGACUUGUGGAUAGGGGU	1	no match			
9281	GGCGGUGCGCCGCGACCGGCUCCGGGAC	1	no match			
9282	UCGAGAUGACAACCUUGGACCUGAAU	1	2	transposon	LTR	micropia
9283	UUCCGAAUCAAAUGCUACGUAACC	1	0	transposon	LTR	diver2
9284	UACUGCGAUGGGGGGACGGAGA	1	no match			
9285	CCUCAGCGAUGUGUGGGCCAUGUU	1	2	Su(Ste)		Su(Ste)
9286	CCGAGAUUCCCUUAGUAGUGGCGAGC	1	no match			
9287	CAUUCAUAACCGAGCGACUGUUCA	1	0	transposon	LTR	GATE
9288	UGAGAUGUACAAUGCUGCUGAGU	1	0	transposon	LINE	G2
9289	UCGAGAAGAAGAUGAAGCGCAAC	1	0	transposon	LTR	ninja-Dsim-like
9290	UCGGGAUAAGGAUUGGCUCUGA	1	0	ncRNA		rRNA
9291	AGAGCGUGUGGAGGACCGAACC	1	no match			
9292	GAAUGAAACAAACAAUCUUCUGGGA	1	2	transposon	LTR	diver2
9293	CAAGCUAGCCUAUGUAUCCCCGGGUAA	1	1	transposon	LINE	Doc

9294	GGCCAACUGUGAGGGGAUCCUCAA	1	no match			
9295	CCAACACAUUUUGGAGACCCACU	1	0	transposon	LTR	Stalker4
9296	CCCGGGAUCAAUGCAGUGCUGCCUGACC	1	0	coding gene		CG3124-RA|CG3124
9297	CUGUUUUCGAGGGUACCUAUGUAGGU	1	0	transposon	LINE	Rt1a
9298	GCGAGUAAUAUGAGCGCCCAGAGUGAUC	1	2	transposon	LTR	opus
9299	GUGGGCGGCGGGACCGGGGGGGU	1	no match			
9300	CAUGAAAUGAACCCGAAAGCACUGA	1	0	transposon	LINE	Doc3
9301	GGACAAAAACAACAUCCCGGGAAAUGA	1	no match			
9302	UCAAGUAUAACGACCCAUUUCUGU	1	0	repeat		trf
9303	AUUGCCCUAGCUGUAACAACGUCUAC	1	0	coding gene		CG13591-RA|Ssl
9304	CCAGACGUAAUACUCCACUACCCGAU	1	2	transposon	LTR	HMS-Beagle
9305	AACAACCGAUAUGCUAUCUUGGCUGAU	1	no match			
9306	UGUUUCAUCGUUAGACGUCUCGGGC	1	1	AT-chX		AT-chX-B
9307	CUCCCCCCGCCCCACCCCGCGCGCGCCGC	1	no match			
9308	CCGAGUAUGAUUCUGACUUCGUAAA	1	0	transposon	LTR	springer
9309	AGUAUCUAGGACUACCAUGGUUGCAAC	1	0	ncRNA		rRNA
9310	CGGCCUGAUGACUGUUGAUGAUC	1	1	transposon	LINE	Circe
9311	AACGAGAAUAUACACAGGGGCUUAC	1	0	transposon	LINE	Doc3
9312	AAACUAAAUAAACGAAAUGGAUGACA	1	1	transposon	LTR	invader4
9313	UAGGCCAUGGAAGUUUGAGGCAAU	1	no match			
9314	UACCCGCGAGAAUUGCCAGUUUCGA	1	0	no annotation		
9315	GAGCGAAAACUUAAAUACCGAGAAA	1	0	transposon	LTR	Dm88
9316	AUAGAAGUAACCUUCGACUUGGACGCC	1	0	coding gene		CG31359-RA|Hsp70Bb
9317	AGUCGAGAGUGGACGGGAGCGGCGGG	1	no match			
9318	GCUUGAUAGUGCAGCACUAAGUGG	1	0	ncRNA		rRNA
9319	ACUUCGGGAUAAGGAUUGGCUCUGAAGU	1	1	ncRNA		rRNA
9320	GUCCUGAGUAUUCUCCUGUUGUCCA	1	0	transposon	LTR	Dm88
9321	UACUUCGCUACUACCCAUUUGGAA	1	1	no annotation		
9322	AUAGAAGUUAAUUAUCUACCCACAGA	1	1	transposon	LTR	DM88
9323	AAGGAGAGCAAUUUUGGCCGUCACU	1	0	transposon	DNA	1360
9324	UUUCUUUGAAGAUGGCCCACAUAA	1	0	repeat		trf
9325	AGUUCAUUUCUGACAGGUGGCACC	1	no match			
9326	CAGCAAGAGACUCGAGUAGGGCGGGA	1	no match			
9327	UCUCAUCGUCGUAGAACUAGCCGA	1	2	Su(Ste)		Su(Ste)
9328	UCGAAAAUAAGAAUCAAUGGCUGUGU	1	0	no annotation		
9329	ACUACAUAUGGUUGAGGGUUGUA	1	0	ncRNA		rRNA
9330	ACUCCCCGCGAACCCGCAAAACUGA	1	2	transposon	LINE	TART-A
9331	UGCUCUCGAAGGUAUGCCCGAUAU	1	1	transposon	LINE	R1-element
9332	GUUCAUUUGAAACUAAAGCUAAGAUA	1	0	transposon	LINE	HMS-Beagle2
9333	AUGACAGUUCACGAUUGAUUCCC	1	0	transposon	LTR	opus
9334	CACGCAGUUACCGACCGUUGUGUCUGU	1	0	transposon	LINE	Rt1a
9335	AGUGUGUAAGACAGAUUGAUAGCUCU	1	0	ncRNA		rRNA
9336	AGCUCCACGGAGACGCUGAUGGACCUCGCU	1	0	coding gene		CG12679-RA|CG12679
9337	GCCAACGACCAUACCACGCUGAAUAC	1	0	ncRNA		5SrRNA
9338	AUAGCGUACCGGAGUGGACUGCCGUU	1	0	ncRNA		7SLRNA-RA
9339	GGAUCCUCAACUGUGAGGGGGAUCUCAA	1	no match			
9340	CUGGAUCGAUUGGUUCCUCGGGAU	1	0	repeat		trf
9341	AGAGUUCCCGUGGUAGGCAGUAGU	1	0	ncRNA		7SLRNA-RA
9342	UGCCAAUGGAGAGCGACUGCAGC	1	0	transposon	LINE	R1-element
9343	UUUUAUGAAAUUGAGGUUCUAGAA	1	0	no annotation		
9344	UAGGAGUUCACCGCCACGACGCCCG	1	0	transposon	LTR	GATE
9345	CCGGAGCGAGGAGGAGGACUAGC	1	0	Su(Ste)		Su(Ste)
9346	CACCAGGAGUGUAGCCUGCGGCUUAAU	1	1	ncRNA		rRNA
9347	GGAAAAAGGCGAUGCCCGUGAGGACC	1	0	no annotation		
9348	UCAAGGAUAACAGGACCAUCCAAGA	1	1	no annotation		
9349	UGUUUCAUCGAUAGACGGCUCGGG	1	1	AT-chX		AT-chX-B
9350	UCCUGGGAAACGGGGCGCGGCCG	1	no match			
9351	CUGCGCUGCGGAUCGUAAUCUCAUCGC	1	no match			
9352	CAAACAAGCAAAGUGAACACGUCGC	1	0	ncRNA		alphagamma-element
9353	UGGGCCCGAACCCUUCUGGUGUGUUG	1	0	transposon	LINE	baggins
9354	CGUUCAAUAAUGACAGAGGUACA	1	0	transposon	LINE	Rt1a
9355	GACUCCAUGGCCAAGUUGGUUAAGGC	1	no match			
9356	GAACGUGAAACUACAAACGGAUUGC	1	1	no annotation		
9357	GAUUAACGAGAUUCCUACUGUCCCUAU	1	0	ncRNA		rRNA
9358	GAACCCCAUUCGUAAGGCGCCCGAU	1	no match			
9359	GUUACCUGGAACUAAGAUUGGCU	1	0	transposon	LINE	INE-1
9360	ACCCUUGAUAAACGGUUGACUUUUC	1	2	transposon	LINE	Rt1b
9361	GGCUUCAUCCAAGCGAACUGUGCCCGA	1	1	transposon	LINE	R1-element
9362	GAACAUAUCAGGGGCCUUCGCAGA	1	1	transposon	LTR	HMS-Beagle
9363	UAAACCAAAGGCCGUGAGUUUCCCCU	1	no match			
9364	GCACACAGAGGAGAGCAGAUGGAGAGCA	1	no match			
9365	GGUCACCUGUGCCGCCGGUAUCCGUUCAA	1	1	coding gene		CG11579-RA|arm
9366	ACAGCCAGGCCGAGGCUUUGAAAGAUC	1	0	coding gene		CG32437-RA|CG32437
9367	GUUCGAUUCCGGAGAGGGAGCCUGAAA	1	1	ncRNA		rRNA
9368	GAGUACUUCAGCCAGACACUGGACG	1	0	Su(Ste)		Su(Ste)
9369	UGAGCACAAAACGGUGGACUACCACA	1	0	transposon	LINE	Quasimodo
9370	GGAGAAAAUAAAAUCCGGGAAUGGAA	1	0	transposon	LTR	rooA
9371	UGGUGUCACCAGCCCAAUCCGCGUC	1	0	transposon	LTR	DM88
9372	AAGUAAACAAAGAGAGCGCGAC	1	0	transposon	LTR	297
9373	AACGUAAACAAAAAAGCUGGGUCGA	1	0	transposon	Low_complexity	F-element
9374	CUGGCAAAGAAAAAUUAAGAGUC	1	0	transposon	Unknown	RR50960
9375	GGCACAAUACUUGUUCGAGCGA	1	0	ncRNA		rRNA
9376	ACCUCCAUGCCGAUCAUCCUGUUGA	1	0	transposon	LTR	ninja-Dsim-like

9377	UCGAGUAGAAUGGUGAACGAGCAAG	1	0	transposon	DNA	transib4
9378	GGCGGAGCUGAGAAGACGGUCGA	1	no match			
9379	GGUAGUAGAACACGUUUAUUCUGGA	1	no match			
9380	UGUGUUCGAGUCCCACGCGUAGUGCCA	1	no match			
9381	GAGAACGCACCCGCUUGACCUUACUCA	1	0	transposon	LINE	G2
9382	UCCGAGGAACCAAUUGAUCCAGCUGCG	1	1	Su(Ste)		Su(Ste)
9383	CCAAGAUCAUCAACCCGCGUUAU	1	0	transposon	LINE	Cr1a
9384	CAAGAUAUGACCAAAGUAAAUACAC	1	0	transposon	DNA	1360
9385	AUGACCCACAUUGUGCGCGAGGC	1	0	coding gene		CG4863-RA|RpL3
9386	UAAACCGUCAUCCAGCGCAAGUUGCC	1	no match			
9387	UGCUCUCGAAUGUAUGUCCCGAUAU	1	no match			
9388	ACCAGAGGCAGAACCGAAACCAGAAUGA	1	0	no annotation		
9389	GCCUGAGGUGUGGGUGGCCCGCAUUCGU	1	no match			
9390	UCGAAACGAAAACAUACACAGAGACC	1	0	transposon	LINE	Doc
9391	AGGGACAGUCGGGGGCAUUAGUACU	1	no match			
9392	ACGCGCGGCAGGACGAAAAGACCCC	1	no match			
9393	UUCCUAGCCAUUAUACGGAUAUA	1	0	repeat		LSU-rRNA_Dme
9394	ACCGAAUUAUUCGCGAGGCUGCAAGA	1	1	transposon	LTR	mdg3
9395	GGCCAUACCACCCUGAACGCGCCCGUA	1	no match			
9396	AGGGGUCUGGAGGCGAACCUGGGGAA	1	no match			
9397	AGAAGCUAUAUUCAAGAAGAUGUAA	1	1	transposon	LTR	flea
9398	AUACCACACCAUCCCCGAACCGCAA	1	0	coding gene		CG7450-RA|CrebA
9399	GCCGAAGAUGUCCGGGGCUCAAGUUCA	1	no match			
9400	GAAGAAGAGAUGCCCAGUGAGUUUA	1	1	transposon	LTR	ninja-Dsim-like
9401	UCGACGAGAACGAGAUCAGCGCGA	1	1	transposon	LTR	frogger
9402	AUACGAUGUCAUCCCUGCCAAACCCAUC	1	no match			
9403	CAGGCGUUCGACAGAGUGUGGCAUC	1	0	transposon	LINE	G2
9404	ACUGCCGACCGAUCGCUUCGCUGG	1	0	coding gene		CG5525-RA|CG5525
9405	AACGAGUGUUAUUGUGGGUCGGUA	1	1	ncRNA		rRNA
9406	ACGGGAAAACUUACCAGGUCCGAACAU	1	0	ncRNA		rRNA
9407	ACGAGCGCACGUGUUAGGACCCGAAG	1	no match			
9408	AUUGACGGAAGGGCACCACCAGGAGU	1	0	ncRNA		rRNA
9409	GUCAGGAUGGCCGAGUGGUCUAAG	1	0	ncRNA		tRNA
9410	GACCAUAUACGAAGACGCGGCGC	1	1	transposon	LINE	Rt1b
9411	AAAACCUGAAGAUGAUCCUGUAGUCGC	1	no match			
9412	UGACCGGGAGGAGGUGGUACUGUGGA	1	0	coding gene		CG10590-RA|CG10590
9413	GCCCAAAGGAUACUCGAAGACACC	1	0	transposon	LTR	ninja-Dsim-like
9414	GUUCCUAAUCCCGUGAUUGUUGUUGUCC	1	no match			
9415	UGAGGACUUGGGCGAUUUAAGUGG	1	0	Su(Ste)		Su(Ste)
9416	ACAUUCAUAACCGAGCGACUAUUC	1	1	transposon	LTR	GATE
9417	ACUCAUAAGAGACUGCCGGGGUCAACU	1	no match			
9418	CUCCCACGGAACACCAGAGCAAU	1	0	transposon	LINE	G6
9419	GACCGAACAAACGAAAUGGAUGAGA	1	no match			
9420	CAGGUACGGACCAAUGGCACAAUA	1	0	repeat		LSU-rRNA_Dme
9421	GGCGAAGGCGACAUACUGGCUCAUUUA	1	no match			
9422	GCCUGAACGCCUCUAAGGUCGUAUCCGU	1	0	ncRNA		rRNA
9423	UUUCCGCAUGUGAAUGUUGGCCCCAGA	1	0	no annotation		
9424	ACGCCUCCUCCCCUCCUCCCGCCU	1	no match			
9425	ACGAUUGGGACGACAUAAAACGAGU	1	0	transposon	LTR	opus
9426	AACCCUGAACGCUCCCGAUCUCGUC	1	no match			
9427	UACAUCGAAAAGCGAGUUGACGUGUGC	1	0	transposon	LTR	Max
9428	UCAGUGCCUGACCAGGUGUUCCAGAU	1	0	coding gene		CG9611-RA|CG9611
9429	UGCUCUCGAAUGUAUGCCCGAUUAU	1	no match			
9430	AAGUAAACAAAGAGAGCGCGACA	1	0	transposon	LTR	297
9431	CCUCCCGUUACCUCACAGCAGCCACC	1	0	transposon	LINE	G2
9432	GCAACGGGUAACGGGGAAUCAGGG	1	0	ncRNA		rRNA
9433	GCUGCUCGAGCCGGAGGCACAUGCUCGC	1	1	coding gene		CG14305-RA|CG14305
9434	AACCGGGACCGGGAUGAUUCGCGUAGGA	1	0	coding gene		CG3845-RA|NAT1
9435	CGACUCGUUUUUAAAUUUCGCAACAU	1	1	transposon	LINE	Cr1a
9436	CACGGAACGGCGAAGGCAUGGCUGC	1	0	ncRNA		alphagamma-element
9437	UCAACCGCUACGUGAACGAGACCUGUG	1	0	coding gene		CG1882-RA|CG1882
9438	UGGGAUGGACUGAGACGUGCCCU	1	0	transposon	LTR	GATE
9439	UCAAACUUGACUGAAGGGAUCUU	1	0	transposon	LTR	invader4
9440	AGGAAUUGACGGAAGGGCACCAC	1	0	ncRNA		rRNA
9441	UGCCUUAUGGGCAACAUAACCGACGA	1	0	coding gene		CG5045-RA|CG5045
9442	UGUUUCGCAAAAGUUGACCGAACUUGA	1	0	ncRNA		rRNA
9443	ACCGGAUUAAUCGCGAGGCUGCAAGA	1	1	transposon	LTR	mdg3
9444	ACUCGCCGAAUCCCGGGGCCGAGUGAGC	1	no match			
9445	AUCUAGGCUGAGCGGAGAACCAGAGUU	1	0	transposon	Low_complexity	F-element
9446	AGUAUAUGAGCGCCCAGAGUGAUA	1	no match			
9447	AAGGGAAAACUUACCAGGUCCGA	1	1	ncRNA		rRNA
9448	ACAUAUCAAACGCAUACUUAAACGGCA	1	1	transposon	LTR	frogger
9449	AGAGAGUGAUGCCGAGUAUAAAAGUAA	1	0	repeat		trf
9450	UCCCGCAAUAUGGCAACCGCUCGA	1	1	coding gene		CG40609-RA|CG40609
9451	GGAUCUCAA	1	0	transposon	LTR	diver2
9452	GGGUUCCGGGGGGAGUAUGGUUG	1	2	ncRNA		rRNA
9453	UAGGAAUGCACCCGCUACUUUUCG	1	1	transposon	LTR	297
9454	CCAUCCGCUGCGUUCGGGAGCCGGGAA	1	1	coding gene		CG9393-RA|CG9393
9455	UCGAGUGAGACCGGGGGGGCAUUC	1	no match			
9456	UCCAAGCGAACUGUGCCCGAGGCCGUA	1	2	transposon	LINE	R1-element
9457	CCAGAGCGAGGAGGAGGACUAGCUGGC	1	1	Su(Ste)		Su(Ste)
9458	ACCAACGGGAUACCUCAUACUCGAGUU	1	1	transposon	LTR	micropia
9459	AGUUGAAAAAUUCCGAGAAUUGG	1	0	transposon	LINE	Doc2

9460	AGACCCGUCGCCGCGCUCUCCCC	1	no match			
9461	UCGAUAUUAUCCCAGCACUACGAAGC	1	0	no annotation		
9462	GAGAAUCCGAGCACUUUGUUACA	1	0	transposon	LTR	HMS-Beagle
9463	UGUUUCAUCGUUAGACUACUCGGGC	1	2	AT-chX		AT-chX-B
9464	UGUGAGGUUUUGAUCCUUCGAUG	1	no match			
9465	GCACACAGAGGAGAAGAUGAGAGCA	1	no match			
9466	AGGAUGAUGAUACCGACUAAAGGCAG	1	1	no annotation		
9467	UACAUCGAGAAUCUCACUCCCACAGGC	1	1	transposon	LTR	Dm88
9468	AACUUCUUAGAGGGACAGGUGGCGUU	1	no match			
9469	ACGCGCACCAAUGCUUUUUCGAGAUAGA	1	no match			
9470	AGCUUCAUCCAAGAGAACUGUGCCCGUA	1	2	transposon	LINE	R1-element
9471	CGAGACACAACAGAUCUCACAGGUCGGC	1	no match			
9472	CGUACAGAAAGCCAAUUGGCAAGGAA	1	0	transposon	LINE	G2
9473	UCGACUCGGCUAUUGAUACUAUGA	1	0	transposon	LINE	INE-1
9474	CGCCAGCACCAACCCGAAUCCGGCCG	1	0	coding gene		CG6148-RB|Past1
9475	ACCACCCUGAAUGCGCCCGAUCUCG	1	no match			
9476	CCAUCAGAGACUGCGGUGGCCAA	1	0	coding gene		CG12717-RA|CG12717
9477	GAGGCGCCCGGGGUUAUAAUCGUG	1	0	transposon	LTR	gypsy8
9478	GAAACCAAAGCUUUUGGGCUCCGGGGGA	1	0	ncRNA		rRNA
9479	CCGAUACAAAAGUGUGCUAUACC	1	0	no annotation		
9480	ACAAGAAACAGAAGAAGUGCCAUU	1	0	coding gene		CG40609-RA|CG40609
9481	AACCCUUUCAUGCUUGGGAUUGUGAA	1	2	ncRNA		rRNA
9482	ACGAAGCAUACGAAGAAGCAUACGA	1	0	transposon	LTR	invader3
9483	CCGCAGGUACGCCGUCGCUUCCCGGGA	1	no match			
9484	GGAUCAACUGUGAGGUCAA	1	2	no annotation		
9485	GUAACUUCGGGGUAAGGAUUGGCUCU	1	1	ncRNA		rRNA
9486	CCUGAUUGAUAGCGUAGACGACGAC	1	0	transposon	LINE	Rt1a
9487	GCCCGGGUAGACUCAGUCGGUAGAGCUAUUG	1	no match			
9488	AGUGACGAAAAAUAACAAUACAGGACU	1	0	ncRNA		rRNA
9489	UGACGACGAGACAUCCUCCCAGGACAAC	1	0	coding gene		CG5217-RA|CG5217
9490	UUAUAAGGAUGAUGAUACCGA	1	0	no annotation		
9491	CCGGGGGGAAGGUGGCUCGGGGGG	1	no match			
9492	CCUAUCUGGAUAGUCCCGAAGAAACC	1	0	transposon	LTR	opus
9493	CAACCGAGGCAACAAUAAUAGUAAUAA	1	0	transposon	LTR	Stalker2
9494	UAUGGCUCUCUCAGUCGCUUCCCGGGG	1	1	ncRNA		rRNA
9495	GGAGAGGUGGCAGAGUGGUCGAAUGUU	1	no match			
9496	AAAAUAACUCCAGCAGAGACCGUACCC	1	no match			
9497	AGGGGAAUGAUGAUAAGGUCCGA	1	0	transposon	LTR	roo
9498	CUGGCGGGAGCUGGUGCGCGAGAAG	1	0	coding gene		CG13891-RA|CG13891
9499	UCCCCGCGGAAAGACGGAAAGACCCC	1	no match			
9500	GCUACGGUCAAUGUCCCUGAGGU	1	1	transposon	LINE	F
9501	GAGGGAAGUAUGAGUUCCGAAGGC	1	0	transposon	LTR	blood
9502	CCAGCCGUGCCCGAUGUGCCUUGAGCA	1	no match			
9503	AAUCAAAUGACAGAAGAGCCGUGGACAC	1	0	repeat		trf
9504	GAGUACUUCAGCCAGACACUGGACAC	1	2	Su(Ste)		Su(Ste)
9505	UGGGCCCCGAACCCUUCUGGUGUGUUAA	1	no match			
9506	GUAAUUGGAAUGAGUACACUUU	1	0	ncRNA		rRNA
9507	GUUACCCAAAAGCUAAGGAGGAG	1	0	transposon	LINE	Rt1b
9508	CACGAUAAGACCGGACGCAGCCGCAG	1	0	transposon	LTR	gypsy4
9509	CAAGCCAAUCUGGACGUACGGAAUUC	1	0	transposon	LINE	G2
9510	CAACCAGAGAUCUAGAGGGCCCAAGUA	1	1	no annotation		
9511	CAAUCUAGACUAUGUAUCCCCGGG	1	1	transposon	LINE	Doc
9512	GGCCUCCAAGUCCUACAUCGCUGC	1	0	transposon	LTR	diver2
9513	CACCGAUGGCGAAGGCAGCCCCCUGGGC	1	no match			
9514	CCCGCUAGGAGAUGUUGAGAUUGUGA	1	0	transposon	LTR	roo
9515	CCCACAGAGUACGGGGGCUUACUGAA	1	no match			
9516	UCCGAGCCAGAGGACUGUGAGGCA	1	0	transposon	LTR	roo
9517	ACUGAUUAGGUAACUGGAUAUGCUC	1	0	coding gene		CG15498-RA|CG15498
9518	AAACCCGAGAGGUACAUUUUUGCAGCU	1	no match			
9519	GAGAGUGAUCCGUGACUUUGUUGUGC	1	1	transposon	LINE	INE-1
9520	ACCGUGGAGGACAUGGACGGUGG	1	0	coding gene		CG5214-RA|CG5214
9521	UAAUUGAAAAAGAACACCGUAGAG	1	0	transposon	LTR	opus
9522	GAGGAAUUUGAGCUGCGUUGACGCC	1	0	transposon	LTR	ninja-Dsim-like
9523	UCUUAAACGAGUGUUAUUGUGGGCCGG	1	0	ncRNA		rRNA
9524	AACUACGAAACUCAAACUCCCGGACA	1	0	no annotation		
9525	CGGACUCUAAGACGGUGCUGCUUG	1	no match			
9526	AAAACCUAGAUGAUCCUGAGUCGC	1	no match			
9527	GGAAAACCUGAAGAUGAUCCUGAGU	1	0	transposon	LINE	Helena
9528	UCGAUACCGAGGCUUGAACGUCA	1	0	transposon	LTR	Idefix
9529	ACCGGGACGUUCGCCAAUUUCUGUGGC	1	0	coding gene		CG5193-RA|TfIIB
9530	CAACAACAGACAGAACGGAGCAGC	1	1	coding gene		CG7115-RA|CG7115
9531	ACCCAGGUGAUUACCACUAUUAAUGAAA	1	1	Su(Ste)		Su(Ste)
9532	CAAGAUAUGACCAAACAAAGACACU	1	no match			
9533	AUCGCCUCGAGGAGCUCACUUUCC	1	0	no annotation		
9534	UGAAGACAAUUCUUGUUGCUCGGGC	1	1	transposon	LINE	INE-1
9535	CAAAACGUAACGGCAGAGGUAAA	1	0	transposon	LTR	HMS-Beagle
9536	CCCGACCUUACAAAACCAUUCACGU	1	1	transposon	LTR	accord
9537	GCAGAUGUAAUUCCAAGGCGACCCAUG	1	0	no annotation		
9538	GGAUCAACUGUGAGGGGAUCCUCUAA	1	no match			
9539	UCGGAACUUUUGUAACUUUUUCGAAAA	1	0	transposon	DNA	FB
9540	GGAUCAACUGUGAGGAGAUCCUCAA	1	no match			
9541	CCGAACAUCGAAGAUUGGCCUACCCUU	1	0	transposon	LTR	297
9542	GGGGAACGAUCUCAAGUGACUGAC	1	0	transposon	LINE	Cr1a

9543	GUGGCUUUGAGAGCCGCCGUAGUUA	1	1	no annotation		
9544	GGAACACGAAACGAAGCCCCCUGUC	1	no match			
9545	GUUGCAUUAAACAGAUUCCGCAGUGC	1	no match			
9546	UACUUGUUCCCCGGAUAUUUUAGUUAC	1	0	repeat		LSU-rRNA_Dme
9547	ACAUCAAUCGCUGAUGAGACACCGU	1	1	transposon	LTR	accord
9548	UGAGGUGUUUCAUCGUUAGACGGCUCGG	1	no match			
9549	UGACGAAAGAGUUUGCUAAGCCACA	1	0	transposon	LINE	R1-element
9550	AGUGAAACGACAGUGUCCCGCGGA	1	1	transposon	LTR	springer
9551	CAACCAGAGAUCUAGAGGGCCCAGAAA	1	2	no annotation		
9552	CCGAUCGUGAGGUUAAGGUGCCG	1	no match			
9553	AAGUGUGUAAGACAGAUUGAUAGCUCU	1	0	ncRNA		rRNA
9554	UGGAACGAAAUUGGCCUGAUUAGU	1	0	transposon	LTR	GATE
9555	AUAGAAGUUAAUUAUGUACCCACA	1	0	transposon	LTR	DM88
9556	AGGUGUCCCAAGGCCAGCUCAGU	1	0	ncRNA		rRNA
9557	GCUCGGGGAGGUAGUGACGAAAAAUAA	1	0	ncRNA		rRNA
9558	AAGAUCAAAAGUUGGCGCCUGUAGGA	1	0	transposon	LTR	flea
9559	CCCCGCGGAAAGACGGAAAGACCCCA	1	no match			
9560	GUGGCAUUGAGAGGCGCCGUAGUUACCAC	1	1	no annotation		
9561	AUGCGUAGAAGUGUUUGGCGUAAGCCUGC	1	0	ncRNA		rRNA
9562	AGACCCGUGGACGGUGUCGAGGCCGGUGGC	1	no match			
9563	CACCCUGAACGCGCCCCGAUCUCGU	1	no match			
9564	CCAGGCCCAUCUGGUUGAACGUGU	1	0	Su(Ste)		Su(Ste)
9565	AGGGCAAGGACAAGAAGAAGUAGACC	1	0	coding gene		CG12902-RA|CG12902
9566	ACGAUUGCCAGCUCACAGAACAAGGUCU	1	2	no annotation		
9567	AUUCCGUAAACGAAGCCCAAAGU	1	0	transposon	LTR	Rt1c
9568	AAGCCUGAGAGGUGAUGCGUAGCC	1	no match			
9569	ACACUGAGAACUGGGACUGCCUGCUU	1	1	transposon	LTR	GATE
9570	UGCUCUCGAAUGCUUGCCCCGAUAUA	1	no match			
9571	CCCUGCCCGUAUGAAUGUCUCGAUC	1	1	transposon	LINE	INE-1
9572	GUCCUGAGAGAUGGGCGAGCGCCGUU	1	no match			
9573	AAAAUCUCGCGCCGGGCCGUACCCAUA	1	no match			
9574	GGCCGCCGGUGAAAUACCACUACUC	1	no match			
9575	AACACUAGAUGAAAAUGAUCGAGUCG	1	0	transposon	LTR	297
9576	UGCAACUUGAGCGACGCCGUCUCUGCC	1	0	transposon	LTR	diver2
9577	UGGAUUAUGCCUGAACGCCUCUAAC	1	1	ncRNA		rRNA
9578	UACGACGAGAACGAGAUCGGCGCG	1	1	transposon	LTR	frogger
9579	ACGAUGAUUUAAUUCUAGCGUCCCAGU	1	2	transposon	LTR	1731
9580	UUGGGCCCGAACCCUUCUGGUGUU	1	1	transposon	LINE	baggins
9581	AAGGUCUGCUCCCGCAAUCCCAAGUGUCC	1	1	coding gene		CG9130-RA|CG9130
9582	CUGUUCACAUGAACUUGGAAUUCCCAG	1	0	ncRNA		rRNA
9583	CAGUAUUGGUCCAAGAACGCUGCCU	1	0	transposon	LINE	G
9584	CCGACAAACAUCCAACGAACACAACA	1	1	transposon	LINE	Doc
9585	AGGGUUCAUCGUCACCAACAAUGGCGC	1	0	transposon	LTR	gypsy3
9586	AGAACUGGCGAUGCGGGAUGAACCUAAC	1	no match			
9587	GGCCAUGUAACGAAACGGAAGUAC	1	1	transposon	LTR	flea
9588	ACUUGGAUGGGAGACCGCCUGGGAAU	1	no match			
9589	GCAUUCGAGACCCAACGGUACGUG	1	1	transposon	LINE	INE-1
9590	UCUCAUCGUCGUAGAACAACCCGA	1	no match			
9591	UUUAAUGCCUAACGAGGUAGAUGAC	1	0	transposon	LINE	Cr1a
9592	CCAUCCUAAAACUGAUAGACAGAUGGC	1	1	Su(Ste)		Su(Ste)
9593	UUCGAUUUCAAGUACCAGGCGAAC	1	0	repeat		RSP
9594	AAUUACAUUAGACGAGACGGCCCUAU	1	1	transposon	LTR	opus
9595	UAGAUCUUCACCGUUGACUAGCCCCA	1	0	Su(Ste)		Su(Ste)
9596	CCACGAGUCGUAUGCGUAAUAAGAGU	1	0	no annotation		
9597	AGCGUCGAACGAUUUCUGCCCA	1	0	transposon	LTR	Quasimodo
9598	UCAGUCUGAGAAGGGAGGAGUUACCGC	1	0	no annotation		
9599	UAGUCCAAAGUUCAAUGCAGGCCU	1	0	transposon	LINE	Circe
9600	AAAAGUGCAUACAAUUCCCGCUUUA	1	0	transposon	LTR	opus
9601	ACCCCAAUUGUGUUUCGUCCUAAAGA	1	0	transposon	Low_complexity	F-element
9602	AGCUAAAUGAGCGAACUAUUC	1	0	transposon	LTR	gypsy
9603	GACUCUAGAUAACCUCGGGCCGAU	1	no match			
9604	GGCAACUAUGCACCAGUGCCUCCUC	1	0	coding gene		CG4836-RA|CG4836
9605	GAAAUCCGACGCAACAAUCCGGUUGUC	1	0	coding gene		CG33936-RA|CG33936
9606	ACCAUGAAAGGCGUUGGUUGCUUAAA	1	0	ncRNA		rRNA
9607	AUGUCCCAAGAGAAGUUGGCCCAU	1	0	transposon	LTR	flea
9608	ACCCUGGACAGAAGGCUGACUUG	1	0	transposon	LINE	G2
9609	AGCGAUGCGAAUGAUGACAGCGAG	1	0	coding gene		CG14040-RA|CG14040
9610	GAAGAAAUGAAUGCUUUCCUGAGCGA	1	0	transposon	LTR	Rt1c
9611	UGCGCGAAGAAGGAGAACGAGAAGGCA	1	0	transposon	LINE	G6
9612	UGCUAAGGGAUCCAGUUGCGAUAAACU	1	1	transposon	DNA	1360
9613	AAUUACUAAUUCGUUGUGGGCCAGAGU	1	0	no annotation		
9614	UGCACCGAAAAUAUAUGCGAGGUAAA	1	no match			
9615	UAGCUAAUUAACACAAUCCCGGGGGC	1	2	ncRNA		rRNA
9616	GUCGUCGAGUCGGUUGCCACCAUCA	1	0	transposon	LINE	Doc4
9617	UGGUUCCUUAGAUCGUUAACAGU	1	0	ncRNA		rRNA
9618	AUUGAGCGAACCCGGGGAACUGAAACAU	1	no match			
9619	UGCGCAGAGUCAUGCCAGUACUGGACGA	1	0	transposon	LINE	G2
9620	AGCUCGAUCAAAAGGCCCUGGCAA	1	0	transposon	LTR	Dm88
9621	GUCUCCGGACGUGAUCACUGUGUGAC	1	no match			
9622	AGAGCAAUAUGACAGCGUGAUCCAA	1	1	transposon	LTR	Rt1c
9623	UACUCGAAGACACCACGGUGAAAGUG	1	0	transposon	LTR	ninja-Dsim-like
9624	AGAUGGCAAACGUCGCGACGGGGAC	1	2	transposon	LINE	Rt1b
9625	UCCGAUUUCAAGUACCAGACACACA	1	0	repeat		RSP

9626	UCCUACUCGCUAGACGCACGGAACGGC	1	0	ncRNA		alphagamma-element
9627	CCGAACGUAACCCACGGUGCCCCAGUGAUG	1	no match			
9628	CGGCUUAUCACAUCCUGGGGCUGGA	1	no match			
9629	UCCGAAGACAACGAGGAGCAGAUCUGA	1	1	transposon	LTR	1731
9630	AUACCAAGGCCAAGAAACAGAAGAAGU	1	0	coding gene		CG12717-RA|CG12717
9631	UAUUCUAUCAAUCAAAACUGGCCAA	1	0	repeat		trf
9632	GGGGUUCCGGGGGGGAGUAUGGUUGC	1	no match			
9633	UCCUACGACACACUCGAGCCAGUCAAG	1	0	no annotation		
9634	CUCGAAGAAGGGCAAGCGUUUGGGCGU	1	0	coding gene		CG1828-RA|dre4
9635	UAGCUUGGUUCAACAGCGACUGACUGC	1	0	transposon	LTR	mdg3
9636	ACUUGAGACCCAACAACGCAACCGA	1	no match			
9637	CGGGAAAACUUACCAGGUCCGAACA	1	0	ncRNA		rRNA
9638	GAGCCUUCCUGUGGCAGGAGAGACA	1	no match			
9639	AACUGAAUAAAGGAAAUGGACGAGA	1	1	transposon	LTR	invader4
9640	CUCUCUCAGUCGCUUCCCGGGGAAAC	1	2	ncRNA		rRNA
9641	ACAGACUCGGAACGCCUCAUUGGUGA	1	0	coding gene		CG31359-RA|Hsp70Bb
9642	AGCUAAGAUAUACCGGACGUUGUGCAAC	1	1	transposon	LTR	gypsy12
9643	GUUCUUUAUAUCGAACGGACCAGCGUAA	1	1	transposon	LTR	GATE
9644	UGGUGGUGCAUGGCCGUUCUUAGUUCA	1	1	ncRNA		rRNA
9645	ACAAAAUUAAGAGUGCCCAAGACGCGC	1	0	transposon	LTR	springer
9646	CACUAUUACACGAUAUACCUUGAGGAA	1	0	transposon	LINE	Cr1a
9647	UUGAUACGAAUGAGACCCACUGUAA	1	0	repeat		trf
9648	GGUUUAGAUUCCGGAGAGGGAGCCUGU	1	no match			
9649	GCACCGAAAAUAUAUGCGAGGAGA	1	0	repeat		trf
9650	UGCUCUCGAAUGUAUGCCGAUAUA	1	no match			
9651	AGACUGCCCAUACAGACGUUAUGCACA	1	0	coding gene		CG2637-RA|Fs(2)Ket
9652	AUCUCCAAAGGCAUAGGGCAAGAC	1	0	transposon	LINE	G2
9653	ACCGUCCGUACCGAGUCAGUCGCAU	1	0	transposon	LTR	GATE
9654	CUGGUGCCGAGGCCUGGUUGGAUGAGU	1	0	coding gene		CG8827-RA|Ance
9655	AAGGCUAAAUACCGGCACGAGACCGA	1	no match			
9656	UCUCAUCGUCGUAGAACAAGCCCGAC	1	1	Su(Ste)		Su(Ste)
9657	AGGUGAAAUUCUUGGACCGUCGUAAG	1	0	ncRNA		rRNA
9658	AGUAAUAUGAGCGCCCAGAGUGAUA	1	0	transposon	LTR	opus
9659	GCCAACGACCAUACCACGCUGAAUACA	1	0	ncRNA		5SrRNA
9660	AAAGCGAAUGAUUAGAGGUCUUGGGGCC	1	no match			
9661	AACUUCGAUAGCGAACUGACUCUG	1	1	transposon	LINE	R1-element
9662	AUCCGGACAAACACACAAGACCCG	1	0	transposon	LTR	opus
9663	AUCGUGGUAACGGAAGAGCAACAGCUG	1	0	ncRNA		snoRNA
9664	CAGACACGAACGACAUUUUGAUUAU	1	0	transposon	LTR	Idefix
9665	CUGGAUCCGAAUAACUAUUAAGUUA	1	1	repeat		trf
9666	CCGAAAACGGUACAAGUGGCUUUG	1	0	coding gene		CG6538-RA|TfIIFbeta
9667	GAUCCUCAACUGUGAGGGGAUCCCAA	1	no match			
9668	GUGGGACAAUGCCCGGAGCGCAAG	1	0	transposon	LTR	diver2
9669	CGCACACACGAGGAGAGCAGAUGAGAGCA	1	no match			
9670	AUCGAGAACAACAGCAACGGAGACC	1	0	transposon	LTR	invader3
9671	GUAACAGACAACUAGACCGAUUUAU	1	0	transposon	DNA	transib2
9672	UCCGAUUUCGAGUACCAGGCGAA	1	1	repeat		RSP
9673	CAGUUCAUUCUUGAGGCCUUCGAGA	1	0	transposon	LINE	G2
9674	GCACACAGAGGAGAGCAGAUGAAAGCA	1	no match			
9675	CAACUACGAAACUCAAACUCCCGGA	1	0	no annotation		
9676	UCCCCCUCUAAGACGUGCAACCGCGAAU	1	1	transposon	LTR	gypsy6
9677	UGAACCACCCCGGUCAGAUUGCCAA	1	0	coding gene		CG8280-RA|Ef1alpha48D
9678	GGAUAUAUCACUAGACCCAUUACUCU	1	2	transposon	LTR	roo
9679	GCUUCAUCCAAGCGAACUGUGCCCGG	1	1	transposon	LINE	R1-element
9680	ACCUAAACGACGAAAACUAAUAAAGAG	1	0	transposon	LTR	opus
9681	CAAAGGAGUAUUGCCAGAUGACAAGA	1	0	transposon	LTR	gypsy8
9682	UAGUAUCUAGGACUACCAUGGUUGCAA	1	0	ncRNA		rRNA
9683	GCCCGAUGUCCGCGAGAUUGCCGCAA	1	0	transposon	LTR	opus
9684	AAAAACCUAGAAGAUGAUCCUGAGUCG	1	no match			
9685	CAAGAUAUGACCAAUGCAAAGACAC	1	0	transposon	DNA	1360
9686	ACUCGCCGAAUCCCGGGGGCCGAGGGAGUC	1	no match			
9687	ACCCUGAACGCGCCCUAUCUCGUC	1	no match			
9688	CAAAGACGAAAUAGAACUGCACUG	1	0	transposon	LTR	flea
9689	ACGGAAUGGAAUCGAGCGCUGAGUU	1	1	transposon	LINE	X
9690	GAUGGCAAGAUGCGAUGCCGCCGCCCCAG	1	0	coding gene		CG4994-RA|Mpcp
9691	AUACCCGAAGACCUUGAAGUGCAAAU	1	0	coding gene		CG33943-RA|BG642312
9692	UAAACUCGAACCCUCUUAACUAUC	1	0	transposon	LTR	opus
9693	GUCCGGACGGACGCACUGUCACAAUC	1	0	coding gene		CG18279-RA|IM10
9694	AAGCAAAACCAAUCCCCAACCAAACCCCG	1	no match			
9695	UGAACGCGCCCGAUCACGGAAGACCCGC	1	no match			
9696	GCGAGAAAAUAUGCAAUAACUGUGG	1	1	transposon	LINE	F
9697	AGUGCGGGUCAUAAGCUUGCGUUGA	1	no match			
9698	UGAGAUAGUCGGGCUUGAUUGGGA	1	0	ncRNA		rRNA
9699	AGACUUCAAAGUCUACGUGCUAAAAGU	1	no match			
9700	AGGCGGAACGAUACGGCAUUGAUCA	1	no match			
9701	CACUUUCAUGCUUGGGAUUGUGAAC	1	0	ncRNA		rRNA
9702	CAGAAAAGUAAAUAUUCGAUACCUCGA	1	0	no annotation		
9703	CUGACCUGUAUUCGGAUAAUGGAAC	1	0	transposon	LTR	3S18
9704	AGCUUCAUCCAAGCGAACUUGCCCGA	1	no match			
9705	CGCUAAACACACAAGAUUGCCCACC	1	0	transposon	LINE	F
9706	CAGAACAUUAGUGGAAAUGGCCCGU	1	0	transposon	LTR	Dm88
9707	UCCGAUUUCAAGUACAAGACGAACCGAA	1	0	repeat		RSP
9708	UACAAGAUGCAUACGAUGUGGC	1	1	transposon	DNA	1360

9709	GGAUCACUCGGUUCAUGGGUCGAUGAA	1	1	ncRNA		rRNA
9710	GCACUCGAGGCUGUGAGCAUGUAGUCC	1	0	coding gene		CG15572-RA|CG15572
9711	GUGGCAUUGAGAGGCGCCGUAGUUA	1	1	no annotation		
9712	AGACCGAAUCGCAAUUCCUACCGAGAG	1	1	transposon	LTR	Quasimodo
9713	AUCCGGUAAAGCGAAUGAUUAGAGGU	1	2	ncRNA		rRNA
9714	AAUGAGAUUGAGCAAUAACAGGUCU	1	0	ncRNA		rRNA
9715	UCAUCCUCAAGACCAAGACCAGCACGC	1	1	coding gene		CG32486-RD|CG32486
9716	CACCAAGAUAAUCAACCCGUGUUAU	1	0	transposon	LINE	Cr1a
9717	CACGAUUACAAACACCGACUUGCCCAU	1	1	transposon	DNA	Ivk
9718	CAACUAUGAGGGAUCCUCAA	1	no match			
9719	CCUGACGAAGAGCGCAAGAAGA	1	0	transposon	LTR	opus
9720	GGAGCCGGGCGUGGAAUGCGAGU	1	no match			
9721	GAACCGUGAGGCUUAACCUUACAACGCC	1	no match			
9722	CCUAAGACGCCUGCAGAGAACUCACC	1	1	transposon	LINE	G2
9723	ACGAGUAAUAUGAGCGCCCUGAGU	1	1	transposon	LTR	opus
9724	UGGUACUAUAGCCCAAAUCCAAU	1	0	transposon	LTR	Quasimodo
9725	UGUUUCAUCGUUAUACGGCUCGGGCG	1	1	AT-chX		AT-chX-B
9726	CUAAGAGAUAUAUUUACUGACCAA	1	0	transposon	LINE	Circe
9727	GCGACAACCAGAGUUCCCAGUGC	1	0	transposon	LTR	DM88
9728	UUCAAGGGUUACCCCAGGAUGCUGUGC	1	no match			
9729	AGGACGUUAAUAAAGAUGCCGUGG	1	2	coding gene		pseudogene
9730	AAAUCCAAUAAGUGCGAAAAACCGG	1	0	transposon	LTR	Idefix
9731	GAAGAGAUAACUGAGGCGCUAAAGGA	1	0	transposon	Low_complexity	F-element
9732	CAACCAUCCCAACCAAACCCGUCC	1	no match			
9733	CGAGCGUGAUGACUCCUGGGGGCAU	1	no match			
9734	CUCCCUAACAAAAAUUCGAGAGAGC	1	0	transposon	DNA	1360
9735	AGGAAUAAUGGAAUAGAACCGCGGUU	1	no match			
9736	ACCCUGAACGCGCCCGUUCUCGU	1	no match			
9737	CGUUGAACACCCCGAACAAUUCCUGCC	1	no match			
9738	UUAACUCACACCACCAACCGUUACC	1	0	transposon	LTR	invader1
9739	AGUUAGGUCUCCGGACGUGAUCACUGU	1	1	ncRNA		rRNA
9740	GCACACAGAGGAGUGCAGAUGAGAGCA	1	no match			
9741	CAAGGCUAAAUACUACUGACUGACCGAU	1	no match			
9742	AUACAUCGCCGACGGGCGCUGACCCCU	1	no match			
9743	UCAACUGUGACGGGGAUCCUCAA	1	no match			
9744	ACCAAGAUUAUCAACCCGUGCUA	1	0	transposon	LINE	Cr1a
9745	UGUGAAAAGAGAUUGAGUCAGUCAG	1	0	transposon	LTR	invader1
9746	AUUUGCAAAAAACGAGAGCGAUAGA	1	no match			
9747	UGAAUAAACAUACGCCCGGUCGCGC	1	0	transposon	LTR	GATE
9748	CAAGCCAAUCCCAACCAAACCCGUCA	1	1	transposon	LINE	G2
9749	GAUGUCGUAUCACCCAGGACGUCCGAC	1	0	transposon	LINE	INE-1
9750	CGCCCGUCGCUACUACCGAUUGGA	1	1	ncRNA		rRNA
9751	GCGGGCGGCCCGGGUUCGUUUCC	1	1	ncRNA		tRNA
9752	AGAACAACAGUGCAGACGGCGCCCG	1	0	transposon	DNA	H
9753	UUGACAUACAGUAUAGAGAAGGC	1	0	transposon	LTR	micropia
9754	AAGGAGUCUAACACGUGCGUGAGU	1	no match			
9755	GAGGUCUCCGGACGUAAUCACUGUGU	1	2	ncRNA		rRNA
9756	AGACCCGUCGCCGCGCUCUCCCCCCCUCC	1	no match			
9757	UGACAUUUACUCAUCCGAAGGCAAA	1	1	transposon	LTR	McClintock
9758	AUCUGUGGGAUUAUGACUGAACGCCUC	1	no match			
9759	ACCGAAAUGAAUUUGGCUAGCGCU	1	0	transposon	LTR	GATE
9760	GCCGACAAGUACGAGAAGAUGAAGGA	1	0	coding gene		CG3213-RA|CG3213
9761	UGCUUGGACUACAUGUGGUUGAGGGUU	1	1	ncRNA		rRNA
9762	GUCAUUAACUCGCAUUGAUUACGUCCCU	1	0	ncRNA		rRNA
9763	UCUCAUCGUCGUAGAACAAGCCG	1	1	Su(Ste)		Su(Ste)
9764	GGAAAUAGAAGAAAUUGCCGUUAC	1	0	transposon	LTR	G4
9765	CGGGCCGUUCAAGCAGUAUCUGAUCC	1	0	coding gene		CG8058-RA|CG8058
9766	UAAAGAGACAAUUGACGGUACCGCA	1	0	transposon	LTR	Dm88
9767	UGCUCUCGGAUGUAUGCCCGAUA	1	1	transposon	LINE	R1-element
9768	UGGGAACACCGCGUGUUGUUGGCCUC	1	0	ncRNA		5SrRNA
9769	UGUCCCGGGACCCACCAAAUUGUGAGUU	1	0	transposon	LINE	baggins
9770	GUUGCAAGAAGUGCGUCCCAACGUU	1	0	no annotation		
9771	CUACCACGUUGAUAGGCCAGGUGUGGAA	1	no match			
9772	AGUUUUAUCCGGUAAAGCGAAUGAUU	1	no match			
9773	CGUAAACUGCAUUGCCCUGAUAAAU	1	0	transposon	LINE	G2
9774	GGUCCUGACCGCUGCUCACUGUACC	1	0	coding gene		CG6467-RA|Jon65Aiv
9775	UGCCAAACCGGAUGAGAUGGUGCCC	1	0	coding gene		CG30376-RA|CG30376
9776	ACCAGAGACGGUGAGAGUCCGGUAGAC	1	no match			
9777	AAACCUGAGAGUUUGAUCCUGGCUUC	1	no match			
9778	UGCGAAUUGAUGAGUGCGAGCGA	1	0	no annotation		
9779	CUGAAAGAAACAACAACGUGCGAGU	1	0	transposon	LINE	G6
9780	AAAGGUGGAAAUGAAGAAAGUACA	1	no match			
9781	CAAAGGUGGAACUUGAGGGGAUCCUCAA	1	no match			
9782	CUAAGUCCAUAUUGGAAAAGCUCGA	1	2	transposon	LTR	Idefix
9783	CUGAGUGCCCGGGGAUUUUGGCUAG	1	0	coding gene		CG14577-RA|ORMDL
9784	UACUGAUCCACACAGUGAAGACGAGAC	1	0	transposon	LTR	frogger
9785	UGAUCCAAUAAAACUGUACACAGACG	1	no match			
9786	GCGGUUAGACGAAGAGACCCCGUGCAC	1	no match			
9787	AGGCUGGGGCGGCACAUCUGUAAAA	1	no match			
9788	ACCCUCUGAUGAUGUGUUGUUG	1	no match			
9789	AGCUUCAUCCAAGAGAACUGUGCCCC	1	1	transposon	LINE	R1-element
9790	ACCCCUCCCACGGCGCGACCGCUCU	1	no match			
9791	AUCGCUCGAUGUCGUAAGGUGAC	1	0	coding gene		CG40609-RA|CG40609

9792	GGAACCUGAGCGACCUACAGGAUGAGC	1	0	coding gene		CG8349-RA|CG8349
9793	GCGGGUAUAGCUCAAUGGUAGAGCUCU	1	no match			
9794	GCGAAUGAUUAGAGGCCUUAGG	1	0	ncRNA		rRNA
9795	AUAAUCGAAUGAGACCUUGGAGGA	1	0	ncRNA		rRNA
9796	GUGGAUGAGAUCUCGUCACCCGUUGGA	1	0	transposon	LINE	X
9797	CUGUACUCCAUCGGAGAUGCUACGC	1	1	transposon	Unknown	RR50960
9798	ACGGAAAUCCAUACAGACGCUU	1	0	transposon	LTR	diver2
9799	AGGAGUCCGUGGCAUUCAGGUUGGCUA	1	0	ncRNA		7SLRNA-RA
9800	AGAGCACUUAAAAGAAAUUGGAAUUGA	1	0	transposon	LINE	Cr1a
9801	AGUAAUAGCAUCUGUCUACUGUCCGCCC	1	0	transposon	LINE	G2
9802	ACCGUGAGGGGAUCCUCAA	1	no match			
9803	AGUACAUCGAAAAGCGAGUUGACGUG	1	0	transposon	LTR	Max
9804	GGCGAGACGGGUGACAGCCCCGUACGGGA	1	no match			
9805	ACUACCUUGAACGAACAAACGCAGC	1	0	transposon	LINE	Doc2
9806	CCCAAGCAUGCAUGAUGAUAACAUCCGA	1	0	transposon	LTR	flea
9807	AUAAUAAAUAAAAGGUGACCGGCAAC	1	0	AT-chX		AT-chX-A
9808	AAGAGGGCUUGAAACCGUUAAGAGGU	1	no match			
9809	AUAGCUCAGUCGGUAGAGCAUUG	1	0	ncRNA		tRNA
9810	GGGACGGCGGAGCGAGCGCACGGGG	1	no match			
9811	UUGACUAUUAUUUGGAGAACCACAGC	1	0	transposon	LTR	opus
9812	GCCGUCCCUUGACCUGCCCAAGAACAA	1	0	coding gene		CG12736-RA|CG12736
9813	GCUGUCCCUAGUACGAGAGGACCGGG	1	no match			
9814	UACGACUUUGGAUUUAACUAACGGU	1	0	transposon	LTR	diver2
9815	ACCCUUGAUAAACGGUUGGCUUUCG	1	1	transposon	LINE	Rt1b
9816	CAUCGAGAACGUGGCCGACUGGGAGG	1	0	coding gene		CG11020-RB|nompC
9817	UGUUUCAUCGUUAGACGGCCCGGGUUCGUUUCCC	1	no match			
9818	GAGAGACAAAUGGCAUAGCUUCCGAA	1	0	transposon	LINE	Rt1b
9819	GGAGGCCAUGAAGAAAUUGGACGCG	1	0	coding gene		CG32425-RA|CG32425
9820	GUAUCCGCGGCUAGACGGAAAGACCCCG	1	no match			
9821	UGUUGAUCGUUAGACGGCUCGGGC	1	no match			
9822	CCGCGAACUGACGGUACCAUGCGUAA	1	no match			
9823	AGGAGAUCGUGAGUUCGCAUCU	1	1	ncRNA		tRNA
9824	CAGAUAUGACCAAAACAAAGACAC	1	1	transposon	DNA	1360
9825	GAUUCCGGAGAAGAACCUGAGAAA	1	no match			
9826	AGUCGUCUCAGUUCGGAUUGUCCUCUG	1	no match			
9827	UAUUAAGGGAUUACUAGCACACGAG	1	0	transposon	LTR	invader1
9828	UAAGGCGAUAUACCAAUAGAUUGG	1	0	transposon	LINE	Circe
9829	AUGGCUCUCUCUGUCGCUUCCCGGGA	1	1	ncRNA		rRNA
9830	GGUCCUGAGAGACUGGGCGAGCGCCGUU	1	no match			
9831	CUGGGAAACAAAAAUUCAGGUUGG	1	0	transposon	LTR	Idefix
9832	GCGGAGACGUCGGCGGGGGCCCUG	1	no match			
9833	GUCAUUGAAAAAUAACAAUACUUGGGA	1	0	no annotation		
9834	GCCACACCAAGACAUAUUGUACCC	1	0	transposon	LINE	F
9835	CAGGAUAAAUAAGUAGACUUCGAGG	1	no match			
9836	CGACAUCGGAAAACUCUCAGCAGA	1	1	transposon	LINE	Circe
9837	GGAUCUUGGGAGCGGGCGGGCGG	1	no match			
9838	CACUGCAAUACUAAGUCGAUCCAGA	1	0	transposon	LINE	Doc3
9839	UGAUCUUCACCGUUGACUUGCCCC	1	1	Su(Ste)		Su(Ste)
9840	GAGAUAUGCUAGCAGACUGGAGA	1	0	transposon	LINE	G2
9841	AUGGGUAAAACAAAGCGUGUCGUU	1	0	transposon	DNA	RR51476
9842	CGGCCCGAACCCUUCUGGUGUGUU	1	0	transposon	LINE	baggins
9843	GGAUCCUCAACUGUGGAGGGGAUCCUCAA	1	no match			
9844	AGAACCAUCACUGGGGCACCGUGU	1	1	transposon	Low_complexity	F-element
9845	CAGACCGAAUCGCAAUUCCUACCGA	1	0	transposon	LTR	Quasimodo
9846	UAGUUACUGCGAUGGGGGGACGGAGAA	1	no match			
9847	GUUCGUUCCUGGGCGAAAUCACCA	1	no match			
9848	AGCCACCAUACAGCGAGGCCCGUUAUGA	1	0	coding gene		CG8280-RA|Ef1alpha48D
9849	CCGAACUUGAGACCCCAGCCGCUCC	1	0	no annotation		
9850	CCAAGGUUAUGAUUUUGAAAUUGUG	1	0	transposon	Unknown	RR50960
9851	ACCGAACGGUGCCUGGUUACGAGCUCC	1	0	coding gene		CG17344-RA|CG17344
9852	UAGUAGAGGUCUCCGGACGUGAUCACUGU	1	no match			
9853	AAGAGUGCGUGAAACCGUUAAGAGGU	1	no match			
9854	GGAGACGUGGCCGAGCGGCUGAAGGCA	1	no match			
9855	UCCACAACUGUGAGGGGAUCCUCAA	1	no match			
9856	GCAUUCAAUACAGCGAUAGUGGUCCC	1	no match			
9857	AACAUUCAUAACCGAGCGACUGUUCAAUC	1	1	transposon	LTR	GATE
9858	UUAGUGAGGUCUCCGGACGUGA	1	0	ncRNA		rRNA
9859	CAAGCGUUGGAUUGUUCACCCAU	1	0	ncRNA		rRNA
9860	CUUGGACUACAUAUGGUUGAGGGUUGUA	1	0	ncRNA		rRNA
9861	UCCGGAGUGUUGAUCCGAGCCUCAAA	1	0	transposon	LTR	GATE
9862	CUCAAAUAAUAACGGAGGUGUCCCAAGGC	1	0	ncRNA		rRNA
9863	CCCCUAUGACUGUAGACUGUCGCGCC	1	0	transposon	LTR	Rt1c
9864	CUAUCCUGAAUUCACGCCCGCUGUGCC	1	0	transposon	LTR	3S18
9865	GCUGCGAGGUGUGGGAAACCCAGUAA	1	0	transposon	LTR	copia
9866	GGAUCCUCAACUGUGAGGGGAUCCUCAG	1	no match			
9867	UGCCGGGGCGAGGAGGAGGACUAGCUA	1	1	Su(Ste)		Su(Ste)
9868	GCAUCAACGUCAAGGCCCGCGACUAC	1	0	coding gene		CG18594-RA|CG18594
9869	ACCGAUGGACCAAUACACACAUAUC	1	1	transposon	LTR	diver
9870	GGACUACAUAUGGUUGAGGGUUGUA	1	0	ncRNA		rRNA
9871	ACUUGAGACCCAACAACGCAACCGAA	1	no match			
9872	GCUACCGAGACGCCCUCAACGGAGUGC	1	0	transposon	LINE	G6
9873	AAAACAAGCAUUGUGAUGGCCCU	1	2	ncRNA		rRNA
9874	AUAUCAACCAAACCGGGUCUGGUAAAA	1	0	coding gene		CG18397-RA|ssp3

9875	AAGAGCGCUACGGAGAAUAUUACUC	1	2	coding gene		CG3354-RA|Mst77F
9876	CCCGGGGCUACGCCUGUCUGAGCGUCGCU	1	no match			
9877	GACACAGAGGAGAGCAGAUGAGAGCUA	1	no match			
9878	AAAAACCUGAAAGAUGAUCCUGAGUCGC	1	no match			
9879	CUCGGUGUUCAGUACACACAGGGU	1	1	repeat		LSU-rRNA_Dme
9880	CAACCAAAGAGGCAUUGCAUUAGCGAAU	1	0	transposon	Simple_repeat	I
9881	AAAUUACAUUAGACGAGACGGCCCUA	1	1	transposon	LTR	opus
9882	UGUUUCAUCGUUAGACGGCUCGGGUCGG	1	no match			
9883	UACCCUAAUAUGUACAAUGUAAAUUCGU	1	0	transposon	LINE	Cr1a
9884	AAUACAAACAGAGUUCCCGAAUGAGA	1	0	transposon	LINE	Cr1a
9885	GGCUUUGAGAGGCGCCCGUAGUUACCACCC	1	no match			
9886	AAAGAAACGAAGACGAGAACAGCUCGA	1	1	transposon	LTR	invader6
9887	CCGUUAAACAUUUUCGCCAUUACC	1	0	transposon	LTR	micropia
9888	UACCAAAAAACCCGAUGCUUCUGGC	1	0	transposon	LTR	297
9889	CUUGAAAUGAUGAUGACUAAGGU	1	0	no annotation		
9890	AGGACACGUUCAACCAGAUGGGC	1	0	Su(Ste)		Su(Ste)
9891	CAGUUGAUCAAAGCGCCUUGCUGGA	1	2	transposon	LTR	flea
9892	GCACACAGAGGAGAGCAGAUGUGAGCA	1	no match			
9893	CAACGUGGCGGCGUGAAGGGUUCCGGCA	1	0	coding gene		CG6652-RA|CG6652
9894	UGCAACGUUCACCAAUCUUCGUCAGCA	1	0	transposon	LTR	3S18
9895	CAAUACGAACUCGGACGAAGACAUU	1	0	transposon	LTR	Max
9896	AGCCCCGGAAAAUAAAAUAGAAC	1	1	coding gene		CG17342-RA|Lk6
9897	CGGACACCGAACCGGGAAGGCGA	1	0	transposon	LTR	Tirant
9898	AAUCAAGUAACCGAGAAACGUGUUGA	1	0	transposon	LTR	invader2
9899	AAAAACCUCUCCGACCGUUAUCCGAU	1	2	transposon	LTR	accord
9900	GAUGCCAAGACGACAAUCAAAAAA	1	no match			
9901	UUCCCCGGAUAGUUUAGUUACGC	1	1	ncRNA		rRNA
9902	GCUGAGAUAAUUGAGGCUCUGAAG	1	0	transposon	LINE	Doc
9903	UGUGCUAGAAGGCGAUUUUCGGGCC	1	0	transposon	DNA	1360
9904	AUCUCAACUGUGAGGGGAUCCUCUAA	1	no match			
9905	AACGCGACCGAUCCCGGAGAAGCCGGC	1	no match			
9906	CACACAGACAAAAUAUCCUAUGG	1	0	transposon	LINE	INE-1
9907	CUGCGAGAGAUUUGGAGGCAGAAGA	1	0	transposon	LTR	ninja-Dsim-like
9908	CGGGACAUCAUUCUGGCUGAUGAC	1	0	transposon	LTR	GATE
9909	AGCGACAAGAUAGACAGGCGAAACCGCA	1	1	transposon	LTR	micropia
9910	ACGGCGGACGGACGGCGAGGCC	1	no match			
9911	ACAAAACAAAACAUGACACCACGGGA	1	0	transposon	LTR	297
9912	CAAGCUGAUAUCCGACCACAGCGCU	1	0	transposon	LINE	G2
9913	CCAGAUGAUAUUCCGAAAACAGCC	1	0	transposon	LINE	Doc4
9914	CAAAGCGAGAAACCGACCGACUUAGAC	1	0	no annotation		
9915	UAUGCCUGAACGCCUCUAAGGUC	1	0	ncRNA		rRNA
9916	UCUGCUCCGAAAACAAGGCCCGUCGC	1	1	transposon	LINE	G6
9917	AUGCUCCGAAAACAAGGCCCGUCGCUAA	1	1	transposon	LINE	G6
9918	CGUGCGGUCCUUGUAGUCCAUGCGAU	1	0	coding gene		CG11719-RA|Mst98Ca
9919	AACGUGACGAUCCCCACGGCUGGAG	1	no match			
9920	AGGUAUAGAAAGCACCUGCAGCGC	1	0	transposon	LTR	copia
9921	UAACGAGUAAUGGUCUUCAGGG	1	1	coding gene		pseudogene
9922	AUGUCGACUCAUCACAUCCUGGGACUG	1	no match			
9923	CAGACAAGAAGGUGGCAUCGGUGA	1	0	transposon	DNA	H
9924	GGACUGUGAGGGGGAUCCUCAA	1	no match			
9925	AAUCAAUUCAGAACUGGCACGGACUUG	1	0	ncRNA		rRNA
9926	CUGGACUCAUGAGAGUUGGUGGCAGA	1	0	transposon	LTR	diver2
9927	AAGAGGGCGUGAAACCGUUAGGAGGU	1	no match			
9928	GCUUGGACUACAUAUGGUUGAGGGU	1	0	ncRNA		rRNA
9929	AUCGGCCCGAACCCUUCUGGUGUGUU	1	1	transposon	LINE	baggins
9930	CUCAACUGUGAGGGGAUCCUUAA	1	no match			
9931	CGAGUUACGAAGAGAACAGUUGAUGC	1	0	transposon	LINE	Circe
9932	AUGACGAGUAAUGGUCUUCGGGGCGU	1	0	coding gene		CG12717-RA|CG12717
9933	UGCCAUACUGCGAUCCACUGAAGC	1	0	ncRNA		rRNA
9934	AGCAGGACGGAAAGACCCCGGGACCUUUU	1	no match			
9935	GCUUUAAUCACGAGCAAUGCGGUUUA	1	2	no annotation		
9936	UCGAGAACAAACGAUACUGUUCGGUC	1	0	transposon	LINE	G2
9937	UGCACUCGAAUGUAUGCCCGAUAU	1	1	transposon	LINE	R1-element
9938	CGCAAACCCAAAAAGACAAUACACGAU	1	1	repeat		trf
9939	GUGGAGCCGUACCUGUUGGUUU	1	0	ncRNA		rRNA
9940	UCCGCAAAGACGACAAAUUGGCCUAU	1	1	transposon	LTR	rooA
9941	GUUACUAUGAUGACGAGUCUCAAGC	1	0	transposon	LTR	opus
9942	GCCAGGAUGGCCGAGUGGUCAAA	1	2	ncRNA		tRNA
9943	CAUACCAACAGUAGACAAUGUUCCGG	1	1	transposon	LTR	accord
9944	AGCCCGGGAUAGCUCAGUCGGUAGAGCAUUG	1	2	ncRNA		tRNA
9945	UCCGCAAUAUGACAAUCGCUCGAUG	1	2	coding gene		CG40609-RA|CG40609
9946	CAUCUUAUAAAUGAUGAGUCGCUGU	1	0	transposon	LINE	Cr1a
9947	CUGACAAAAAAAAACCGGCGCAUAGCGA	1	0	transposon	LTR	Rt1c
9948	AUCUCAUCGUCCUAGAACAAGCCCGA	1	2	Su(Ste)		Su(Ste)
9949	GACCAGGGGAAUCCGACUGUUU	1	2	repeat		LSU-rRNA_Dme
9950	AAGACAAUCAAGGCUGUAUUAGCA	1	0	transposon	LINE	Cr1a
9951	UCAAGCGAUAUUCGAGGCCCAACA	1	0	transposon	LINE	Doc4
9952	UGCACAAGUUUAGUCUCUUCCGACA	1	0	transposon	LTR	diver2
9953	CACCUGAAGAUCCGCAACCCAGCA	1	0	transposon	LINE	G2
9954	AGAGAGAGGACGGGAGCGGCGGGGGC	1	no match			
9955	UCGCCGGCCGAGGUGGGAUCCCGA	1	no match			
9956	CAAAAGCUCAGCCUAUCGAUCCUUU	1	1	ncRNA		rRNA
9957	GCAUGCUUAAAUUCAAGGCCAUU	1	0	transposon	LTR	roo

9958	GGGGAAUAAGGGUUCGAUUCCGGA	1	1	ncRNA		rRNA
9959	UACGUAAACAACCCCAGGCUGACUCCAU	1	1	transposon	LTR	Rt1c
9960	GUCCUUAACAAGAACGACGUUGUCC	1	0	transposon	LTR	diver2
9961	CUCGCUUGAAAUGAUGAUGACUAAGGU	1	0	no annotation		
9962	UUCGUGGAGUGAUUGUCUGGUU	1	no match			
9963	UAUUAUUAAGAUAGGGCUUGUAGGU	1	no match			
9964	GGCGGCGACGACCCAUUCGAACGU	1	no match			
9965	GAUUCCGGAGAGUGAGCCAGAGAA	1	2	ncRNA		rRNA
9966	ACCAACGUACAAAUCGAGCAACUCGUC	1	0	transposon	Low_complexity	F-element
9967	GAAGUGGGUUUCGGCUACGAGUAGG	1	0	transposon	LINE	G6
9968	UCAGGUCGGAGCGCGGUGAGAAUGA	1	2	Su(Ste)		Su(Ste)
9969	UAUGUGCUCGUUCUGGGUAAAUA	1	0	ncRNA		rRNA
9970	AACUACAUUGAUGCUUAUCCCGAA	1	0	transposon	DNA	Ivk
9971	AGAGAGCUCUGAUGACGAACCCG	1	0	transposon	LINE	G6
9972	AGAGUCUACACGUGCGCGAGUC	1	no match			
9973	GGCACGAAGACGUUGUUAUAGCGAG	1	0	no annotation		
9974	GAACUUGAGAUCGCCCCUGGACGAC	1	0	coding gene		CG42398-RA|SteXh:CG42398
9975	AGAGGGCGUGAACCGUUAAGAGGU	1	no match			
9976	UGACGCCAGAGGUAGACGACCGG	1	0	transposon	LINE	R1-element
9977	AAGGAAAGUGCGAAAUUCAUCUGGA	1	0	transposon	LINE	Doc3
9978	AAUCAACUGUGAGGAUCCUCAA	1	no match			
9979	AGGGGAUGAGCUGUGGGUAGG	1	no match			
9980	AACACAUCUCGAUCCCAUGUCCGUUGA	1	0	transposon	LTR	Idefix
9981	AGGGUUACUAUGAUGACGAGUCUCAA	1	0	transposon	LTR	opus
9982	UUAAUGCCGGAAGCGAUCCCACUG	1	0	AT-chX		AT-chX-B
9983	UACUGAGACAUAGUGACAUCGGCCA	1	1	no annotation		
9984	GGAGCCGGCGUGGAAUGCGAGUGCCU	1	no match			
9985	UCCGAAGUAGACCGACCGAAACAGUCGC	1	no match			
9986	CUCAGCGAUGUGUGGGCCAAGUCAAC	1	0	Su(Ste)		Su(Ste)
9987	AAUCUCGCGCCGUGCCGUACCCAU	1	no match			
9988	CUGGAUACUAUGUAUCCCCGGGUAA	1	no match			
9989	UCCUGACACUGGAAUUUCGUCGAAGGC	1	1	coding gene		CG17949-RA|His2B:CG17949
9990	AUCUCCAAAGGCAUAGGGCAAGACAG	1	0	transposon	LINE	G2
9991	UUAAGUCACGAUAGCCUAUGAUAUACA	1	no match			
9992	ACAGAAGGCAACCGAGGGAUCUCAU	1	0	transposon	LINE	X
9993	CGACUCGUUUUAAAUUCGCAACAAG	1	no match			
9994	AAACUGAAUAAACGAAAUGGAUGA	1	0	transposon	LTR	invader4
9995	AAGAAGAGGAAGGCGACGACGGGCG	1	1	coding gene		CG11857-RA|CG11857
9996	AGCGGGGGAAGAAGACCCUGUUGAGC	1	2	repeat		LSU-rRNA_Dme
9997	AAGGCCGAGAACUGAUGACGAGUUA	1	no match			
9998	UAGACUUAAAGAUGAUGAACCCAUU	1	1	transposon	DNA	NOF
9999	UAUCAAGAAAUCCUUUAUCACCCGA	1	0	transposon	LTR	micropia
10000	UGCCCUGAAUGUAUGCCCGAUAUAC	1	no match			
10001	AACACGGGAAAACUUACCAGGUCC	1	0	ncRNA		rRNA
10002	AUGAAUGAGAUGCAGCUAUAUAGUAGCA	1	0	no annotation		
10003	UACCAUGGUUGCAACGGGGUAACGGGGGA	1	no match			
10004	AGCCGAAGUCAUUGUGCGCCAGAGUUA	1	1	transposon	Simple_repeat	I
10005	ACGAGCGCACGUGUUAGGACCCGUAA	1	no match			
10006	CCGGACUAGAAGUGCCCGCCCCACC	1	0	transposon	LINE	Doc
10007	UACCCCCAAUGAGAUGGCCUUUGGCGA	1	0	transposon	LTR	Quasimodo
10008	AAGCCCGAUGAACCUGAAUAUCC	1	0	ncRNA		rRNA
10009	GGUCCCCGCGGAUGCUCACGCAAUGUGAU	1	no match			
10010	UGUUUCAUCGUUAGACGGCUACGGGCGG	1	no match			
10011	AAAGAAAGUAUUGGAUUUUCGGGUU	1	0	transposon	DNA	HB
10012	CGCUCAAAAACUCCCAUCCUUAUG	1	no match			
10013	AUUAGAAAGUGACCCUGUUGU	1	1	transposon	LTR	Idefix
10014	UCCGAGGAACCCAUUGAUCCAGCUGC	1	0	no annotation		
10015	AUUGUGGGCCGGUACUAUUACUUU	1	0	ncRNA		rRNA
10016	CACAAAGUUAUUCUUCUACCGAUGGAGG	1	0	transposon	LTR	Rt1c
10017	GAAAACCUGAAGAUGAUCCUGAGUCGC	1	0	transposon	LINE	Helena
10018	GCGACGAGAAAAUCUGGAUACCCUGU	1	1	transposon	LINE	G5
10019	AUGGCAUAAAAUAGAUAACCUGAUUG	1	0	transposon	LINE	Cr1a
10020	CCACAGAGUACGGGGCUUACUAAAGU	1	0	transposon	LINE	baggins
10021	UCGGCCAUAGUGAUCCGGUGGUCCCGA	1	no match			
10022	UAGAUGUCCUGGGCUGCACGCGCGCU	1	0	ncRNA		rRNA
10023	GUGGGUCUUGUUCUUAGUUCGUGGAGU	1	no match			
10024	UGUCCCAAGGCCAGCUCAGUGCGGAAA	1	1	ncRNA		rRNA
10025	UGUUCCCCCGGAUAGUUUAGUUACG	1	no match			
10026	CACACUUGACGAAAAUGAUCGAGU	1	0	transposon	LTR	McClintock
10027	GGUCCUGAGAGAUGGGCCAGCGCCGUU	1	no match			
10028	AGAGAGAUAGUUAUCAGGGUUCACCAAA	1	0	coding gene		CG32396-RA|CG32396
10029	GUGCCAACGUCCCGAGAACUGAAAGAA	1	0	coding gene		CG3800-RA|CG3800
10030	UCAGACGAAGUUCAAGGUUACCCCAGGA	1	0	repeat		trf
10031	UGUUUCACUCGUUAGACGGCUCGGGC	1	no match			
10032	UGGAAUGUUUUCGUCCCACCCAAGUUC	1	0	transposon	LTR	GATE
10033	GUGAUCGUCUAGUGGUUAGGACCCC	1	0	ncRNA		tRNA
10034	AGAACAACAGUGCAGACGGCGCCC	1	0	transposon	DNA	H
10035	UUAAAUGUAAAUGACCUGCGACGCGG	1	0	no annotation		
10036	AUGGGGCCCCGAAACCCCUUUCUGGGUGUGU	1	no match			
10037	GCCGCGGAGCCUCGGUUGGCCUCGGAU	1	no match			
10038	GGCUCCUCAACUGUGAGGGAUCCUCAA	1	no match			
10039	UAAACUACUACAACAACGCCGACUG	1	0	transposon	LTR	Stalker2
10040	GACUAGCUGGCCGAGUAUUUGAUAUUC	1	0	Su(Ste)		Su(Ste)

10041	UCCUCAACUGUGAGGGGGAUCCUCAA	1	no match			
10042	AUCAACUGUUCAAUC	1	1	transposon	LINE	G3
10043	UAGCUAAUUAACACAAUCCCGGGGCGUU	1	0	ncRNA		rRNA
10044	UAGUUCACCGAAGCGCCUUACUGGA	1	2	transposon	LTR	diver2
10045	CUGUCUAUUCGCCAGGUCCUGAC	1	1	transposon	LINE	Cr1a
10046	AAGGGAAUCAACGAGGCGGAC	1	2	coding gene		CG31029-RA|CG31029
10047	UGAUUAAUAGGGACGGUCGGGGGC	1	no match			
10048	AAACGCAUGCACUCCCGCCUGGA	1	no match			
10049	AGGAGAGAAAGAAGUAACGGGGCCG	1	0	coding gene		CG6770-RA|CG6770
10050	ACCUGAUUGAUAGCGUAGAUGACGACC	1	0	transposon	LINE	Rt1a
10051	UGCGGCUUAAUUUGACGCAACACGGGA	1	1	ncRNA		rRNA
10052	AGCUAACGACAAGUAGGGCGGGACA	1	no match			
10053	AUUGAGAUAGUCGGGCUUGAUUGG	1	0	ncRNA		rRNA
10054	CGCGAUGAUUAAGAUGGAUGUCAUU	1	no match			
10055	AAAACCAAUAUGGACGUAUGGAA	1	0	transposon	LINE	Porto1
10056	AAGAACGAAAGUUAGGGGAUCGAAGA	1	no match			
10057	CGAGACUACAAACGGAUUGCGGACCG	1	1	no annotation		
10058	CGCAGAGUCAUGCCAGUUGGAC	1	no match			
10059	UGGUUCCCUCCGAAGUUUCCCUC	1	1	ncRNA		rRNA
10060	GCGGAUUAUUACAGAUGAUGGAUCA	1	2	transposon	DNA	H
10061	CAAACCAAUCCCAACCAAACCCGUCAUU	1	2	transposon	LINE	G2
10062	GCUGGUGCAUGCCUGUGGUCCCAACU	1	no match			
10063	CGCAAACCCAAAAUUACAAUGCACGA	1	1	no annotation		
10064	AAGAGCGAUAUCCGAUGCCGAU	1	2	transposon	LTR	diver2
10065	GCGCCCUGGGAGGCACCAUCCCCGA	1	0	coding gene		CG18211-RA|betaTry
10066	GAAAAUGUGACGAAAUCUGGAUACCG	1	2	no annotation		
10067	GAAAUGACUCUGUCCCGAUUCAAGA	1	0	transposon	LINE	R1-element
10068	GAAAGAAGCCCCUGUUGAGCUUGACUCU	1	2	repeat		LSU-rRNA_Dme
10069	GAUGCAAACUCCGAAUACCAGCAAGUU	1	no match			
10070	UGCUGAAGAAAGGAGAUGCGUUGCCAA	1	0	transposon	LTR	diver2
10071	GAUGAUCGAUGACGAACUACGAAGG	1	0	transposon	LTR	ninja-Dsim-like
10072	CGCAAACCCAAAUCGACAAUGUACGA	1	1	repeat		trf
10073	UUUAAUCCGAAUGUAGCAGAAGUGG	1	0	coding gene		CG30438-RA|CG30438
10074	CAAAGGCAAAAUAUAUUGUGGGUGCA	1	1	transposon	LINE	TART-A
10075	GACGAACUACCGAGCCCGAGCCUUAAU	1	0	transposon	LTR	Max
10076	UGCCCCACUACGUACCGCACCGCU	1	0	coding gene		CG11567-RA|Cpr
10077	CUGCGAUCCACUGAAGCUUAUCCUUUGC	1	0	ncRNA		rRNA
10078	UCUCAUCGUCGUAGAACAAGCCCGGGG	1	1	Su(Ste)		Su(Ste)
10079	UGCCCUCGAAUGUAUGCCCGAUA	1	1	transposon	LINE	R1-element
10080	UUCGGGAUAAGGAUUGGCUCUAAGG	1	2	ncRNA		rRNA
10081	AACCUAAUCACGUUCGGUGCCCGAAAUU	1	no match			
10082	CGUAAACAAUUCAAGAUCUGGCCGG	1	no match			
10083	GGGGCCGUAGCUCAGUUGGUAGAGC	1	no match			
10084	UGCUCUCGAAUGUAUGCCCGAUAUUAU	1	no match			
10085	GAUCCCCCUGAAGUCCAAUGCAC	1	0	transposon	LINE	baggins
10086	UCCCACGGAACACCAGAGCAAUCCCA	1	0	transposon	LINE	G
10087	UUCUGACGAGAACUCGAGCCGCGUAGC	1	0	no annotation		
10088	CCCGGGGGCGUUCUAUAUAGUUAUGUAU	1	2	ncRNA		rRNA
10089	AACCAGCUAGAGUACAGAUCGAAA	1	2	transposon	LINE	jockey
10090	UCCGAGCUAAUAUGGCACCAAGCU	1	1	no annotation		
10091	GAACGAGAUUCCCACUGUCCCUAC	1	2	repeat		LSU-rRNA_Dme
10092	UGUAUCAUCGUUAGACGGCUCGGGC	1	1	AT-chX		AT-chX-B
10093	GAAAGCACCGUUUCCCGUCCGAUCAA	1	no match			
10094	AAUGCCGUUCGUCCUGGAGUCCCGA	1	0	coding gene		CG10252-RA|CG10252
10095	GAGGUUCGAAGGCGAUCAGAUACCGCCC	1	0	ncRNA		rRNA
10096	AAAGUCGAACUGGAUCCAUCGAGUC	1	0	transposon	LINE	Doc3
10097	GUGCGCAAGUCCCGGAAUGAGUA	1	0	coding gene		CG8954-RA|Smg5
10098	CGCACCGUGAUUGGCUCAACCGGCGAA	1	0	transposon	LINE	G
10099	AGCAUUCGAGACCCAACGGCACGUGA	1	1	transposon	LINE	INE-1
10100	GCUCCCUGGCGAAACUGGCCCCCA	1	no match			
10101	GCAAAAACCAGCGAUAACGAAGAAAGU	1	1	transposon	DNA	RR51476
10102	CGUGAUGGGGAUCGGGGGAUUGC	1	no match			
10103	AUGCCGGGGCGAGGAGGAGGACUAGCU	1	0	Su(Ste)		Su(Ste)
10104	ACUGGCGCCGUGGGAUGAACCAA	1	1	ncRNA		rRNA
10105	CAGUUACGAAAAAUAUUUCCCGCAG	1	0	transposon	Low_complexity	F-element
10106	GUAUCUGAUAGUCGCGGACUCGA	1	1	transposon	LINE	INE-1
10107	UAAACCUGAAUUGAGCACUAAAUCGU	1	0	transposon	LINE	BS
10108	UCCAGAUAAAUCACCGUGCCUACGU	1	0	transposon	LINE	INE-1
10109	AUGUCUGCCCUAUCAACUUUUGAU	1	0	ncRNA		rRNA
10110	GAGGAAUUUGAGCUGCGAUGACGCCGAA	1	0	transposon	LTR	ninja-Dsim-like
10111	GAAGCUCGCGAAGGAAAAAGAC	1	0	coding gene		CG1979-RA|CG1979
10112	AAGUUAACUAUCCCAAAGGCAUCAC	1	0	no annotation		
10113	AUGAUAUGCACCGAAACUAUAUGA	1	0	Su(Ste)		Su(Ste)
10114	UGUUCAUACAAUUGAGGCUGGAG	1	0	transposon	Low_complexity	F-element
10115	GUGGCCUGAUUGCUAUGCACCGAAAA	1	0	repeat		trf
10116	CGGGACAUCAUGCUGGCUGAUGACC	1	0	transposon	LTR	GATE
10117	UAAGCCCGAUGAACCUGAAUAUCCGU	1	0	ncRNA		rRNA
10118	GAAUCACCGAGAGCCACAUUCUGCCGU	1	1	transposon	LTR	gypsy6
10119	ACCGAAGAAAACAAAGCCUCCGCC	1	0	transposon	Low_complexity	F-element
10120	GCAAUCACCGCCACUGCAGCCCAUGUCC	1	0	coding gene		CG5987-RA|CG5987
10121	AUGGAACACAUAGAGGCUUCGAGGA	1	0	transposon	LTR	HMS-Beagle
10122	CUACGGGCUGGACCGGACCGGCAAGGG	1	no match			
10123	CAGAGGAGAAGACCGCAGUACUGAACU	1	0	coding gene		CG7004-RA|fwd

10124	GCGGAUAUAAAAGCAUCGGGGCGG	1	1	transposon	LTR	opus
10125	UGCUUCCGGGAUUUUGACGCCGUUAAU	1	0	transposon	LTR	opus
10126	AAUAAUCAAAGACAAUCUCAGCCCGAU	1	1	transposon	Low_complexity	F-element
10127	UGAGGGAACCACUUUCAUGCUUGGGAUU	1	no match			
10128	AAGGAGAGCAAUUUUGGCCGUCACUA	1	0	transposon	DNA	1360
10129	UCCCGCAAUAUGGCAAUCGCUCGAU	1	0	coding gene		CG40609-RA|CG40609
10130	CAAACCAAUCCCAACCAAACCCGUCC	1	0	transposon	LINE	G2
10131	AAACUGCUGCAAAUGGACGAGCCGA	1	0	coding gene		CG4799-RA|Pen
10132	CAAGCUAGCCUAUGUAUCCCCGGG	1	1	transposon	LINE	Doc
10133	UGCGGGAUGAACCGAACGCCGGG	1	no match			
10134	CUGUGCAAAAGCAAAAGCGAACGCCCG	1	no match			
10135	AUGGAACAGAAACCCGCGCCAGACGG	1	2	transposon	LTR	ninja-Dsim-like
10136	AAAAACCUGAAGAUGAUCCUGAGUCU	1	2	transposon	LINE	Helena
10137	CUACAUUCGGCUCUGUAACUCGGCAAU	1	no match			
10138	ACCACCCUGAACGCGCCCGAUCUCCUC	1	no match			
10139	UCCAAAUGCAAAAAUCAAGAGAUGGA	1	0	transposon	LTR	Burdock
10140	CUCUCAACGGGAUUGAAGGCGAGCGGC	1	1	coding gene		CG7361-RA|RFeSP
10141	UGGGAACACCGCGUUUUGUUGGCCU	1	1	ncRNA		5SrRNA
10142	GGAUCCUCAACUGUGAGGGGGUCCUCAA	1	no match			
10143	ACGGGAAACCUCACCCGGCCCGGACACGG	1	no match			
10144	ACCCAUGAGCGAGUCAUUCCGACAA	1	0	transposon	LTR	blood
10145	AACGAGAGAAGCAUUUUUGCCGCAAAC	1	0	transposon	LTR	gypsy12
10146	AAUACUGUGGCGACUUUAGGUAGGAGA	1	0	coding gene		CG12250-RB|CG12250
10147	AAAUCAACAUACACUACUCCGAAUG	1	2	transposon	LTR	Idefix
10148	GUCUUCCUUAAACGUUACCACCUUU	1	no match			
10149	UGUCGGCUCUUCCUAUCAUUGUGAAGC	1	0	ncRNA		rRNA
10150	UCAACUGUGAGGCGAUCCUCAA	1	no match			
10151	CGGACCACCAACCGUUGACAUUCGCG	1	0	transposon	LTR	gtwin
10152	CAACCAGAGAUCUAGAGGGCCAAA	1	1	no annotation		
10153	GUUUGACUGGGGCGUACAUCU	1	no match			
10154	CAAAAGGGGAACUGAGGAACGCAU	1	0	transposon	LTR	invader3
10155	UAACGAGAGUAGUGAAGAGGAAGAG	1	1	transposon	LTR	Dm88
10156	CUCCGGGGGAAGUAUGGUUGCAAAGCUG	1	0	ncRNA		rRNA
10157	GUCUGGUUAAUCCCGAUAACGAACGAGA	1	1	ncRNA		rRNA
10158	GGAGGGAUGGGACCGAUGAAUCCGCGC	1	0	coding gene		CG7187-RA|Ssdp
10159	GACAAAUCUACGAACAUACGACGUCAA	1	1	transposon	LTR	diver2
10160	GGCUGCAAUUGUUUCCCGUGGGACUAC	1	0	transposon	LINE	baggins
10161	AGUGUGACCCACCGCCCGAUCUUA	1	0	no annotation		
10162	GCAGCCCAACGAGCCAAUAAACGGAUCA	1	1	coding gene		CG3779-RA|numb
10163	AUUAGUGACGCGCAUGAAUGGA	1	0	ncRNA		rRNA
10164	CAUGAUGCAACAUGCUGACCAUAGGU	1	1	transposon	LTR	diver
10165	ACUCGCCGAAUCCCGGGGCCGAGGA	1	no match			
10166	UGACCCGUGAUGGAGGCGGCCGGGAU	1	no match			
10167	UGCGAAAUAAUUGCCAGAGUCAC	1	0	transposon	LTR	Quasimodo
10168	CAAAGCGGGACCUCUUAGGCAAGAC	1	0	transposon	LINE	G3
10169	GAUCCUCAAUUGUGAGGGGAUCCUCAA	1	no match			
10170	ACGAACGAGACCUUAACCUACUAAA	1	no match			
10171	CCGCCCCCGGCCCCGCGUCCUCC	1	no match			
10172	GCCUUCGAGAGAUAUAUGGCCCAGUUGA	1	1	coding gene		CG11552-RA|CG11552
10173	AUACAUGCCGACGGGCGCUGACCCCU	1	no match			
10174	ACUACUUUCAGGACACUUUCAACCAGA	1	1	Su(Ste)		Su(Ste)
10175	UUGGACCGAACACUUUGAACCAAGUAUU	1	0	Su(Ste)		Su(Ste)
10176	ACGAUGGCCACACUGUCUCCUCCUGAAA	1	no match			
10177	UCACGUUAUCCUAGAAGAACAUUGAAGC	1	no match			
10178	AACACUAACUGGAGGACCGAACCGACU	1	no match			
10179	AAUACAAACAGAAUUCCCGAAUGAGA	1	0	transposon	LINE	Cr1a
10180	CGACCCAUUUCUGUUACUUUGUGUC	1	0	repeat		trf
10181	ACAAUAGCUGAGCCGUACUAAGUGGCC	1	no match			
10182	CCCGAUGAGAAGAAUGGCCCGUAAA	1	0	transposon	LINE	RR49783
10183	CGCUAGAAUCCGCAAGAAGGGCGAGG	1	0	coding gene		CG4164-RA|CG4164
10184	GAUGGUGUAGCAUCGAUCGCUGUGUGGC	1	no match			
10185	ACAAACGUUCUUAGAUGAGUAUUCCGA	1	0	transposon	LTR	Rt1c
10186	UAAUGUUGGCCAAUUUCGAGUAUA	1	0	transposon	LINE	Cr1a
10187	AUAAUAAAAGGUGACCGGCAACGACGU	1	2	no annotation		
10188	CGGUCUGGUGGUUUGAGCGGUG	1	no match			
10189	GACCACGCCGUCACCGAGGUCCUGCC	1	0	transposon	LTR	ninja-Dsim-like
10190	UCACACUCAUCAACCAAAGGACGU	1	1	no annotation		
10191	UCUUCUAGGGAUGGCGUGAUUUUAAA	1	2	Su(Ste)		Su(Ste)
10192	UUGAGACCGCCCCUGGACGACCCAG	1	1	Su(Ste)		Su(Ste)
10193	CUGUGAGGGGGAUCCUCAA	1	2	no annotation		
10194	AUAGCUAAUUAACACAAUCCCGGGGCG	1	0	ncRNA		rRNA
10195	GUUCAUCUCCGGUAGCAUAAAGA	1	0	transposon	LTR	McClintock
10196	GCCGUUCUUAGUUCGUGGAGUG	1	0	ncRNA		rRNA
10197	ACAAGCUAGACUAUGUAUCCCCGGGU	1	0	transposon	LINE	Doc
10198	GAUCCAGCUGUGAGGGGAUCCUCAA	1	no match			
10199	AAAAGGAGCGAUUUUUGGCCGUCACUA	1	no match			
10200	CACUUCGAACCAGGUGCCAGAGGG	1	no match			
10201	UAUAGCUAAUUAACACAAUCCCGGGGCG	1	0	ncRNA		rRNA
10202	GUAACUUCGGGAUAAGGAUUGGCACAAU	1	no match			
10203	UGCCAAUUACAUUAGACGAGACGGCCCA	1	1	transposon	LTR	opus
10204	GUCGGAGGCUGGCGGCGGGGGGGU	1	no match			
10205	GGAGCCGUACCUGUUGGUUUGUCCCAUU	1	0	ncRNA		rRNA
10206	CCCAAACGCAGACAUCGAGCGGUCA	1	0	transposon	Low_complexity	F-element

10207	CGCAAACCCAAAAUUACAAUGCACGAC	1	1	no annotation		
10208	CGCACUCUACGAAGCAGUCCAAGCCGA	1	0	transposon	LINE	G2
10209	CUCAUCCUGGCAACAGGAACGACCAUA	1	0	ncRNA		rRNA
10210	GAUCCUCAACUGUUAGGGGAUCCUCAA	1	no match			
10211	GGCGGCCCGGGUUCGUUUCCC	1	1	ncRNA		tRNA
10212	ACCGAUGGCGAAGGCAGCCUUCUGGC	1	no match			
10213	GCCCGGCUAGCUCAGUCUGUGGAGCA	1	2	ncRNA		tRNA
10214	CGAAAAUGCAGACCUAUUGUCAUCCG	1	0	transposon	LTR	flea
10215	UCCAACGAGAGAGCGGAGAGCCAGGC	1	1	transposon	LTR	mdg3
10216	CUAAAGUCCAUAUUAGAAAACUCGA	1	no match			
10217	CGUCGCCGAGAUGGCCGAGCAGGUUGAUUC	1	2	coding gene		CG17927-RA|Mhc
10218	AAAAUCAACAUACACUACUCCGAAUG	1	no match			
10219	GGGAAACUUAAUGACCGGCAAAAUU	1	0	no annotation		
10220	GAUGUCCUGGGCUGCACGCGCGCU	1	0	ncRNA		rRNA
10221	CACAGAGUACGGGCUUACUAAGUCA	1	no match			
10222	AACCAAGCGCGGGUAAACGGCGGGA	1	no match			
10223	AUCUGCAUGUGGGUGACCGGGUCGU	1	0	coding gene		CG4836-RA|CG4836
10224	AUGCUCUCGAAUGUAUCCCCCGAUA	1	no match			
10225	UCCCAUCGUCGUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
10226	UGUUUCGCAAAAGUUGACCGAAC	1	0	ncRNA		rRNA
10227	ACUCAACCUGAGAGUUUGAUCCUGGCUC	1	no match			
10228	GCAUUCGAGACCCAACGGCACUUGACAC	1	1	transposon	LINE	INE-1
10229	ACCCUGAACGCGCCCCGAUCUCGU	1	no match			
10230	UGUUUCAUCGUUAGACGGCUCGUCG	1	no match			
10231	GCAUCAGUUAAAGGCGAAUCGCCCGAU	1	0	transposon	LINE	RR49783
10232	UGGUGACCACGGGUGACGGGGUAA	1	no match			
10233	UUUGAGAGGCGCCGUAGUUACCACCC	1	0	no annotation		
10234	CUGCAGGAACGCUACUCUCGGGAGA	1	1	coding gene		CG30456-RB|CG30456
10235	UGGGAACACCGCUGUUGUUGGCCU	1	no match			
10236	GAAGCAAUCAGAAAAACGACGGUCGAA	1	2	coding gene		CG40271-RA|CG40271
10237	UCUGAUGUUACGCCCGUAGAGAUA	1	1	transposon	Low_complexity	F-element
10238	GAACUUGAGACCGCCCCUGGACGACC	1	0	Su(Ste)		Su(Ste)
10239	UGGAAAUUAACCGAUGCAGACAAGAGG	1	0	transposon	LTR	rooA
10240	GAGACCAGUCGCCGCGCUCUCCCCCCUC	1	no match			
10241	GCCAAGAGUUCCCGUGGUAGGCAGUAG	1	0	ncRNA		7SLRNA-RA
10242	ACUCAUUUUACCCAAUUACAACAUG	1	0	no annotation		
10243	UGACGCGAUGUGAUUUCUGCCCAGUGC	1	2	ncRNA		rRNA
10244	CCACAAAGAAAAUGGCAUCCUCUGA	1	0	transposon	LTR	Rt1c
10245	AAAUGCUGGCGGAGCAGGCGGUCCGGGA	1	1	coding gene		CG2252-RA|fs(1)h
10246	UGAGACUACAAACGGAUUGCGG	1	0	no annotation		
10247	UGAAAGUAGUUGGUGGGGCACGUGGC	1	no match			
10248	UCGCAGACCCGACGCACCCCCGCCACGC	1	no match			
10249	UAUCCAUCAGUCCCAGCCCAAGU	1	0	transposon	LTR	McClintock
10250	CAGUAUUGGUCCAAGAACGCUGCCUU	1	0	transposon	LINE	G
10251	UUACUGACGAAUUUAGUUACAAGCAG	1	0	no annotation		
10252	CCCGGUAGCACGAGUCCAUGCGC	1	0	transposon	LTR	ninja-Dsim-like
10253	AGCAAGUGGACCCGGUUUCUGGAGA	1	0	coding gene		CG31226-RA|CG31226
10254	ACGACGCCAGGGGCUGCCCGCCG	1	1	coding gene		CG2049-RB|Pkn
10255	AAGCCCGACCGUUCUGCCCAGCUGCCGC	1	0	coding gene		CG14011-RA|CG14011
10256	GAGCCGGGCGUGGAAUGCGAGU	1	no match			
10257	AACCCUGAACGCGCCCGAUCUCGUUC	1	no match			
10258	GCUCUCGUAAUGUAUGCCCGAUAU	1	no match			
10259	UGAUCCAAUAAAAACUGUACACAGAC	1	0	transposon	LTR	invader6
10260	UAAUAAGGCAAUGAUUGUUGAGUGC	1	0	transposon	LINE	Cr1a
10261	CACACAGAAGGAGAGCAGAUGAGAGCA	1	no match			
10262	UGUUUCAUCGUUAGACGGCUUGGG	1	1	AT-chX		AT-chX-B
10263	GCAAGAACACCCGGUGGACCCCGAA	1	0	coding gene		CG9821-RA|CG9821
10264	AAACCAAUCCCAACCAAACCCGUCAA	1	0	transposon	LINE	G2
10265	UAGUACAAUAUAUCCGACUGGGCCAA	1	2	transposon	LINE	baggins
10266	CACAGGUACGGGCUUACUGAA	1	no match			
10267	ACAAAAACGAUUAAGGAGACAACAG	1	0	transposon	LINE	G2
10268	CCUGAGAGAUGGGCGAGCGCCGUUC	1	no match			
10269	CAGAUAUGACCAAAGCAAAGCAC	1	2	Su(Ste)		Su(Ste)
10270	GCAGGGUACCAGUCCCGGAAUG	1	no match			
10271	UUCGGGGCUGGUCCGGGUUGAAGA	1	no match			
10272	AUUCACAUCAAAGACCACAUGCAC	1	0	transposon	LTR	Stalker2
10273	CGCGGGCGGCCCCGGGUUCGUUUCCC	1	no match			
10274	AAAAACCUGAAGAUGAUCCCUGAGUC	1	no match			
10275	AAAUUCUCAACCUGGACGCUUCUUA	1	1	transposon	LTR	3S18
10276	GUAACAACGAAGAUGAUUCUGGCCCC	1	1	transposon	LTR	invader4
10277	ACGUACUAAAACAAAAACCUCCG	1	0	transposon	LINE	Rt1b
10278	GUUCCCCGGAUAGUUUAGUUACGU	1	0	ncRNA		rRNA
10279	AAAAGUGUCCAACAUGGCGAGUGA	1	2	transposon	LTR	opus
10280	UAGCAUUGCGAUGGUCAGAAAGUGAU	1	no match			
10281	GCUAAUACCGGAUGAUGACGUA	1	no match			
10282	AUGGCUUUCUCAGUCGCUUCCCGGGA	1	1	ncRNA		rRNA
10283	UACGUAAACAACCCCAGACUGACUU	1	2	transposon	LTR	Max
10284	GCAAAAAUAACUCCGAAUUAUGGCA	1	0	transposon	LINE	Cr1a
10285	GGAACGCCCGUCAUGAUCAUCGAUGA	1	0	coding gene		CG32022-RA|CG32022
10286	AUACAAACAAAUAAAGAGUUUCCGA	1	0	no annotation		
10287	CAAACCAAUCCCAACCAAACCCGUCAAU	1	1	transposon	LINE	G2
10288	ACGAGAAGACCAGUUCGCUGGAGGCGGAG	1	2	coding gene		CG10542-RA|Bre1
10289	CCCAGAUAUAAACGCUACGCUAGCCA	1	0	transposon	LTR	opus

10290	CGCCCCUGGACGACCCACGUACCUA	1	no match			
10291	CAACCAGAGUUCCCAGUGCUUGG	1	0	transposon	LTR	DM88
10292	UGACAUCGAAUGGAGUGCCAGAACCCGU	1	0	transposon	LINE	R1-element
10293	GGACGAGGCGCCGCCGCCCCCCCCAC	1	no match			
10294	GAGACAAACAUGUUCGAGCUACAGC	1	0	transposon	Unknown	RR50960
10295	UGCUAGAAGACGAUUUUCGGGCCG	1	0	transposon	LINE	Cr1a
10296	UGUGCUAGAAGACGAUUUACGGGCCGA	1	0	transposon	DNA	1360
10297	GCUUAAUUCCCUGGGAAGAUGCCCUGUU	1	0	coding gene		CG17991-RA|CG17991
10298	CAGCGUUUGACGUAGCAGCCCGG	1	0	transposon	LINE	R1-element
10299	GCAAAAGUUGACCGAACUUGAUUAUUU	1	0	ncRNA		rRNA
10300	AGAACGAUGACGAUCCCACGGCUGU	1	1	transposon	LTR	ninja-Dsim-like
10301	AAGAACAACAGUGCAGACGGCGCCCGA	1	1	transposon	DNA	H
10302	CCAGACGUAAUACUCCACUACCCGGAUA	1	2	transposon	LTR	HMS-Beagle
10303	GUGGCUUUGAGAGGCGCCGUAGUUUC	1	1	no annotation		
10304	ACUCGGCUUGGUCCAGUCCUGGGAAUG	1	1	coding gene		CG17673-RA|Acp70A
10305	CAAGCCAAUCUGGACGUACGGAAUUCA	1	0	transposon	LINE	G2
10306	ACCCUGAACGCGCCCGAUCUCAUC	1	no match			
10307	ACCGACGAAAUUUGCGUUCCAGCGAGA	1	0	transposon	LTR	3S18
10308	GCGUAAGCCUGCAUGGAGCUGCC	1	0	ncRNA		rRNA
10309	AGAACCUGAAACCGUAUGCGUUAC	1	no match			
10310	GCUAAGUGGGAAAGGAUGUGGA	1	no match			
10311	AUAAUAAUAAAUAAAAGGGUGACCGGC	1	no match			
10312	CAAAGCUUUUGGGCUCCGGGGGA	1	0	ncRNA		rRNA
10313	GCCGAAAUGACCGAUAGACGAAUGGCC	1	0	no annotation		
10314	AUUGCACUCCGGAUGUGCUGACCCCU	1	no match			
10315	AAGAUGGCAAAGCUCUGCUGAAGAC	1	0	coding gene		CG2238-RA|Ef2b
10316	GCCGAACUUGAGACCGCCCCUGGACA	1	0	Su(Ste)		Su(Ste)
10317	UGUUUUCAUCGUUUAGACGGCUCGGGC	1	no match			
10318	AACCCUGAGGCGUGAUGACGAGGCACU	1	no match			
10319	CAACCAGAGAUCUAGAGGGCCCAUU	1	2	no annotation		
10320	UCUCUUAUCAUCCGAGCGAUACGCCUAC	1	no match			
10321	GGGGAAAACAACGCAGUGGCAGACGC	1	0	transposon	LINE	Doc4
10322	GUGAAUGAUAUGCACCGGAAGUAUA	1	0	Su(Ste)		Su(Ste)
10323	ACGUCUGUGAUGCCCUUAGAUGU	1	1	ncRNA		rRNA
10324	CGAGUAAUAUGAGCGCCCAGAGUGAAAA	1	1	transposon	LTR	opus
10325	GCGCUAUUGAUUCAUCCAGGGAA	1	0	transposon	LTR	ninja-Dsim-like
10326	GUGAAUAAUAAAUAGCCGAGGAAAGU	1	0	transposon	LTR	diver2
10327	AAAAACCUUAAGAUGAUCCUGAGUC	1	2	transposon	LINE	Helena
10328	CCCCUAUCCAAGAUACAACACACC	1	1	transposon	LINE	F
10329	AUAGUCAUUUUGAGAGUUCGCUAU	1	0	transposon	LINE	Cr1a
10330	CCAGUCAGUAGAGCACUACCGCCGGCC	1	0	repeat		trf
10331	UGCAAACCCAAAAAGACAAUACACGAC	1	1	repeat		trf
10332	CAGGGGGAUCUUUGAGUGAUGAAAUU	1	no match			
10333	CAACUGGAGAGUUUGAUCCUGGCUCA	1	no match			
10334	UGUUUCAUCGUUUAGACGGCUCGGGC	1	no match			
10335	GCCUGAUGACCAUAGCAAGUUGGUACC	1	no match			
10336	UGCAUUUCAAUCGGAAGAGUUCUCUGC	1	0	transposon	LINE	Circe
10337	GACUAGCUGGCCGAGUAUUUGAUAG	1	1	Su(Ste)		Su(Ste)
10338	AACUGAAUAAACGAAAUGGAUGACC	1	1	transposon	LTR	invader4
10339	AAUACAUGCCGACGGGCGCUGACC	1	no match			
10340	GGCGAGACGGGUGACAGCCCCGUACGG	1	no match			
10341	CGACUCAGUAUCCCGUUUAUUUGGGG	1	0	transposon	LINE	Rt1b
10342	ACAACAAACAACAAAUACCCGCUG	1	1	transposon	LTR	GATE
10343	CGCAGAACCUACCAUCCCUUGACA	1	no match			
10344	AUGCUCUCGAAUGUAUGCCCGAUAUU	1	1	transposon	LINE	R1-element
10345	CGACAGAAUAGAGAGGGACCUCAA	1	no match			
10346	CAGAUACCCGUUACUCAGCUAGU	1	0	transposon	LINE	INE-1
10347	UCGGACAAUUAUUUUUUUAUGUUGUGGC	1	no match			
10348	CUCAUUCCGAACAUACACCGACA	1	0	transposon	LINE	Doc
10349	UUCGAGAAGCGUUUGAUCGAGUGUGC	1	1	transposon	LINE	BS3
10350	CCGAAGAACAAUUUUACCUACAAGACC	1	0	transposon	LINE	G6
10351	GACCACAUGGAAACCAAGGUCCCU	1	no match			
10352	GAAGACGUUCAUUUCGGACAUGGGU	1	0	transposon	LTR	gypsy8
10353	ACCGACCCGACCCCUUUAUGUAAGGCC	1	0	transposon	LTR	ninja-Dsim-like
10354	GUGAUCGUCUAGUGGUUAGGACCCCA	1	0	ncRNA		tRNA
10355	CGAGAUCUUGCUGACGUGCGACC	1	0	coding gene		CG5276-RA|CG5276
10356	UCCUCAACUGUGGAGGGGAUCCUCAA	1	no match			
10357	CGCAAACCCAAAAUGACAAUGCACGAC	1	0	no annotation		
10358	UAAACAGUACGUGAAACUGCUUA	1	0	ncRNA		rRNA
10359	CAGCUCCUGCCGGUAGAACGUGGGCC	1	no match			
10360	CUGUUUCAUCGUUAGACGGCUCGGGCGG	1	0	AT-chX		AT-chX-B
10361	UUAGAUACAAAGCUAGAUUGGUUGCAC	1	0	transposon	LINE	Cr1a
10362	GCCGAAGUCAUUGUGCGCCAGAGU	1	0	transposon	Simple_repeat	I
10363	AACGGACACAUAACGGAAUCGCUA	1	1	transposon	LTR	opus
10364	ACAGACUCGGAACGCCUCAUUGGU	1	0	coding gene		CG31359-RA|Hsp70Bb
10365	GUACCCGACGGCAAGACUCCUUC	1	0	transposon	LINE	G2
10366	GAGGAAACGAAAACCCAUUCCGGCAC	1	no match			
10367	UACGACGAGAACGAGAUCAUCGU	1	2	transposon	LTR	frogger
10368	CACGCGGGCGGCCCGGGUUCGUUUCC	1	1	ncRNA		tRNA
10369	GUCUUUGUAGAGGCGCCGUAGUUACCAC	1	no match			
10370	UGGACCAAGGUGAAGCCUACGCGC	1	0	transposon	LTR	HMS-Beagle
10371	UGCGAGAUGGACCUUUAGAGCAC	1	0	transposon	LTR	GATE
10372	CUCAAUAAGAAUAGACACGAUUUUGA	1	0	transposon	LTR	Idefix

10373	UGACGAGUAAUGGUCUUCGGGGCGU	1	0	coding gene		CG12717-RA|CG12717
10374	UAUUGGCAACCGAAAAUAUAAACGA	1	0	transposon	Unknown	RR50960
10375	GGUUCCCUCUACCUAGACGACGAGA	1	0	coding gene		CG14476-RA|CG14476
10376	CCGCGUCCUGGGGCUGAGCCAGUUUUGU	1	0	coding gene		CG17097-RC|CG17097
10377	AAUUCUAGAGAACGAAAGCGAGGAC	1	0	transposon	LTR	flea
10378	UGAGUAACGCGUAUCUGAUAGUCG	1	0	transposon	LINE	INE-1
10379	AUAGGGAUGAUGAUACCGAUUAAAGG	1	1	no annotation		
10380	GGUUGCGAGAGGUCCCGGGUUC	1	1	ncRNA		tRNA
10381	GAGACCCGUCGCCGCGCUCUCCCCCCU	1	no match			
10382	ACCGAGAGAAGACUUUUCACCCGAUGU	1	no match			
10383	GACACCGGAACAUGCGGCUGCGAGU	1	2	transposon	LTR	diver2
10384	AAGUGGAGCCGUACCUGUUGGUU	1	0	ncRNA		rRNA
10385	AGCGAAUACCUCCGCUUGUCCCAAG	1	0	transposon	Simple_repeat	I
10386	AUUCGUGUAUGAAAUGACCCGUCUAA	1	0	no annotation		
10387	ACCCUUGAUAAACUGUUGACUUUCG	1	1	transposon	LINE	Rt1b
10388	GAGGUCGAGGUGGGUGGAUUGC	1	no match			
10389	UCGCCCCGAACCCUUCUGGUGUGU	1	1	transposon	LINE	baggins
10390	GCGGAAAGAGAGCGAUUUUGGCCGU	1	0	transposon	DNA	1360
10391	UGUGACUGAUCUAGAGUACAAUGU	1	0	transposon	LTR	DM88
10392	CAAUCCCAACCAAACCCGUCAAACCC	1	0	transposon	LINE	G2
10393	AGCUCGAUCAUGUGACUCCGGGUG	1	0	transposon	LINE	Doc2
10394	GGCGGGUGUUGACGCGAUGUGAUUUCU	1	no match			
10395	CUGUAAGGGGAUCCUCAA	1	2	no annotation		
10396	ACAAGCUGUGUACCCUACGGCAUA	1	2	transposon	LINE	G2
10397	UGGAACAAAUUGGCCUGAUUAGCGGA	1	no match			
10398	CAAGCUAGACUAUGUAUCCCCAGG	1	1	transposon	LINE	Doc
10399	GGCGUUUUUCCCGUUGUCAUUAAGG	1	no match			
10400	GAUCCUCACUGUGAGGGGAUCCUCAA	1	no match			
10401	CAGGUGGGGAGUUUGGCUGGGGCAGCA	1	no match			
10402	UGCAUUUUUAAGUUACCUGCAUCGAGG	1	0	transposon	DNA	pogo
10403	CUUGAUCUCGGUGUUCAGUACACACA	1	0	repeat		LSU-rRNA_Dme
10404	CUUUAACAAGGACCAAUUGGAGGGCAA	1	0	ncRNA		rRNA
10405	GAACUCAAAAUUAUUCCUGAUGGCG	1	0	transposon	LTR	flea
10406	CAAGGGUGCACCAUCGACCGAUCCUGA	1	no match			
10407	AGAGAACGCACCCGCUUGACCUUA	1	0	transposon	LINE	G2
10408	UCACCCUGAACGCGCCCGAUCUCGUC	1	no match			
10409	ACCCUGAACACGCCCGAUCUCGU	1	no match			
10410	UCGUUGAGCAGAACAGACAUGUCU	1	0	transposon	LTR	invader1
10411	AUGUUCGAACGAUCUUUUAACCCAGGU	1	0	transposon	LTR	invader4
10412	AUACAACUGAAAAAGACGCUAUAGU	1	0	transposon	LINE	RR49783
10413	ACGAGUAAUAUGAGCGCCCAGAGUGUA	1	2	transposon	LTR	opus
10414	AAAACAACUAACACCUACCUGAGAUG	1	1	transposon	LINE	TART
10415	UUAGAAGACACAUCAUACCCAGGAAG	1	0	transposon	LTR	Idefix
10416	GAUGGAUACGAGUUCUUUGCCAACCGAC	1	0	coding gene		CG10930-RA|PpY-55A
10417	GUCAAGGAUGCCAUACAGACGGUUGGGA	1	no match			
10418	AAAACCUGAAGAUGAUCCUGAGUCGCAC	1	1	transposon	LINE	Helena
10419	CAUCUUCUUAACGCAGCUGCCGAAA	1	1	Su(Ste)		Su(Ste)
10420	CUGAGAGGACGACCAGCCACACUGGGA	1	no match			
10421	ACCACCCUGACGCGCCCGAUCUCGU	1	no match			
10422	AAAAGUUUACCGGAACGGUCUGGAAUGGCC	1	no match			
10423	GCCUGCUGACGGGGCCCGAAAACACGA	1	0	transposon	LTR	opus
10424	UAAGCUCUCGAAUGUAUGCCCGAUA	1	1	transposon	LINE	R1-element
10425	UAGUCGGUCCUACGUUCAAGGCGAA	1	1	ncRNA		rRNA
10426	UCCUCGAUAGUAUAGUGGUUAGUAUCC	1	0	ncRNA		tRNA
10427	CUAGGACUACCAUGGUUGCAACGGGU	1	0	ncRNA		rRNA
10428	UACGAUACGGAAGAUGGCCCAAAAA	1	0	transposon	LINE	F
10429	UGUUUCAUCGUUAGACGGUCUCGGGCG	1	no match			
10430	UCCUCGGAGGCGUGGGUUCGUAUCCC	1	1	ncRNA		tRNA
10431	UUCCAAAGUGUAAAGGCCCAUAUGU	1	0	transposon	LTR	flea
10432	UUAAACGAGUGUUAUUGUGGGCCGGUA	1	0	ncRNA		rRNA
10433	CACGUACAACGGGGCUUGAAGC	1	0	transposon	LINE	Doc4
10434	ACCGAAUUAAUCGCGAGCUGCAAGACA	1	no match			
10435	UGCCAAAAGAUAAAACGUUUACGAAA	1	no match			
10436	AUAUACGUUCGUCGAUGCCCGUGGCA	1	0	coding gene		CG7713-RA|CG7713
10437	UGCACAUUAAGUUUUGAUCGCCGAAA	1	0	transposon	LTR	invader1
10438	GCGGAAUCAGCGGGGAAAGAAGA	1	no match			
10439	CACUGGAUGGUUGCGGGGGUCGCG	1	no match			
10440	CAGGCAAGGUGUUUAAGUGGUUGAGC	1	no match			
10441	GGUUUUGUAAACCAGGGGUCGUGA	1	1	ncRNA		tRNA
10442	ACCACCCUGAACGCGCCCGAUCUACGUC	1	no match			
10443	CACCCGAAGGUUCUUAUCGUGCCA	1	0	coding gene		CG7728-RA|CG7728
10444	UUUGAAUUGAUCGACGUACUCUUUG	1	0	transposon	LTR	Dm88
10445	GGCGAGCGAAAAGAAAACAGUUCAGCA	1	0	ncRNA		rRNA
10446	ACCAACGGGAUGCCUCAUACUCGAG	1	1	transposon	LTR	micropia
10447	AGAACUAUUAGAGUACGAUUUCGAGG	1	0	transposon	LTR	diver2
10448	UAGGAAACUAAAUCAUAAGACAACCGA	1	0	transposon	LTR	gypsy4
10449	AAGGAAAUGCUCGCGAUAAUUUGGUC	1	0	transposon	LTR	opus
10450	CAGGACCAUAUCCGAAGACGCGGCG	1	0	transposon	LINE	Rt1b
10451	UGCCGAUGACAACGCCUUCCUAGCCUCC	1	1	transposon	LINE	G
10452	CCCAAUAAAAGCCACGGCACUCCU	1	0	transposon	LINE	Doc
10453	UAGACUAUGAAGAGACAUUUGCUC	1	0	transposon	LINE	Cr1a
10454	AAUCUCAACUGUGAGGGGAUCCUCAA	1	no match			
10455	UACGGGAGGCAGCAGUGGGGAA	1	no match			

10456	GGUGCGCAAUUCCGUGGGCGAGGA	1	0	coding gene		CG9285-RA|Dip-B
10457	AAGGCUUUUCCCGUAGUUGAUUUUG	1	no match			
10458	GGCGGCGACGUCGGCUACCCACCCGAC	1	no match			
10459	CAACCAUCCCAACCAAACCCGUCAA	1	no match			
10460	UACACACCGCCCGUCGCUACUACCGAU	1	0	ncRNA		rRNA
10461	UCAAGACCGAAAAGAACGAUGGCC	1	0	coding gene		CG4049-RA|CG4049
10462	AAACAGACUUUGAUCCGGAGAUGUCU	1	no match			
10463	CGAGGCCAAGUUUGCGUUGAAGAA	1	0	coding gene		CG4599-RA|Tpr2
10464	ACCUAUAAUAGACGGUGACUUAACGAU	1	0	transposon	LTR	gypsy12
10465	AAAACCUGAAGAUGAUCCUGAGUG	1	1	transposon	LINE	Helena
10466	CGAGCGGAAGGACAUCAUGGACACC	1	0	coding gene		CG9130-RA|CG9130
10467	UUUGUCAAUAAACAAGAUUCCCGGA	1	1	transposon	LTR	micropia
10468	CUGACAAACAAGUAGGCAAUGAAA	1	0	transposon	Simple_repeat	I
10469	GCCCGAAUGAUAACCAAGUUUCUAA	1	1	transposon	DNA	BS
10470	CCGAAACACACUCCAUUCCAGCGAUA	1	2	transposon	LTR	GATE
10471	UCACCAAGCGUUGGAUUGUUCACCCG	1	1	ncRNA		rRNA
10472	GGCGGCGUCCGGUGAGUUCUCGCUGGCCCU	1	no match			
10473	AAAACCUUCGUAGACGACCUGCUUC	1	no match			
10474	GGAUCCUCAACUUGAGGGGAUCCUCUAA	1	no match			
10475	CAAGAUAUGACCGAAACAAAGACAC	1	1	transposon	DNA	1360
10476	UCCGAUUUCAAGUACAAGACGAACC	1	0	repeat		RSP
10477	CCAAUGAAAACUACACCCAGACGGA	1	no match			
10478	GACAAUCGAACGAAGUCAUAGAACAUA	1	1	transposon	LTR	gypsy8
10479	GAAGGCGUGAAACCGUUAAGAGGU	1	no match			
10480	GCGAUCCACUGAAGCUUAUCCUUUGCU	1	0	ncRNA		rRNA
10481	UACCAAUCACGCCGUUUGAACCCU	1	0	transposon	LINE	X
10482	ACAACUCAAGUCCCACCUUCCCA	1	2	transposon	LINE	G2
10483	UUGUUCCCCGGAUAGUUUAGUUACGUA	1	0	ncRNA		rRNA
10484	AUCCUCGAGAAUGGAAAAAACCCCAA	1	0	transposon	LTR	Idefix
10485	AGAGAUAGAAAAAGGCGACAACGA	1	0	transposon	LTR	Idefix
10486	ACAAACACAGUUCCCGAAUGAGA	1	1	transposon	LINE	Cr1a
10487	UUCCCAACCAACUAAAACCCGUAGA	1	0	coding gene		CG10377-RA|Hrb27C
10488	UGGAGAGUUAUGCGAUGAGUGGA	1	1	transposon	LTR	GATE
10489	UUGGGCUCCGGGGGAAGUAUGGU	1	0	ncRNA		rRNA
10490	UCUCAUCGUCGUAGACAAGCCCG	1	no match			
10491	UACAAUGAUCUGAUCGGGUUUUU	1	0	transposon	LINE	Cr1a
10492	CGACUCGUUUUAAAUUUCGCAACAAGC	1	no match			
10493	CCAUCCUAAAACUGAUACACAGCC	1	0	Su(Ste)		Su(Ste)
10494	AGUACUUCAGCCAGACACUGGACGU	1	0	Su(Ste)		Su(Ste)
10495	CGAACAACGAGACCACGUGUUGCAUU	1	0	transposon	LTR	HMS-Beagle
10496	CAAAACCAGAACCAACGGACGUGGAC	1	0	transposon	LTR	HMS-Beagle
10497	GCGAGAUAGAGAUAUUCCCGUUAA	1	0	transposon	LTR	invader1
10498	UCGAGAUGGAGAGGCUUCAGAGU	1	0	transposon	LTR	ninja-Dsim-like
10499	GGCACUGGGGAGUUCUCUUCCACCUCGA	1	0	no annotation		
10500	AGCCCGUGGACGGUGUCGAGGCCGGUGGC	1	no match			
10501	GUCUCGACGAAUACCGAUCCAGUGG	1	no match			
10502	AAGCACAAGUACGAGUUCAUCCUGGAU	1	1	coding gene		CG12261-RA|mRpS22
10503	UCCGAUCAGACGAGAUGGCCAACAG	1	0	transposon	LINE	Rt1b
10504	UGAAGUAAAAGAACCAACCACGGGA	1	0	transposon	LTR	diver2
10505	GACUACAUAUGGUUGAGGGUUGUA	1	0	ncRNA		rRNA
10506	GCGGAAUCAGCGGGGAAAGAAGAC	1	no match			
10507	AAAAACCUGAAGAUGAUCCGGAGUCGC	1	2	transposon	LINE	Helena
10508	CGACUAGGGAUCGGGUGGUGUUUUUUU	1	no match			
10509	AAGUCUUCCGUCGCAUCUCGCAGCU	1	0	coding gene		CG33715-RB|Msp-300
10510	GAACGUACCGGCAAUCUGACUGUGG	1	1	coding gene		CG8430-RA|Got1
10511	AUAAUCUGAAGGUCGAGAGUUCG	1	2	ncRNA		tRNA
10512	AUAAGCAAAAUGCAGAACACCGG	1	0	transposon	LINE	Doc4
10513	CAACGCCUCGAAAGCGAUUGCUGCAGAU	1	0	coding gene		CG2841-RA|ptr
10514	ACCCUGAACGCGCCCGAGUCUCGUC	1	no match			
10515	GGAUCUCAACUGUGAGGGUAUCCUCAA	1	no match			
10516	CAAAAAACCCAGUGCUCGAAGCAGCGG	1	0	transposon	LINE	G6
10517	CAACCGAGAUUCCCUUAGUAGUGGC	1	no match			
10518	AUGGCUCUCUGUCGCUUCCCGGGA	1	no match			
10519	GAGAGGCGCCGUAGUUACCACCCCACC	1	0	no annotation		
10520	UCUCAUCGUCGUAGAGCAAGCCCGA	1	1	Su(Ste)		Su(Ste)
10521	AUGAGAUAUGCUAGCAGACUGGAGAU	1	1	transposon	LINE	G2
10522	UAGACCAGCGACGACACCUUUGGA	1	0	transposon	LTR	297
10523	GCCAAUGAAAACCACACCCAAACCAGA	1	0	transposon	LTR	McClintock
10524	GACCUAGAUAUACGAUUCAUUAAGGC	1	0	transposon	LINE	Circe
10525	UAGAUCUUCACCGUUGACUUGGCC	1	0	Su(Ste)		Su(Ste)
10526	AUCUGUGGGAUUAUGACUGAACGCCU	1	no match			
10527	AGAAGACAUACCUGAGCGAACAGACC	1	2	transposon	LTR	McClintock
10528	GAUCGCCCAUCUCAAGUCUCUGCUGC	1	0	coding gene		CG6030-RA|ATPsyn-d
10529	AGACGACAUAAUUGUCUUUCGACAUCC	1	no match			
10530	UUCCGGAGUGUUGAUCUGAGCCUC	1	0	transposon	LTR	GATE
10531	GGGGAGUCCGAAGUGGUCUUAGUACAA	1	no match			
10532	CUUGAACAUAAACGAAGACCCACAAA	1	1	transposon	LTR	INE-1
10533	CCGGUGGAGUUCUUAUAUGUGAU	1	0	ncRNA		rRNA
10534	UGCCUGGGCCUCAGCAAUGUGUGGG	1	no match			
10535	CCAGGUCCGAACAUAAGUGUGUAAGA	1	0	ncRNA		rRNA
10536	GGGCUCCGGGGGGAAGUAUGGUUGC	1	no match			
10537	CUGUCUACGAAAACAAUCGCAGGGUU	1	no match			
10538	GGAUCCUCAACUUAGGGGAUCCUCAA	1	no match			

10539	CCCCUGGACGACCCACGUCCCUAUGCU	1	no match			
10540	ACAUCCCAGGGGGUAGCGACUGUUU	1	no match			
10541	ACAUUCAUAACCGAGCGACUGUUCCA	1	0	transposon	LTR	GATE
10542	UUCUUACAUGUGGUAUUUACGAAGU	1	0	no annotation		
10543	CGCUGAUCUCGGACAAGUUUGGUGC	1	0	coding gene		CG6186-RA|Tsf1
10544	UUCAAGGUUGCCCCAGGAUGCUGUGCCAU	1	1	repeat		trf
10545	UGAAAUCCGAGCUAAUCCGAGCACCA	1	no match			
10546	CUGCAUCCGAUUGGGCCUUCCGA	1	0	transposon	DNA	H
10547	UUGUGAAGCAAUUCACCAAGCGUUG	1	no match			
10548	UACGACGAGAACGAGAUCAGCGCGAUU	1	0	transposon	LTR	frogger
10549	CAUCAAGCCUGAGAGGUGAUGCGUAGUCC	1	no match			
10550	GUGAUGUCAACGUUUGAUCCCCUAGGA	1	1	transposon	LTR	ninja-Dsim-like
10551	UAACGCGUGAAGUUGACCGCCUGGGG	1	no match			
10552	CGGACAGAGACGGCGCAUUUUCCAG	1	0	transposon	LTR	McClintock
10553	CGCAGCGCUCGAGAGGUGUCCG	1	0	coding gene		CG6831-RA|rhea
10554	CAGUCCGAAACACACCCCCGAUAAU	1	1	no annotation		
10555	CGCAGAGUACGGGGCUUACUGAAG	1	0	transposon	LINE	baggins
10556	UGGCUCCGGUGCGCUUGUGACGGCCCG	1	no match			
10557	CAGAUACCAUCUUCCUACGAACCGA	1	0	transposon	LTR	297
10558	ACCCUGAACGCGCCGGAUCUCGU	1	no match			
10559	AACUCCAGCAGAGACCGUACCCGA	1	no match			
10560	GGCUCUCUCAGUCGCUUCCCGGAAAC	1	0	ncRNA		rRNA
10561	AAAGGAAUUGACGGAAGGGCACCA	1	0	ncRNA		rRNA
10562	GGACGAGGCGCCGCCGCCCCCCCCCAC	1	no match			
10563	GGCGAUGAAAAGAAGUAGUACGGCA	1	1	repeat		trf
10564	GCGCGGGACCCUUCACAAUUGCUUCG	1	0	transposon	LINE	TART
10565	UGAAGUGGGGAAGGGUUCCAUGGUAAC	1	no match			
10566	AUAGAUGACUUGAGUUCGUGUACAUUA	1	0	ncRNA		snoRNA
10567	GACACAAUCGAUGACGAUCGUCUGCGC	1	0	coding gene		CG5119-RA|pAbp
10568	GAGAGUGAUGCCGAGUAUAAA	1	0	repeat		trf
10569	CACCACAAGGGAACAUUGAGUCCGGU	1	0	transposon	LINE	Cr1a
10570	GCAAAGUGAACACAUCGCGAAGCA	1	2	coding gene		CG18743-RA|Hsp70Ab
10571	UACGAAUUCAACGAGGUAUGCCAU	1	0	repeat		SAR_DM
10572	UCCGAGGCCCUGUAAUUGGAAUGAGUA	1	0	ncRNA		rRNA
10573	AUGCGAGAGGCCCUGGGUUCAUUUCCCA	1	no match			
10574	CUGCCCUUUGAUUGCAUUCCGAAUGA	1	0	coding gene		CG2995-RA|G9a
10575	UGGAGCACUCGAAGAAGACCCACGCCC	1	0	coding gene		CG31546-RA|CG31546
10576	CGACUCAGUAUCCCGUUUAUUUG	1	0	transposon	LINE	Rt1b
10577	UCCGAUUUCAAGUACCAGGCGAACAGA	1	0	repeat		RSP
10578	GCGGUGGGCUGAAUCCUUUGCAGACGA	1	no match			
10579	CCGACAAACAUCCAACAAACACAACA	1	0	transposon	LINE	Doc
10580	AAGUACGAAAUUUCCAUUCCUUUUAA	1	0	transposon	LTR	GATE
10581	AAAUUCCGGCGCCAGGAUGUGGAUGA	1	no match			
10582	AUGAAAACGAAGGUGACCUUUUUGGG	1	0	transposon	LTR	flea
10583	CUACUUAGUUCCCCGGACUAGUUUAGUUACG	1	no match			
10584	GGAUGGGAGACCGCCUGGGAAUACCG	1	no match			
10585	CACGAAUGGCGUAACGAUCUCCCCGCU	1	no match			
10586	GCCGAUCACAGGUCCGCACCUACGC	1	0	transposon	LINE	R1-element
10587	AAAAUACAAAGACGACGGCCCACC	1	1	transposon	LTR	springer
10588	CUGGGAAGGUGGGACUUGAGUUU	1	0	transposon	LINE	G2
10589	GUUCGAUUCCGGAGAGGGAGCCUGA	1	0	ncRNA		rRNA
10590	CUUCGAUUCCGUAGAAGACACAUUCGA	1	no match			
10591	AAAAAACCUGAAGAUGAUCCUGAGUCG	1	2	transposon	LINE	Helena
10592	GUGAACUACAUCCCAACGACCGAGAUGC	1	0	no annotation		
10593	GAAGAAAGGAGAUGCGUUGCCAAGAGU	1	0	transposon	LTR	diver2
10594	UCGGAUCCUACGCCGCGACGACGCCGA	1	0	transposon	LTR	ninja-Dsim-like
10595	UACAUACGAAAAGCGAGUUGACGUGU	1	no match			
10596	GUCAAGUGCGGCGAUGUGGUGACCCUGC	1	0	coding gene		CG4750-RA|loopin-1
10597	GAAUAGAGAGUUAAACAGUACGUGAAA	1	0	ncRNA		rRNA
10598	CUAACAUAUCUGCGAGUAUUUGGG	1	no match			
10599	GACAAGCUGACCCGCAUCACGGGACAAA	1	no match			
10600	UAUAUUUUGAUGGAGUACGGAGUGU	1	0	transposon	LTR	gypsy12
10601	AUAAGUGGGAGAUAUUAGACCUCGGU	1	0	repeat		LSU-rRNA_Dme
10602	ACUACAUUCAGGACCCGUUCAACCGGA	1	0	coding gene		CG40635-RA|CG40635
10603	UACGAAGGGGGCUAGCGUUGCUCGGA	1	no match			
10604	CCGUCUCCGCCCCCCGGCCCCGCG	1	no match			
10605	AAGGAGCACCAUCGACCGGCCCU	1	no match			
10606	AACCGAAUUAAUCGCGAGGCUGCAA	1	0	transposon	LTR	mdg3
10607	AACCAAGAUUAUCAACCCGUGUUG	1	2	transposon	LINE	Cr1a
10608	CAAACUGAAUAAACGAAAUGGAUGACA	1	0	transposon	LTR	invader4
10609	GGGAUGAAAGGAUCCUCAA	1	2	no annotation		
10610	CUUGAGACCGCCCCUGGACGACCCACGA	1	0	no annotation		
10611	AACUGGAACGUACCUUGAGCAUAU	1	0	ncRNA		rRNA
10612	GCCGGGGCGGGUUCCUGAGGGGGUCCU	1	no match			
10613	GGAAUAAGAACACGUUGACGACGAA	1	0	transposon	LTR	HMS-Beagle
10614	UUUUCAAGAAAAGAUGCCCGAACGAC	1	0	transposon	LINE	RR49783
10615	ACCAUCACGAUUGGCGAAGACCAAGC	1	0	transposon	LTR	G4
10616	UACUGACGAAUUUAGUUACAAGCAGA	1	0	no annotation		
10617	CAUAGAGCAAAAGGGCAAAUGCUGA	1	0	ncRNA		rRNA
10618	GUUAUGGUUCCUUAGAUCGUUAAC	1	0	ncRNA		rRNA
10619	ACCACCCUGAACGCGCCCGAUCUCGC	1	no match			
10620	AGAGACUAAAUCAACAUGGCUCAC	1	0	no annotation		
10621	AAACUGAUAGACAGAAGGACGGAG	1	1	Su(Ste)		Su(Ste)

10622	UCGUGUUGGACGAGAUUACAUUAAGA	1	0	ncRNA		Hsromega-RC
10623	GUUGCAAUUAAACAGAUUCCGCAGUU	1	1	Su(Ste)		Su(Ste)
10624	CAGAUACCCGUUACUCAGCUAGUAUAA	1	0	transposon	LINE	INE-1
10625	ACCAAGAUCAUCAAACCCGUGUUAU	1	0	transposon	LINE	Cr1a
10626	GCGACGACAACAAAGAAGACUGACAC	1	1	coding gene		CG31224-RA|CG31224
10627	GGCAAAACACAUCCCGGGAAAUGAGC	1	no match			
10628	CAAACCAAUCCCAACCAAACCCGUUU	1	2	transposon	LINE	G2
10629	UACACACUCAUCAACCAAAGGACGUUA	1	2	no annotation		
10630	CCCAGCUCGGGGAGGUAGUGACGGAA	1	1	ncRNA		rRNA
10631	GCACACAGAGGAGAGCAGAUGAAGCA	1	no match			
10632	UUUCACGAAACUGGAACGGGAUUAU	1	0	no annotation		
10633	GGGUUCCGGGGGGAUAUGGUUGC	1	no match			
10634	CACUGAAAUCCAAAAAGACAUGCCCGC	1	1	coding gene		CG14967-RA|CG14967
10635	UUUGCAAAAAACGGAGCGAUAGAGC	1	no match			
10636	AACAUUAACAUCCGCUGCCGAAUC	1	0	transposon	LTR	Quasimodo
10637	CAGGAAUCAACUGCAUGACUCUCGC	1	0	transposon	LTR	Quasimodo
10638	GACGUUAAUGAAGAUGCCGUGGUGU	1	1	coding gene		pseudogene
10639	GGGUACGGCCAUCCAGUCGAAGCAGCC	1	0	coding gene		CG6773-RA|sec13
10640	GCCCCCAUCGUCUAGAGGCCUAGGACAUCU	1	no match			
10641	AUCGAAUGGAGUGCCAGAACCCGUUAU	1	2	transposon	LINE	R1-element
10642	GCUUUAAUCACGAGCAAUGCAGU	1	1	no annotation		
10643	UGCUCUCGAAUGUAUGCCCGAUAUAAC	1	2	transposon	LINE	R1-element
10644	AAAUCCUUGAUAUGACGGACGAG	1	0	no annotation		
10645	UCAAGCCAAAGAGGGAGCCCGGA	1	0	transposon	LTR	flea
10646	CAUCUUCUUAACGCAGCUGCCGAACUU	1	0	Su(Ste)		Su(Ste)
10647	AUGAGAUAUGCUAGCAGACUGGAG	1	0	transposon	LINE	G2
10648	ACUAACGUACAAAUCGAGCAACUCGUC	1	0	transposon	LINE	F
10649	GCAUUCAAUACAGCGAGAUGGCCCAAC	1	0	transposon	LINE	Rt1a
10650	AAGGAGUCUAAUACGUGCGCGAGUC	1	no match			
10651	CCGGGGCGUUCUAUAUAGUUAUGUAU	1	0	ncRNA		rRNA
10652	UGGCUCUCCCAGUCGCUUCCCGGGA	1	1	ncRNA		rRNA
10653	CCAGACGUGAUACUCCACUACCCAG	1	0	transposon	LTR	HMS-Beagle2265
10654	UCAUUUCCCGGGAUGUUGUUGGGUCC	1	0	transposon	LINE	baggins
10655	UCUGGAGAGCUUCUUCACUGCCACACG	1	no match			
10656	ACCACCUUGAACGCGCCCGAUCUCUUC	1	no match			
10657	ACUCGCCGAAUUCCCGGGGCCGAGGGAGC	1	no match			
10658	CCGUCACAGAGGGCAACCUACGGGGCA	1	0	transposon	LINE	R1-element
10659	GAGACCCGUCGCCGCGCUCUCCCCCC	1	no match			
10660	AAAACCUGAAGACGAUCCUGAGUAU	1	no match			
10661	CGUUGCGUCGCGGUGCAUGGAGCA	1	no match			
10662	CUGAGAAGACGGUCGAACUUGACU	1	no match			
10663	UUCGAGAACAACGAUACUGUUCGGC	1	no match			
10664	AAAACCUGAAGAUGAUCCUGGGUCGC	1	1	transposon	LINE	Helena
10665	ACCAAAAUCAUCAAACCCGUGUUA	1	0	transposon	LINE	Cr1a
10666	GUUGAAAACUUUAACCAAUACCCCGC	1	0	ncRNA		snRNA
10667	UGCAAUCUCACCCCCCACCUCCCCG	1	0	transposon	LINE	S
10668	CUGCCUACAAUACAGAUACCCACU	1	1	transposon	LINE	Rt1b
10669	ACCAAUAAAACGAUUAAGACCAGCUAAA	1	0	transposon	LTR	McClintock
10670	AACCCAAGAUCAUCAACCCGUGUUAUUC	1	1	transposon	LINE	Cr1a
10671	GCCAAUUGCAUUAGACGAGACGGCCC	1	1	transposon	LTR	opus
10672	AGAACUUUAAAUUAGCCGACCCCGAUU	1	0	transposon	DNA	HB
10673	CGCCCCUGGACAACCCACUGCCUUGA	1	no match			
10674	AAAAGAGGAGAAGGUGGCUUUGAGA	1	1	no annotation		
10675	CGGGGCGUUCCUUCGGGACCGAGUCACA	1	no match			
10676	CCAAAGCUUUUGGGCUCCGGGGGAAGU	1	0	ncRNA		rRNA
10677	AAUCGCUCGAUGUCGUGAGGUGAC	1	0	no annotation		
10678	CGAGAUGUUCAACUACACGGACUCGCC	1	1	transposon	LTR	3S18
10679	AUGGGGCGAAGCUACCAUCUGUGGGAUU	1	no match			
10680	UUGUACACACCGCCCGUCGCUACUAC	1	0	ncRNA		rRNA
10681	UCCUGAAAGUAGUUGGUGGGCACGUG	1	0	Su(Ste)		Su(Ste)
10682	AUGCGAAAUUGGCCCAAUCUGA	1	no match			
10683	CAGUCGCGAAAUAGGUAUCCCCGAGU	1	2	transposon	LTR	diver2
10684	GAAAGCACCGUUUCCCGUCCGAUC	1	no match			
10685	AAGUGACAGACCGAAUCGCAAUUCC	1	0	transposon	LTR	Quasimodo
10686	CGUUGCCAUCGUUGGGCCCCCGGUUC	1	no match			
10687	AAGGACAAAGCAGGCUGCACGAAA	1	2	transposon	LTR	micropia
10688	GUAUCCUCAACUGUGAGGGGGAUCCUCAA	1	no match			
10689	GGUGAGCAUACGGUUACUGGAGU	1	0	transposon	LINE	R1-element
10690	CAAAACAAAAGUACACUGUGGCGAGCC	1	no match			
10691	UUGAAGAGCAAGAACCGGAGUCGU	1	0	transposon	LTR	blood
10692	CACAGAGUACGGGGCUUACUAAAAAA	1	no match			
10693	AGCAAACCGUGUUCUACCUCGAUGACC	1	0	coding gene		CG6554-RA|Art1
10694	ACCCGUCAAACCCCACCUGAAGAUCC	1	0	transposon	LINE	G2
10695	CUGUAAUUCAAGAGGACACAAAGC	1	0	transposon	Unknown	RR50960
10696	CCAUGGGAGUUGGUCUUACCCGACG	1	no match			
10697	AACUUAACUUGACCGCUCUGACCA	1	no match			
10698	CGACAUAGUACAGCGAACACAGU	1	0	transposon	LINE	Doc3
10699	UUCAGAAUAAUGUGAUUUGUCUUGAGU	1	0	transposon	LINE	Cr1a
10700	ACUGCCAACAGGCCCCGCACGAUAAGA	1	0	no annotation		
10701	GGAUCCUCAACUUGAGGGAUCCUCUAA	1	no match			
10702	CUGGAUAAUAGCCAAUCCCUAAGACGCC	1	0	transposon	LINE	G2
10703	GCGCAUGAAUGGAUUAACGAGAUU	1	0	ncRNA		rRNA
10704	AGAGACGAGAACGCGUAUAAGUGUGCGU	1	0	transposon	LINE	Cr1a

10705	GGCACAAUUACCAGAAGAUGUUGA	1	0	transposon	LTR	roo
10706	GAGAUAUUGAGCGACUUUCCCCGUAGC	1	0	transposon	LTR	3S18
10707	GUGGGUGGUGGUGCAUGGCCGUUCU	1	1	ncRNA		rRNA
10708	UAGAACAACUACACCGAGUGGUCGAG	1	0	transposon	LINE	BS3
10709	GGCGCGGGUAACCCGUUGAACCCCAUUC	1	no match			
10710	AACGAGCGGAGUGAACCUAGUGAAC	1	0	no annotation		
10711	CGAACUUGAGACCGCCCCUGGACGA	1	0	Su(Ste)		Su(Ste)
10712	CAGGCCCACCUGGUUGAACGUGUCCUG	1	1	Su(Ste)		Su(Ste)
10713	AAAGCGAAUGAUUAGAGGCCUUAGGGU	1	0	ncRNA		rRNA
10714	UACAAUGAUCUGAUCGGGUUUUUUCU	1	no match			
10715	UUAACAACGAUGAUGAUUCUGGCCCA	1	0	transposon	LTR	Max
10716	GCGACCUGGAGCGAAUCCCCAAAAGCC	1	no match			
10717	AUCAUCAACGAGCCCACGGCGGCCGUCC	1	no match			
10718	UUUAACCAAAGACCCUAGAAUAACGA	1	2	transposon	LINE	Cr1a
10719	ACGUUGCAAGACAGGGCGGAAUGGC	1	0	no annotation		
10720	GAUAGAGAGAACAUACGAAACACC	1	0	transposon	LTR	flea
10721	GCCGAAUUAAUCGCGAGGCUGCAAGA	1	0	transposon	LTR	mdg3
10722	AAACCAACCGGGAUUGCCUUAGU	1	no match			
10723	AAAAACCUAGAAGAUGAUCCUGAGUCGCA	1	no match			
10724	UCCUGACCGCGAAAACCCGGGUAGCA	1	0	transposon	LTR	GATE
10725	UCCAAUUCAAGGACAUUGCCAGGU	1	0	ncRNA		rRNA
10726	UUCCGGAGAGGGAGCCUGAGAAACGGCUA	1	0	ncRNA		rRNA
10727	CAGCAAACCAAUCCCAACCAAACCCG	1	0	transposon	LINE	G2
10728	UGGUGUCACCAGCCCAAUCCGCGU	1	0	transposon	LTR	DM88
10729	AAAGGCGAGAUAGAUCCACGGAUCGGUC	1	0	transposon	LTR	flea
10730	GGGAGACAGCGACGAAGAGGAUGAGUU	1	1	coding gene		CG10198-RA|Nup98
10731	CGGCGGCAAUGGCACAGCACCCUGGGG	1	1	repeat		trf
10732	CGGCGAUGCGGUGGUGCAGUCCGA	1	no match			
10733	CGACACCAUCCCUACCUCCAGCACCC	1	0	transposon	LINE	G2
10734	GAUCCUCAACUUUGAGGGGAUCCUCAA	1	no match			
10735	UCAGACUAAAAACGUGUUAGUAAA	1	no match			
10736	CUGGAGAAUAACCGAUGGACCAAUGC	1	0	transposon	LTR	diver
10737	AUUCCGAAAAUGCGAGUUCCUCAACC	1	0	transposon	LTR	Rt1c
10738	CAUAGUCAUGCGAGGCGAACCGGGU	1	no match			
10739	UAGCUAAUUAACACAGUCCCGGGGCG	1	1	ncRNA		rRNA
10740	UUCGAUUUCAAGUACCAGGCGAACG	1	1	repeat		RSP
10741	CACAUCAACAGAAAGCGUAUCCAGGCCC	1	0	transposon	LINE	G2
10742	CACACGGCUAGAUCCUGCUAUACGCAAGG	1	0	transposon	LTR	ninja-Dsim-like
10743	GUUCCUCGAGAACAAAGGCAACGAG	1	no match			
10744	UGAGACCGCCCCUGGACGACCCAC	1	0	Su(Ste)		Su(Ste)
10745	UGCGUUAAGAAAAUGUCCGGACAGC	1	0	Su(Ste)		Su(Ste)
10746	GCAUUCGAGACCCAACGGCACGUGAUUA	1	no match			
10747	GGCUCCGGGGGGAAGUAUGGUUGC	1	no match			
10748	AACCCUGACCAACAGCUAAGGCCCC	1	no match			
10749	AUCCUCAAGAACCCUUUGAAUUCCGA	1	0	transposon	LINE	Doc
10750	AUCAAACAGAAAAUGACGACCCAGA	1	0	transposon	LTR	roo
10751	UUGAAAAUCCGGGGGAGAGGGUGUAAA	1	no match			
10752	AGAGCCGUCUAACGAUGAAACAAAGAUC	1	1	no annotation		
10753	GGUGAAGAGUCUCUAGUCGAUA	1	0	ncRNA		rRNA
10754	CCCGUAGAAAACCUUUGGAGUGCUU	1	1	transposon	LTR	HMS-Beagle
10755	AGCCCCGAUAAACGCAUUCAAGAAUC	1	0	transposon	LTR	HMS-Beagle
10756	AAAAGCUCGGCCUAUCGAUCCUUU	1	0	ncRNA		rRNA
10757	CCAGGCCCGUAUAACGUUAAUGA	1	0	ncRNA		rRNA
10758	CCGUUCUUAGUUCGUGGAGUGAUUU	1	0	ncRNA		rRNA
10759	CUGUGAACGAGUGAACUGGCAAC	1	0	Su(Ste)		Su(Ste)
10760	UGGAAUGUUUUCGUCCCACCCAAGC	1	0	transposon	LTR	GATE
10761	CCAUCCUAAAACUGAUAGACAGCUGGAC	1	1	Su(Ste)		Su(Ste)
10762	UCAACCAACAUCUGCACAUUGUG	1	0	transposon	LINE	Circe
10763	AGGGACAUUAUUGUGGAAAAAUGUUAAA	1	no match			
10764	ACGUGAGCUGGGUUUAGACCGUCG	1	0	ncRNA		rRNA
10765	UCCGAACGUACCACGGUGCCCCCAGUGA	1	no match			
10766	UGUAGGUGUACUGACCCAGUCGGA	1	0	transposon	LINE	baggins
10767	UCAAGAAUAACAGGACCAUCCAGAGU	1	no match			
10768	GGGAAACGGGGCGCGGCCGGAAAGG	1	no match			
10769	GAGGUCAAGCCCUAUGUCCUGGACGAUC	1	0	coding gene		CG5735-RA|orb2
10770	GUCAGGAUGGCCGAGCGGUCAAA	1	1	ncRNA		tRNA
10771	UCCGUGGCAGAAAAACGACUACC	1	0	coding gene		CG11621-RA|Pi3K68D
10772	CAGGUCACCAAGGCCGCCGAGAA	1	1	coding gene		CG8280-RA|Ef1alpha48D
10773	ACUGUGCCCAAGUUAUUUCCGGAUA	1	1	transposon	LTR	micropia
10774	GGACCCUCAACUGUGAGGGGGAUCCUCAA	1	no match			
10775	CAUUCAUCAAGCGGUGGAUCCUGCGUUU	1	no match			
10776	CGGGACAUGGUGUCGGACCGACUUGG	1	0	transposon	Simple_repeat	I
10777	CAAACAGUUAAUCUUGUCAUUCGAGA	1	1	transposon	LTR	invader3
10778	GUGCGAACGGGAUCUUGAUCGACUU	1	no match			
10779	UCAGAGACUUUGGUGGCCAACAA	1	0	coding gene		pseudogene
10780	GCCGACGAGAUGCCCAGGGCGUGCA	1	0	transposon	LTR	ninja-Dsim-like
10781	GGGUUCGGAUCCCACCGCUGUCACCA	1	no match			
10782	AUCAAACGCUGAUAAGGCACCGGG	1	0	transposon	LTR	qbert
10783	ACGGGUAACGGGGAAUCAGGGUUCGAU	1	0	ncRNA		rRNA
10784	AACUAUAUAAUGAAAAGGGAAUGCC	1	0	transposon	LTR	micropia
10785	GGAGCACGACUAGAUUGCCUUUAG	1	0	no annotation		
10786	UUAGACCUCGGAGAAUUUAGUAG	1	no match			
10787	AAACCCUCUAAUUCGGCCCGAACCC	1	1	transposon	LINE	baggins

10788	UGGAGCCGUACCUGUUGGUUUGUCCCAU	1	0	ncRNA		rRNA
10789	CAAUGUGAUUUCUGCCCAGUGCUCUGA	1	0	ncRNA		rRNA
10790	AAACCUGAAGAUGAUCCUGAGUCGCAA	1	1	transposon	LINE	Helena
10791	GAUAACCAAGGACGUUCACACGAG	1	0	transposon	LTR	opus
10792	GGAUACUCGAAGACACCACGGUGU	1	1	transposon	LTR	ninja-Dsim-like
10793	CAAAUCAAUCCCAACCAAACCCGUCA	1	1	transposon	LINE	G2
10794	AUCGCAACUGCUACCACUGACUACAA	1	2	transposon	LTR	diver2
10795	AUAGCUAAUUAACACAAUCCCGGGGCGU	1	0	ncRNA		rRNA
10796	UCUUGAAUCACCGAAUUGAAAUUGGA	1	0	no annotation		
10797	UGGACAACUUGGACACCUGGUGGC	1	0	transposon	Unknown	RR50960
10798	UCCCUGAACGCGCCCGAUCUCGU	1	no match			
10799	ACCACCCCUGAACGCGCCCGAUCUCGUC	1	no match			
10800	CACCCCUGAACGCGCCCCGAUCUCGUC	1	no match			
10801	CACAACAGUAAUUUAGAUUUUUACCC	1	0	transposon	LINE	Cr1a
10802	UCCUUCUCUUCGUUGCGCUGGCUCUGUG	1	2	no annotation		
10803	AAUAGGGACGGUCGGGGGGCAUC	1	no match			
10804	AAGAGGGACGGCCGGGGGCAAUUCCGAU	1	no match			
10805	AAACACGAGAGCCGAGCGCUGUUG	1	1	transposon	LTR	opus
10806	UCUGCAGGCGUCUUAGGGAUUGG	1	0	transposon	LINE	G2
10807	AACCGAAUUAAUCGCGAGGCUGCAAG	1	0	transposon	LTR	mdg3
10808	CCGAACUUGAGACCGCCCCCUGGAUG	1	no match			
10809	GCAGUCGUGGCCGAGCGGUUAAGGCU	1	1	ncRNA		tRNA
10810	CGGCGGGGAAGGUGGCUCGGGGGG	1	no match			
10811	AUGUACAAUGCUGCCGAGUUUGCCUAA	1	2	transposon	LINE	G2
10812	GCGAAUGAUAAGUUUCAUGCCUCCGG	1	0	no annotation		
10813	GCGGGCCUCACGAUCCUUCUGA	1	no match			
10814	UGUAAUUGUCCGCGAGCUUGCCC	1	0	transposon	LTR	3S18
10815	CUGGGCCCGAACCCUUCUGGUGUGUU	1	1	transposon	LINE	baggins
10816	UACUAACUUCCCGAUUACUAAACGA	1	0	transposon	LTR	297
10817	ACAACAAACAACACAUACCCACUGG	1	1	transposon	LTR	GATE
10818	CUAAUACAUAUCGAUCAUUACGGCCC	1	0	transposon	LTR	flea
10819	UACUAGGCUGAGCGGAGAACCAGAGUU	1	2	transposon	Low_complexity	F-element
10820	GACCAUAUCCGAAGACGCGGCGCUAG	1	0	transposon	LINE	F
10821	AACCAACAACGACAACGAGACGCAUGA	1	2	coding gene		CG3585-RA|CG3585
10822	GCAAGAAGUACCUCCCGUAGAUGAC	1	0	transposon	LTR	gypsy
10823	GACUAGCUGGCCGAGUAUUUGAUA	1	0	Su(Ste)		Su(Ste)
10824	AACGGGAAAACUUACCAGGUCCGAA	1	1	ncRNA		rRNA
10825	CAACUGUGAGGCGAUCCUCAA	1	no match			
10826	UCGUGUUGCUUGAUAGUGCAGCACUAA	1	0	ncRNA		rRNA
10827	AUACCAGCAAGUAUCAGCACGGGAGAC	1	no match			
10828	GCACACAGAGGAGGCAGAUGAGAGCA	1	no match			
10829	UGUUUCAUCGUUAGACGGCCCGGG	1	1	AT-chX		AT-chX-B
10830	AACUAUCUUCUCAACCCGCUGAUAU	1	0	no annotation		
10831	UACGGGCUCGACAUGUUGCCAGUUCUAC	1	no match			
10832	CAAUUCAAGGACAUUGCUAGGU	1	1	ncRNA		rRNA
10833	UGGGGCGGUACACCUGUCAAACGGU	1	no match			
10834	ACACCGCCCGUCGCUACUACCGAUU	1	0	ncRNA		rRNA
10835	GAAGCCCAAACGUGAACCAGGUUC	1	0	transposon	Unknown	RR50960
10836	AAACAAAAAUAAAAAUCGGGGCAA	1	no match			
10837	ACGUGUAUCAUUUCCUAGCCAUUAUACG	1	0	repeat		LSU-rRNA_Dme
10838	GACCGGCUCCGGGACGGCUGGGAAGGC	1	no match			
10839	AUCGAGAUCAAGAUGUCAAAAGGGG	1	1	transposon	LTR	flea
10840	CGAGCACAACAGAUCUCACAGGUCGGUC	1	1	transposon	LTR	gypsy3
10841	CGGUGCGGAGUGCCCUUCGUCCUGGGA	1	no match			
10842	UCCUUCCCAAGGCGUCUGGAUCCAGU	1	0	transposon	LTR	flea
10843	CAUUGAUCCAGCUGCUGUUGUUGUUG	1	0	no annotation		
10844	AGAAUGGUGUGGCGCCCAGACUGAAGC	1	0	no annotation		
10845	UUGAGAAGAACACCGACCUUUUACAA	1	0	transposon	LTR	Idefix
10846	AAGGAUUGGCUCUAAGGGUCGGGU	1	no match			
10847	CCAGAAGCAGGGAUAACACUUGGAAACA	1	no match			
10848	AACGCUGAUAAAUCCUCCCACACCACC	1	0	transposon	LINE	G2
10849	AUCAACUGUGGAGGGGAUCCUCAA	1	no match			
10850	AAAUAAGACCUCUGUUCUGCUU	1	0	ncRNA		rRNA
10851	AGGAGAGCGUUCUGUAAGCCUGUGA	1	no match			
10852	AAUCACUGAGGCGGAGUUGUACGAUCU	1	1	coding gene		CG13526-RA|CG13526
10853	UAAUAGCAAAGGGAACAUGGAGAA	1	0	transposon	LTR	micropia
10854	CCCCGCGGAAAGACGGAAAGACCCCACG	1	no match			
10855	CAGAUCAACAAAGUACGGUGGGGAA	1	1	no annotation		
10856	GGUUUCAUCGUUAGACGGCUCGGGC	1	1	AT-chX		AT-chX-B
10857	UCCUAUCUUAUCCCGUAGUACAUCAUC	1	0	transposon	LTR	gypsy
10858	ACACCAUUCGUAGACGACCUGCUU	1	no match			
10859	UGCUCUCGUAUGCUUGCCCGAUAU	1	1	transposon	LINE	R1-element
10860	GAAGCCGGUGAUCGAUAGUUCCCC	1	1	repeat		trf
10861	UGUGGCCUUGUUUCGAGACAAAU	1	0	ncRNA		snoRNA
10862	UUUGUUGAUCAGUUUACACAUUAU	1	0	transposon	LINE	Cr1a
10863	GAGAGUGGACGGGAGCGGCGGGGCG	1	no match			
10864	AGCUUCAUCCAAGCGAACUGUGCCCGAA	1	1	transposon	LINE	R1-element
10865	CGAGGCCCUGUAAUUGGAGUGAGUAC	1	1	ncRNA		rRNA
10866	AUCAGAAACCCACUUUUGUCCCAGA	1	0	transposon	LINE	G2
10867	CUGUAACUCGGCUGCAACUCGGGUCAA	1	no match			
10868	CACUAUAGGAGGACAGUCGACCA	1	0	transposon	LINE	baggins
10869	UGCUAGAAGACGAUUUUCGGGC	1	0	transposon	LINE	Cr1a
10870	GUCGUGUUGAUCGGUACUUGCUGUC	1	2	transposon	LTR	invader4

10871	GCCCGUCCCCGCCCCUUGCCUCUCGG	1	no match			
10872	AAGCGUAUUACCGGUGGAGUUCUUAUAU	1	0	ncRNA		rRNA
10873	GAGAAUAGCAGAAUAUCCGAGAUUGAA	1	0	no annotation		
10874	ACUGGAGUCGACGGAAGGAGGAA	1	0	transposon	LINE	R1-element
10875	AAUUCCGUCAACGAAGCCCAAAGU	1	0	transposon	LTR	Rt1c
10876	AUCUCUAAAUCUAUGGGGUGGUGGUGCAUG	1	no match			
10877	UUCUGAGAAAAUUAAGUGCUGGAA	1	0	transposon	LINE	Circe
10878	CCAGUUCCUCUGCCGCGUGCCCAC	1	0	repeat		trf
10879	UCUGUGAGGUUGCGCCUGAGCCCCGA	1	0	transposon	LTR	blood
10880	AACAGAGAGAAGACUCACCCGACGA	1	0	no annotation		
10881	GCCCGGAUAGCUCAGUCGGUAGAGCAUUGU	1	1	ncRNA		tRNA
10882	GAAGACGAAAUCCAAACUGACAAUGG	1	0	transposon	LINE	Porto1
10883	GGUCUUCCUGCGCUUGCGUCCCGUUA	1	0	coding gene		CG12298-RA|sub
10884	AAAAGGAGCAAUUUUUGGCCGUCG	1	no match			
10885	AGAGACCCGUCGCCGCGCUCCUCCCCCCCCU	1	no match			
10886	CCGGAGGGGGAGCCUGAGAAACG	1	1	ncRNA		rRNA
10887	UCUCCGCGCAUGUUGACUGGGUA	1	0	transposon	LTR	opus
10888	UCAGCAGCGGGAUGAUCAGUAGCAC	1	0	transposon	LINE	R1-element
10889	UAGCUUGCACCAUCACCAUCACCAU	1	no match			
10890	UUGUUUCGCAAAAGUUGACCGAACUUGA	1	0	ncRNA		rRNA
10891	AGGCCAGAAACCACCCUCGUGAAGU	1	0	transposon	LTR	opus
10892	UGCCGGAGCGAGGAGGAGGACUAA	1	1	Su(Ste)		Su(Ste)
10893	CCAGGAAAACGGUCAGGUACGGAG	1	0	transposon	LTR	roo
10894	CCAGGCCCGUAUAACGUUAAUGAU	1	0	ncRNA		rRNA
10895	AGCCCCCGGGGAGCCGCCCUUGGAAU	1	no match			
10896	GGAGUUCUCGAUGGCGCCCAGA	1	0	coding gene		CG10252-RA|CG10252
10897	CGAAUCAACAAAUUGCGCCGAUA	1	0	coding gene		CG41545-RA|CG41545
10898	CAAGAUAUGACCAAAACAAAGCA	1	0	transposon	DNA	1360
10899	CGAAAACCGCAAUUACUUCUGCACCAGUCC	1	no match			
10900	GCGCGUCUGCCCCCCCCCGCCGGGUGCGC	1	no match			
10901	UCCCACGGAACACCCGAGCAAUGCCC	1	1	transposon	LINE	G2
10902	GUGCUCCUAAACGAGUGUUAUUGUGGG	1	1	ncRNA		rRNA
10903	AAACCUGAACAUGAUCCUGAGUCG	1	1	transposon	LINE	Helena
10904	ACAUAAGCAAGGACCCGUAGAGC	1	0	transposon	LTR	GATE
10905	AGGCGCGCAAAUUACCCACUCCCGAC	1	no match			
10906	CUUGUUCCCCGGAUAGUUUAGUUACGC	1	1	ncRNA		rRNA
10907	ACCCCUCCCACGGCGCGACCUCU	1	no match			
10908	UACCAACCCAAAAUAUGACUGCAUU	1	no match			
10909	CAAAUUCCGAGAAUGCAGGUUCCU	1	0	transposon	LTR	Rt1c
10910	UUCGAUUCCGGAGAGGGAGCCUGA	1	0	ncRNA		rRNA
10911	CCAAAGCUUUUGGGCUCCGGGGGAAG	1	0	ncRNA		rRNA
10912	UGAGAGGCGCCGUAGUUACCACCC	1	0	no annotation		
10913	GCUGGAAAAAAACGAAGUAUAACGA	1	no match			
10914	GGCUCCGGGGGAAGCAUGGUUGC	1	1	ncRNA		rRNA
10915	CAGCAGCGAUGCGGGUCCUGAGG	1	0	coding gene		CG6061-RA|mip120
10916	GGUGCGCGAGAUAGCCCAGGACUU	1	0	coding gene		CG5825-RA|His3.3A
10917	UGAACGAUGACGAUCCCACGGCUGGAGC	1	1	transposon	LTR	ninja-Dsim-like
10918	UACAAGUACACGUAGGCUGCGAAA	1	1	repeat		trf
10919	UAUUACGACGCGAGAGGUGAAAUUCU	1	0	ncRNA		rRNA
10920	GACAACCAGAGAUCUAGAGGGCCCU	1	1	no annotation		
10921	UAAGGUAAAACUUAUCAUGCAAGCAG	1	0	transposon	DNA	NOF
10922	CCCCAAGAUGUCUGCCACAAAG	1	1	coding gene		CG13315-RA|CG13315
10923	GAGGAAUUUGAGCUGCGAUGACGCC	1	0	transposon	LTR	ninja-Dsim-like
10924	CAAGAUAUGACCAAAGCAAAGGCA	1	1	transposon	LINE	Cr1a
10925	AAUGCUCGCAAAUGGAUUGUGGGACCA	1	0	coding gene		CG2263-RA|CG2263
10926	ACAUCUGUGAGACUACGCUUGAGUU	1	1	transposon	LTR	gypsy2
10927	UAUUUGUGCACAACACGAGCCCCAGGU	1	1	transposon	LINE	Circe
10928	CACGAGGAGCUCGGGAUUCCAUGGUUA	1	1	transposon	LINE	G2
10929	AUCAACAACGAUGUGGUAGUGAGUG	1	0	coding gene		CG3249-RA|yu
10930	GCAACCCAAAACGUAACGGCAGAGG	1	0	transposon	LTR	HMS-Beagle
10931	AUGCCCUGCAGACCAUUACUCGUCAU	1	0	no annotation		
10932	AACCUAACUAAAUGAUGUUUCCCAG	1	2	no annotation		
10933	GACCAUAAAGAAGCCGUCGAAA	1	1	ncRNA		rRNA
10934	CUGUGAACGAGUGAACUGGCAACA	1	0	Su(Ste)		Su(Ste)
10935	UGAGCGAACCCGGGGAACUGAA	1	no match			
10936	AUCCGCGGCUAGACGGAAAGACCCCGUGA	1	no match			
10937	AGAGCAUAAAGACGACGAAAGUGAUC	1	2	transposon	LINE	Doc3
10938	GUAUCUAGGACUACCAUGGUUGCAACG	1	0	ncRNA		rRNA
10939	AGUGGACCCGGUUUCUGGAGAGGUG	1	0	coding gene		CG31226-RA|CG31226
10940	UAGUACGAGAGGACCGAAUUGGACAAA	1	no match			
10941	UGAUAACUCAUAUCGUAAACGGCCUC	1	0	transposon	LTR	flea
10942	AACGUGACGAUCCCACGGCUGGAG	1	no match			
10943	AGGAUGCCGCUAAGUUCUCGCUGAGAG	1	1	ncRNA		alphagamma-element
10944	UCGAUCUAGAGGAGCUCCUUGAAA	1	1	transposon	LTR	flea
10945	CUGACUAAGAGGGCUGCACAGUUUCC	1	0	transposon	LTR	HMS-Beagle
10946	GACACCAGGAAGAUUGAUGAGGCC	1	0	transposon	LINE	Rt1b
10947	AAAAGACUUAAAGAUGAUGAACCCA	1	1	transposon	LTR	gypsy8
10948	GAACGAAAGUUAGAGGUUCGAAGG	1	0	ncRNA		rRNA
10949	AGGAUAGUCCCGAAGAAACCUAAACC	1	1	transposon	LTR	opus
10950	GCGGUUAAACCAUAUCACGUACAGUC	1	0	ncRNA		snoRNA
10951	ACCAAAUUAAUCGCGAGGCUGCAAG	1	1	transposon	LTR	mdg3
10952	GCCCAGUCCCAAUCGCAGCCGAUGGCCACC	1	0	coding gene		CG11866-RA|CG11866
10953	UGCAACAUAAUGACGGUUUAGUAAA	1	0	transposon	LINE	Cr1a

10954	GCCGAGAAAAAUGAACACGAAACGA	1	0	transposon	LTR	Quasimodo
10955	UUUGCAAAAACGAGAGCGACAGAGC	1	1	transposon	LTR	micropia
10956	AAGACCUAUAAACACACCUCAACAGU	1	no match			
10957	CGCAUAAUCAAAACAUCCGAUGACG	1	0	transposon	LTR	McClintock
10958	CCCGCGGAAAGACGGAAAGACCCCA	1	no match			
10959	CUACUCCGGAUGUACCCAACAGAUACCAU	1	1	transposon	LTR	Idefix
10960	AUGUCUAUGAGAUGACCGUAAUG	1	0	transposon	LTR	ninja-Dsim-like
10961	UCCAUUCACUCCAUGGCGUUACCACGA	1	0	transposon	LTR	springer
10962	AUAGUACGUAGGCAUAAUUCCAAUA	1	1	transposon	LINE	Rt1b
10963	UACAAUACCCAGUUACUCGUGCC	1	0	no annotation		
10964	UCUCAUCGUCGUAGAAACAAGCCCCGA	1	no match			
10965	AAAAAACCGAAUGCGAUCCGGAUU	1	1	coding gene		CG4714-RA|CG4714
10966	CACACGAAAAUACACGUUCGAAGAGC	1	0	transposon	LTR	GATE
10967	CGCUAGCUCAUUUCCCGGGAUGUUU	1	0	transposon	LINE	baggins
10968	AUGAAAACAGUUCUCAUUGGAGGCCA	1	0	transposon	LTR	flea
10969	AAAACCUGAAGAUGAUCCUGAGC	1	1	transposon	LINE	Helena
10970	GCAAGACGACACUCUGCAGGAACCCAAU	1	0	repeat		trf
10971	CAAGAUAUGACCGAAAACCCAGACAC	1	no match			
10972	CGGAGAAGCCGGCGGGAGCCCCGGGGAG	1	no match			
10973	AUAUAAAAAUCCUCAACUGUGAGG	1	no match			
10974	AAUCGCUCGAUGUCGUGAGGUGA	1	0	no annotation		
10975	UUAGACAGAAAAUAAGGGGACACGCC	1	0	transposon	LTR	opus
10976	ACCAUGAAAGGCGUUGGUUGCUU	1	0	ncRNA		rRNA
10977	GUAAAUGAAGAAAAUAUUGUCGAUGU	1	0	transposon	LTR	invader1
10978	AAACCAAGCUAGAGUACAGAUCGAAA	1	1	transposon	LINE	jockey
10979	CGUUCAAGGACCACGCGAGCUACGCCAU	1	1	transposon	LINE	Rt1b
10980	CCCAGAAACACAGUGUUCGACGCCA	1	1	coding gene		CG18743-RA|Hsp70Ab
10981	AAGUCCCGCCACCGCACCCAUAGAA	1	0	transposon	LINE	BS
10982	CCCCGGAACUUGGACACGGCGAUGCCGGU	1	0	coding gene		CG3982-RA|CG3982
10983	GAUGCAAACAAACAAGAGACGGCA	1	0	transposon	LINE	I
10984	GUUGCAAGAAGUGCGACCGAUCCCA	1	0	transposon	LTR	GATE
10985	UUGGGAACACCGCCGUGUUGUUGGCCCU	1	no match			
10986	AGGGACAUUAUUGUGGAAAAUGUGACA	1	1	no annotation		
10987	UGCUCGAGGGCCCGUGGUUACCAG	1	0	coding gene		CG3800-RA|CG3800
10988	ACCCUACUGAUGAUGUGUUGUUGU	1	no match			
10989	CUUUAAUGUAUUAAACGGGCGGCC	1	0	transposon	LTR	invader6
10990	CAGUGGCGAAGGCGUCUAUCUGGUUCA	1	no match			
10991	AAUCUCGCGUCGGGCCGUACCCAU	1	no match			
10992	CGCCCAAUCUGCGGAUCCGAACACCAAC	1	0	coding gene		CG5016-RA|Mst57Db
10993	AAAACCUGAAGACUGAUCCUGAGUCGA	1	no match			
10994	CCUGCAUGAAGAUGAUUCCGAAGAUC	1	0	transposon	LINE	Circe
10995	UGUAAGCGAUCUUGAGGCGACAGU	1	0	transposon	LINE	G3
10996	UGCCUUCGAAGAGUCGAGUUGUUUGGG	1	no match			
10997	AUACUACGGAACCACCGUGCCCAGUGC	1	0	transposon	LTR	opus
10998	AUGGUUGCAACGGGUAACGGUGAAUCA	1	1	ncRNA		rRNA
10999	GGAGUGAUUUGUCUGCUUAAUU	1	1	ncRNA		rRNA
11000	ACUGAUAACGAAAUGGAUGAC	1	no match			
11001	GCGCUUGUGACGGCCCGUGAAAAAUCC	1	no match			
11002	UCGGGAGCGAGGCAUGUCUGUCUU	1	0	transposon	LTR	invader1
11003	UCCCCGUGGCGUAGGUCCCCACCGA	1	no match			
11004	ACCUCUCGAGCAUCGAUGCAAGC	1	0	transposon	LINE	INE-1
11005	GUUUAAAGUCGAAUAGGUUUUCCGUGU	1	1	coding gene		CG18449-RA|CG18449
11006	ACGAUCAAUAAAACAUACGCCCGGUU	1	1	transposon	LTR	GATE
11007	UAGAGUAUUCGAUCUUGAGGCGUGUGA	1	no match			
11008	CGACAAAGGAGUAUUGCCAGAUGAG	1	0	transposon	LTR	gypsy8
11009	UGCUCUCGAAUGUAUUCCCGAUAU	1	1	transposon	LINE	R1-element
11010	ACCCUUGAUAAACGUAUGACUUUCGGU	1	1	transposon	LINE	BS
11011	GAUAGGCCGAUAGCGUACAUUUCCC	1	0	transposon	LTR	opus
11012	UUCGUGGAGUGAUUUGUCUGGUUAAU	1	0	ncRNA		rRNA
11013	CCAUCCGGUAAGAUUGUGCCCGCGU	1	0	transposon	LTR	Rt1c
11014	AAUACCACCCUGAACGCGCCCGAUCUCGU	1	no match			
11015	UCCGAGCUAAUAUGGCACCAAG	1	1	no annotation		
11016	CCAAGAGUUGACCGAACUUGAU	1	2	ncRNA		rRNA
11017	UACCAACCCAGAAUAUGACCCAGC	1	0	transposon	LTR	McClintock
11018	CCAAAAGAUAAAAACGUUUACGAACAAU	1	0	transposon	LTR	mdg3
11019	AACGAGUGUUAUUGUGGGCCGG	1	0	ncRNA		rRNA
11020	CCGAAGAUGACCAAGUUGGCAACC	1	1	transposon	LTR	gypsy6
11021	ACCACCCUGAACGCGCCCGAUCUCGUCU	1	no match			
11022	AAAACCUGAAGAUUAUCCUGAGUCG	1	1	transposon	LINE	Helena
11023	GCUCCGGGGGGAAGUAUGGUUGC	1	no match			
11024	UGGAAGAUUAGCUCGAGUAGAACUGU	1	1	transposon	LTR	3S18
11025	UUCUGCCCGUCGAUGUGCCUAUGCGCCC	1	0	coding gene		CG8349-RA|CG8349
11026	AAGAGGGCGUGAAACCGUUAAGAGUU	1	no match			
11027	AGCACGAGCGCACGUGUUAGGACCC	1	no match			
11028	GCUUUAAUCACGAGCAAUGCGGUGG	1	1	no annotation		
11029	UGCUUGCACCAUCACCAUCACCAU	1	no match			
11030	GAAGAAGUAAACAGCGACUACACG	1	0	transposon	LTR	micropia
11031	CGCACAAACGCAACGCUCCUGUACAA	1	0	transposon	Low_complexity	F-element
11032	CAUUGGAGGGCAAGUCUGGUGCCA	1	1	ncRNA		rRNA
11033	UUAAACAUGACACUUCUGGACA	1	1	transposon	LTR	297
11034	CACCACAAAUGAUUAUUGAUACCGGA	1	0	transposon	LINE	Cr1a
11035	UGUCCGCUAGCUCAUUUCCCGGGAUGUU	1	0	transposon	LINE	Fw2
11036	ACUCGCCGAAUCCCGGGGCGAGGGAGUC	1	no match			

11037	UGUUCCCGGCGGUGGCGUCGUCCCUG	1	no match			
11038	UAGAUGUGGCCUGAAGAUGAUUUUCC	1	0	transposon	Low_complexity	F-element
11039	AUCUCAUCGUCGUAGAACAAGCCCGAGG	1	1	Su(Ste)		Su(Ste)
11040	CUCGAGAUGUGUUCGGAAGGAUCGG	1	1	coding gene		CG3506-RA|vas
11041	AUUGUGGGCCGGUACUAUUACUUUG	1	0	ncRNA		rRNA
11042	UACGACGAGAAUGAGAUCAGCGCG	1	0	transposon	LTR	frogger
11043	UCGGGCCCGAACCCUUCUGGUGUGUUGA	1	2	transposon	LINE	baggins
11044	UCCAGAGAGUCGUGUUGCUUGAUAG	1	1	ncRNA		rRNA
11045	AAAAGGGGAACUGAGGAACGCAUU	1	0	transposon	LTR	invader3
11046	UCCAUCUCGAAAACGUUACCAAAG	1	0	transposon	LTR	ninja-Dsim-like
11047	CCCGCCGAGGUACGAUUCCUGCACC	1	1	transposon	LINE	R1-element
11048	GGAGUGCCCUUCGUCCUGGGAAACGGGGUC	1	no match			
11049	UGACCAUUAAAGUGUGACCCACCAAA	1	0	transposon	LTR	Quasimodo
11050	UAAACCGUCAUUCCAGCGCAAGUUGCCCU	1	1	transposon	LTR	ninja-Dsim-like
11051	ACCGCAACGAACUCCCAGCCCAGC	1	0	transposon	LINE	Doc
11052	UGAGACUACAAACGGAUUGCGGACCGAA	1	1	no annotation		
11053	UCCCAUAUGAAGAAGCUCCUGAGCGU	1	0	transposon	LTR	blood
11054	GAUCUCAACUGUGAGGGGGAUCCUCAA	1	no match			
11055	UCAACUGUGAGGGGAUCCUCCAA	1	no match			
11056	CAGACAAGUAGAGGGCCGUCGCCG	1	0	transposon	LTR	accord
11057	UGCUACUCGAAUGUAUGCCCGAUAUU	1	no match			
11058	UGAGGACUUGAGCGAUUUAAGUGGCA	1	1	Su(Ste)		Su(Ste)
11059	UGGACAUCCGGAAGAGUUUGCGAC	1	0	coding gene		CG6963-RA|gish
11060	CAAACCAAUCCCAACCAAACCCG	1	0	transposon	LINE	G2
11061	GUCCCGAAGAAACCUAAACCAAA	1	0	transposon	LTR	opus
11062	UCUGCCGAACGGUCUGCGGAUUGCC	1	1	coding gene		CG8728-RA|CG8728
11063	GUGGAAUCUAACCGACCGCCAUGACCC	1	no match			
11064	GUUCAGUACACACAGGGACAGCAAAA	1	0	ncRNA		rRNA
11065	CCGAACUUGAGACCGCCCCCUGGACGACCC	1	no match			
11066	GUCGGAGGCUGGCGGCGGGGGGU	1	no match			
11067	UUGCCUCUGGAUGAUGGCUGAUGCUCG	1	0	transposon	LINE	G2
11068	UGUGGAAUUAUGACUGAACGCCUCU	1	no match			
11069	GCGCACAGAGGAGAGCAGAUGAGAGCA	1	no match			
11070	GGGGGUAUAGCUCAGCUGGGAGAGCGCU	1	no match			
11071	AGGAGGCUAACGAGGGGAACUGAAA	1	no match			
11072	AUCCGCCUGGGGAGUACGGUCGCA	1	no match			
11073	AGAACUGGCGCUGCGGGAUGAACCGAA	1	2	ncRNA		rRNA
11074	ACGGAGGUGUCCCAAGGCCAGCU	1	0	ncRNA		rRNA
11075	UAUCUAGUACUACCAUGGUUGCAACGG	1	1	ncRNA		rRNA
11076	UGGCCAGCUAGUCCUCCUCCUCGCCCCG	1	1	Su(Ste)		Su(Ste)
11077	UGAGAUAGUACAAAUCGGUUACGAAAA	1	1	transposon	LINE	I
11078	UCCCAGCUCGGGGAGGUAGUGACGAAAA	1	0	ncRNA		rRNA
11079	GACUUCCGAACUGGCCGAUAUGAA	1	no match			
11080	UCCGAGGAACCAAUCGAUCCAGCUG	1	1	Su(Ste)		Su(Ste)
11081	GUGAUCACUGUGACGCCUUGCGUGU	1	0	ncRNA		rRNA
11082	GCAUGAUUCACGCCCGAUACAUAGUGUC	1	0	coding gene		CG40635-RA|CG40635
11083	GCUACCGGGCGUGAGCAUCCACAAUGC	1	0	transposon	LTR	ninja-Dsim-like
11084	AAUGGCUCUCUCAGUCGCUUCCCCGGGGGA	1	no match			
11085	UAGUUUCAUCGUUAGACGGCUCGG	1	1	no annotation		
11086	GCUGCACGAAAUGAACCCGAACGAA	1	0	transposon	LINE	Doc3
11087	ACGACUGGGAGGAUCAUCUGGCCGUC	1	0	coding gene		CG1242-RA|Hsp83
11088	CAACGAGUUCCUCUGCCGCGUGCCC	1	0	coding gene		CG42398-RA|SteXh:CG42398
11089	UCCGCACUUCGUCCCUCAAGAUAUGAAC	1	0	transposon	DNA	1360
11090	AAACUUGAAAUCCAGGUGAUCCGGC	1	no match			
11091	AACGAGUUCCUCUGCCGCGUGCCCAC	1	0	coding gene		CG42398-RA|SteXh:CG42398
11092	ACCGGUCGAUUCCGAAUUCAGACGUGCU	1	0	coding gene		CG8649-RA|Fim
11093	UCCAGAAUGAGGACCUGAGGGGCAU	1	0	transposon	LINE	Rt1b
11094	AGUUAACUAUCCCAAAGGCAUCACGC	1	0	no annotation		
11095	AACUUCGAUAGCGAACUGACUCUGUAA	1	no match			
11096	ACGAGAGGAACCGCAGGUACGGACCA	1	0	ncRNA		rRNA
11097	UAAGCUAGACUAUGUAUCCCCGGGUAU	1	2	transposon	LINE	Doc
11098	ACCAUGAAAGGCGUUGGUUGCUUAAAA	1	0	ncRNA		rRNA
11099	UGCUGACUCGGCUGAUGUUUCUCC	1	0	transposon	LINE	Rt1b
11100	UUUGAGAGGCGCCGUAGUUACCACC	1	0	no annotation		
11101	AAACGAUGCCGACCGGCGAUGCGGC	1	no match			
11102	UCAACCAGCACGACAAGGCCCAUACG	1	0	transposon	LINE	G2
11103	AGGAUCCCAAGCUGACUCCCGCCGGCCA	1	0	coding gene		CG13340-RA|CG13340
11104	CAUGAAUGAAUACAUGGAUUUAGGAC	1	0	transposon	DNA	HB
11105	ACACCGCCCGUCGCUACUACCGAUUG	1	0	ncRNA		rRNA
11106	AAGGCGCGGGUAACCCGUUGAACCCCAUUC	1	no match			
11107	UCCAUCUCGAAUACGUCGCCAGAGCGCCU	1	1	transposon	LTR	ninja-Dsim-like
11108	GACCGUAGCACCGAACGAGCUCCUUUC	1	0	transposon	LINE	Rt1b
11109	CACAGAGUACGGGGGCUUACUAAC	1	no match			
11110	UCGAACGAUAAACGGUGCUAACCAGCCGG	1	0	transposon	LTR	GATE
11111	UGCUCUCGAAUGUAUGCCUGAUA	1	1	transposon	LINE	R1-element
11112	GUUCCUCGGGGAUCAAGGGCAACGAG	1	1	no annotation		
11113	UUAGAUGUCCUGGGCUGCACGC	1	0	ncRNA		rRNA
11114	ACCUAAAUGAUGCUGGUAAAAAC	1	0	transposon	LINE	Doc
11115	CCGACACCGAGCGCCUCAUCGGA	1	no match			
11116	CAGAACUGGCGAUGCGGGAUGAACCG	1	no match			
11117	AACACGGGAAACCUCACCCGGCCGGAC	1	no match			
11118	UUCUUGACAACCCGAUCGGACCUCACGC	1	0	transposon	LTR	invader4
11119	AGGUAGACGACCGGGCGAUGGCGA	1	0	transposon	LINE	R1

11120	UGCACAUUGACAUAUUUUCAACCGA	1	0	transposon	LTR	gypsy
11121	AAAGCAGAAAUGUUUGUCCGAUAU	1	0	Su(Ste)		Su(Ste)
11122	GCGGGGAAAGAAGACCCUGUUGAG	1	1	repeat		LSU-rRNA_Dme
11123	CGAGAUGUUCAACUACACGGAUUCG	1	0	transposon	LTR	3S18
11124	ACUCGGCUCGUGCGUCGAUGAAGAACG	1	2	ncRNA		rRNA
11125	UCCUGGGAAAACGGGGGCGCGGCCGGAAA	1	no match			
11126	ACUCGCCGAAUCCCGGGGCCGAGGGACG	1	no match			
11127	UUGUGAGUUAUCCGUCAGGUGAGGCGA	1	0	transposon	LTR	micropia
11128	GCAUUCAAUACAGCGAGAUGGCUC	1	1	transposon	LTR	Quasimodo
11129	AAGACGAGGUUGACGCCGCGGGUUC	1	0	transposon	LTR	gtwin
11130	UGGCUCUCUCAGUCGCUUCCCCGGGA	1	2	ncRNA		rRNA
11131	UGGUAACCAGCUCGUCGACCUUAGC	1	no match			
11132	UGCCUGAGAAACUCGCGCCCGUUGAGG	1	1	coding gene		CG3373-RA|Hmu
11133	AUCAUCAGUUUAUUGCGGGAUGA	1	0	transposon	LINE	Circe
11134	UGUCCCAAGGCCAGCUCAGUGCGGA	1	0	ncRNA		rRNA
11135	ACGAAAACUCAAUAUGGUAACAAUGGCA	1	1	transposon	LTR	HMS-Beagle
11136	UGAGCUGUGGACCGAGCGGAAUGA	1	0	coding gene		CG4821-RA|Tequila
11137	CUCGGAGGUGGACUCGCCAGCAACA	1	0	transposon	LINE	X
11138	UGCCAUUCCAGGUGGUUCAAGCCC	1	0	transposon	LTR	GATE
11139	AUGCCGGAGCGAGGAGGACUAGCU	1	no match			
11140	UGCACAUUAAGUUUUGAUCUCCGAAU	1	2	transposon	LTR	invader1
11141	CAUCUUCUUAACGCAGCUGCCGAA	1	0	Su(Ste)		Su(Ste)
11142	AACACCACAAGGAGACGCAUGAUCCGCA	1	0	coding gene		CG10366-RA|CG10366
11143	AACCGAAUGAGACCUUGGAGGACU	1	1	ncRNA		rRNA
11144	UGUGACUACAAACGGAUUGCGGACC	1	1	no annotation		
11145	ACCGAGAGAAGACUUUCACCCGAUG	1	0	coding gene		CG12717-RA|CG12717
11146	GUCCCGGAUACAUUGUCUCGUGUA	1	0	transposon	LTR	gypsy12
11147	CAGAUCAACAAAGUACGAUGGGGAACCAC	1	0	no annotation		
11148	CACAGAGUACGGGGCUUACCGAAGUCA	1	1	transposon	LINE	baggins
11149	UAAACCUGAAGGAGUGAAAAUGCU	1	0	no annotation		
11150	GCCUAAGUACUCCUCAGCGACCGAUA	1	no match			
11151	CAGAUCAACAAAGUACGAUGGGAA	1	1	no annotation		
11152	ACCUGGUUGAUAGCGUAGACGACGACC	1	1	transposon	LINE	Rt1a
11153	AGACCCGCAAGGACCCGUUCAUCCUGGA	1	0	coding gene		CG6213-RA|Vha13
11154	UCAAGCGAUAUUCGAGGCCCAACG	1	0	no annotation		
11155	GGUAGAGAUAGGCCGAUAGCGUACAU	1	0	transposon	LTR	opus
11156	ACCAUUUACAAAAUGUAUUCCGACCA	1	0	transposon	LINE	Doc
11157	CACGAAAUGUCGUUACCGAGAAGA	1	0	transposon	LTR	copia
11158	CAGGAUCAAAUAAGUAGACUUCGAG	1	0	transposon	LTR	micropia
11159	UCGAGAAGCGAACCCGGCGAACUGA	1	no match			
11160	CACGAUUACAAACACUGAUUUGCCC	1	0	transposon	LINE	S
11161	AUCGAAUGGAGUGCCAGAACCCGUUGU	1	1	transposon	LINE	R1-element
11162	GUCGUGUUGACCAACUGUUGCGACGU	1	0	coding gene		CG4286-RA|CG4286
11163	AACGAGCGCACGUGUUAGGACCCGAAAG	1	no match			
11164	GACAACCAGAGAUCUAGAGGGCCCAG	1	1	no annotation		
11165	AGCUUCAUCCAAGCGAACUGUGCCCGUA	1	2	transposon	LINE	R1-element
11166	UGAAGAAAUAAUUGAAAUGAAGCCG	1	0	transposon	LINE	baggins
11167	AUCCGGUAAAGCGAAUGAUUAGAGGUC	1	2	ncRNA		rRNA
11168	AUCGCAAAGAAGCUGUUUUGAAGGACCA	1	0	coding gene		CG41423-RA|CG41423
11169	CUUGGACCGAACACUUUGAACCAA	1	0	Su(Ste)		Su(Ste)
11170	AAAAACUCAAAGAAAAAGACGAUGAC	1	0	Su(Ste)		Su(Ste)
11171	CUACCGCAUAUCGGACAAAGAGGCCCGC	1	0	coding gene		CG2127-RA|CG2127
11172	UUCAAAGUUACCCCAGGAUGCUGUGC	1	0	no annotation		
11173	GAAACCAUCAGACGCCUCCAGAGG	1	0	transposon	LINE	G
11174	CAUGGAGGAGUCCGUGGCAUUCAGGUU	1	0	ncRNA		7SLRNA-RA
11175	UCGGCCCGAACCCUUCUGUGUG	1	2	transposon	LINE	baggins
11176	GAGUACUUCAGCCAGCACUGGA	1	0	Su(Ste)		Su(Ste)
11177	UGCAUUGCCUUAACGUUAUUAUUA	1	0	transposon	LTR	297
11178	GGGUCCUGAGAGAUGGGCGAGCGCCGUU	1	no match			
11179	CGACCCACCAGGAUGAUUAAGCGGCCAC	1	0	coding gene		CG32435-RA|chb
11180	CAGAGGUAGACGACCGGGCGAUG	1	0	transposon	LINE	R1-element
11181	GCUCUGAAGAUUGAGAUAGUCGG	1	0	ncRNA		rRNA
11182	ACUCGCCGAAUCCCGGGGCCGUGGGAGC	1	no match			
11183	UAACGAGAGUAGUGAAGAGGAAGAU	1	1	transposon	LTR	Dm88
11184	CAUGGGAAUCCGAGGUGCCGUAGAU	1	no match			
11185	ACCCGUCAAACCCCACCUGAAUCUCCGC	1	0	transposon	LINE	G2
11186	AAAACCGAUAUGCUAUCUUGGCUGA	1	1	transposon	LINE	Doc
11187	UCGACAACCAGAGAUCUAGAGGGCCCAA	1	0	no annotation		
11188	GAAGGACGACCACGUUGUAAUGCC	1	0	transposon	LTR	ninja-Dsim-like
11189	AUCCCACGUAUGUUGGCCGGAUUGUCG	1	1	transposon	LTR	invader1
11190	AGUGGCGCGUCGCGAGGCUAACCCGUG	1	no match			
11191	UACAAUUCCAGCUCAACUCCCGAA	1	0	transposon	LTR	Quasimodo
11192	AUACCCUGGUAGUCCACGCCGUAAA	1	no match			
11193	UCUUUGGACAGAAAUACACGGGUUGA	1	1	transposon	LINE	Cr1a
11194	GUGUCCGAUCAGACGAGAUGGCCAU	1	1	transposon	LINE	F
11195	CUAACUGUGAGGGAUCCUCAA	1	no match			
11196	UGAACACUCAUAAGGCCCACGAAGC	1	0	transposon	LTR	flea
11197	UGAUAUCCUGAGCAAGUGCUCUCACCGA	1	0	transposon	DNA	S
11198	UUAGAUGUCCUGGGCUGCACGCGCGC	1	0	ncRNA		rRNA
11199	GUAGUGACGAAAAAUACAAUACAG	1	no match			
11200	GCCCCCUGAUGAAGAAGAGAUGCCCAG	1	2	transposon	LTR	ninja-Dsim-like
11201	UCGGACCUAAGCCAUAAAUGUAACGCG	1	0	transposon	LTR	Quasimodo
11202	ACCAGAUCGAUACCGCCGCCCUGAU	1	0	coding gene		CG2184-RA|Mlc2

11203	UCUGCAAUCAUUACCGAGCUGUCAA	1	0	transposon	LTR	gypsy8
11204	GCCUACCCGUCCGAACCAAACUCC	1	no match			
11205	UCGAUCCUAGGAGAUGGCACCA	1	no match			
11206	GUGGACAAAGACGGCUGCAAGCAUCC	1	0	coding gene		CG4836-RA|CG4836
11207	CACCACAAGUGAUAAUUGAUUCCGGA	1	0	transposon	LINE	Cr1a
11208	CCCCAUCCAAAGCCACAGUACAGCGC	1	0	transposon	LTR	GATE
11209	UGCACAACUAACCGUGUGACGAAGGC	1	1	transposon	LTR	micropia
11210	UCUCACCGUCGUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
11211	GCCGAACGUAGUCUCUCGAGAGCGGU	1	2	transposon	LTR	invader1
11212	AAUGCCGCGUGAGUGAUGAAGGCCCU	1	no match			
11213	UCCUAAAGGGUUCUGUAUUUCCGA	1	no match			
11214	AGAGACUUAAAUAUCCCAUCAGUUA	1	0	transposon	Low_complexity	F-element
11215	ACCCCUGAACGCGCCCGAUCUCGUCC	1	no match			
11216	UAUUACAUAGAUGUAGAUGGGUCG	1	0	no annotation		
11217	UGGAUACCGCAGCUAGGAAUAAUGGAA	1	no match			
11218	ACGAACUAGAGCGACAUCGGGAA	1	0	transposon	LTR	roo
11219	GAUCCACUGAAGCUUAUCCUUUGCUU	1	0	ncRNA		rRNA
11220	UGCGCGAAGAAGGAGAACGAGAAGGC	1	0	transposon	LINE	G6
11221	CGUAAAACAAUUCAAGAUCUGGCCGU	1	1	transposon	LTR	blood
11222	GUGGCUUUGAGAGGCGCCGUAGUUAUCC	1	2	no annotation		
11223	AGCACUCUCAUCCAAUAGACGGACACAC	1	1	no annotation		
11224	ACUCGCCGAAUCCCGGGGCCGGGGGAGC	1	no match			
11225	ACCCACAGAGUACGGGGCUUACUGAAGUCA	1	2	transposon	LINE	baggins
11226	UCUUGAAUCACCGAAUUGAAAUUGUA	1	0	no annotation		
11227	GAAAGAAGACCCUGUUGAGCUUGACUCU	1	1	repeat		LSU-rRNA_Dme
11228	GGCUUCGAGAGGCGCCGUAGUUAA	1	2	no annotation		
11229	UCCUUAACAAGAACGACGUUGUCCAC	1	0	transposon	LTR	diver2
11230	UGUGAAAAGAUCCCGUUUGAAUAUGAC	1	0	transposon	DNA	S
11231	GUCGAGAAGAAGAUGAAGCGCAACAA	1	0	transposon	LTR	ninja-Dsim-like
11232	CCCCAAAAACCGUCUCAGUUCAGAU	1	no match			
11233	UGCUCUCGAAUGCUUGCCCGGUAUU	1	2	transposon	LINE	R1-element
11234	UCUUGUAGGUGAGUCUCGGUGAUCGAGUG	1	0	transposon	LINE	G2
11235	AACGGAGGCGCGCGAAGGUUGGCUCA	1	no match			
11236	CGGACAGAGACGGCGCAUUUUCCAGU	1	0	transposon	LTR	McClintock
11237	UGUGAUUGUAUGAUUGUCCAGUGGU	1	1	transposon	LINE	G2
11238	CUGUAUCUAACUCGUCAUCAGUACGG	1	0	coding gene		CG1009-RA|Psa
11239	UGAGACUACAACGGAUUGCGGACC	1	no match			
11240	UGGAACAUUGCCGUGAACGCAGACAAA	1	0	transposon	LINE	G
11241	CCACCCCUGAAACGCGCCCGAUCUCGUUC	1	no match			
11242	CAUUCCUGCGUAGUACGAGAGGAACCG	1	0	ncRNA		rRNA
11243	UGAGUUAAAAGAAAAAAUUGAGAAAA	1	1	transposon	LTR	mdg3
11244	CCAAUCCGCUGCGUUCGGGAGCCGGGAA	1	0	coding gene		CG9393-RA|CG9393
11245	UGUCACGCGGGAGACCGGGGUUCAAU	1	0	ncRNA		tRNA
11246	UGGGAACACCGCGUGUUGUUGGUCCU	1	no match			
11247	AUUUAAAUAAUAUUAUGAGAAUA	1	0	ncRNA		rRNA
11248	GAUGUCCGGGGCUGCACGCGCGCUAC	1	1	ncRNA		rRNA
11249	GGCUUUGAGAGGCGCCGUAGUUU	1	1	no annotation		
11250	CUGGCAAAGAAAAUUAAGAGUC	1	1	transposon	LTR	mdg3
11251	GCCGAAAAUACGACGACCCCAAGUAA	1	no match			
11252	UGAAGAAAUAAUUGAAAUGAAGCCGA	1	1	transposon	LINE	baggins
11253	UCCGAAAACAUCUGAUGCACCUGU	1	0	transposon	LTR	Quasimodo
11254	CAGAAGAAAAACAAAGCGCCGUUGUCA	1	no match			
11255	GUUGGGAACCCUAAUGGGACUGCCGGU	1	no match			
11256	AGUCUAACACGUGCGCGAGUCGGGGUC	1	no match			
11257	ACGGGAAACCUCACCCGGCUCGGACAC	1	no match			
11258	UGGGACCACCGCGUGUUGUUGGCCU	1	1	ncRNA		5SrRNA
11259	UACGAGGAUGUCCGAGCUUAUCACU	1	0	coding gene		CG6289-RA|CG6289
11260	ACUGAAUAAACGAAAUGGAUGACAC	1	0	transposon	LTR	invader4
11261	AAGACGAGAAGACCCUAUGGAGCUUU	1	no match			
11262	UCUGCAUGAAGACGAUUCCGAAG	1	0	transposon	LINE	Circe
11263	GUAAUACAUCGGAACGUGCCCAAUCG	1	no match			
11264	CGCCACAGAAUAAACUAAUAACGAG	1	2	transposon	LTR	frogger
11265	UAGAUCAGUGAAAUCAUCUUCCGGGA	1	0	transposon	LTR	micropia
11266	CAGACCUAUACAGAUUGGAACAU	1	0	transposon	LTR	McClintock
11267	UGCUACUCGAAUGUAUGCCCGAUAUAC	1	no match			
11268	UCAUCUGUGGAGGGGAUCCUCAA	1	no match			
11269	GUCUCAGAGAGCAAAUCGCCAUGU	1	0	transposon	LTR	gypsy3
11270	GUGUAAGUUACGAACCCUCUUCUUGC	1	0	transposon	LTR	opus
11271	UAAGAUUGGACCCAGUACACUGCCU	1	0	transposon	LINE	Doc
11272	AUGACAUACAGAACCAUGAAGGUGA	1	1	transposon	LTR	McClintock
11273	CCGGAUUACAAGAAGCCAUUCCA	1	0	transposon	LTR	gypsy3
11274	UAUUGUCCGACUACGGAAUGACCAUGAG	1	0	transposon	LINE	Cr1a
11275	AUCUCCUGUGAUAGGCAGAACACCCUG	1	0	repeat		trf
11276	CGCACUUUAAGCGUUCGUUGACGUUGA	1	0	transposon	LTR	HMS-Beagle
11277	UGCACUAAUAUAAGCCCUGACUUCCAA	1	2	transposon	LTR	G4
11278	UUGGGAAUGAUUGCGUUGAGAAA	1	0	transposon	LTR	mdg3
11279	GCCAAACUGAUGCGAUGGUAUUGUA	1	0	transposon	LINE	Doc2
11280	AUCCCCCAGGACUUGAUCGCAUCGA	1	2	transposon	LINE	G2
11281	UAGACAGAAAAUAAGGGACACGCCG	1	no match			
11282	GCCCCCAAACGACGUCGGCGAUACCCUAA	1	1	transposon	LINE	Quasimodo
11283	CAACCAAAAAUGACAAGCGCGAAA	1	0	transposon	LINE	Doc4
11284	AAUUGACGGAAGGGCACCACCAGGAGU	1	0	ncRNA		rRNA
11285	AAAAACCUGAAGAUGAUCCUGAGUCGCAU	1	2	transposon	LINE	Helena

11286	ACCCAAAGAACAAAGACGAAACAC	1	0	no annotation		
11287	ACCGAGAGAAGACUUUCACCCGAUGA	1	0	coding gene		CG12717-RA|CG12717
11288	AGGAGCCGGGCGUGGAAUGCGAGUGCCU	1	no match			
11289	GAGCUCGUUUGAUUUUGACCAGGAUU	1	0	transposon	LINE	X
11290	CACUAUUACACGAUAUACCUUGAG	1	0	transposon	LINE	Cr1a
11291	ACGGGGAAAACUUACCAGGUCCGAAC	1	no match			
11292	AAAGAGAGUAAUUUUGGCCGUCACCU	1	1	transposon	LINE	Cr1a
11293	CCGAAGAAAACAAAGCCUCCGCCA	1	0	transposon	Low_complexity	F-element
11294	GCAUCUUCCAAAGAACGAUCGAUGA	1	0	transposon	LTR	gypsy2
11295	CAAAUUCCGAAAAUGCGAGUUCCU	1	0	transposon	LTR	Rt1c
11296	UGCCAUCUACAAUGUCGUCCUGGG	1	0	coding gene		CG30104-RA|CG30104
11297	CUUGGACUACAUAUGGUUGAGGGUUGU	1	0	ncRNA		rRNA
11298	GUGAGGUCUCCGGACGUGAUCACUGUGU	1	1	ncRNA		rRNA
11299	UGUCUAAGUACACACGAAUUAAA	1	0	ncRNA		rRNA
11300	GUUUGACUGGGGCGGUACAUCUCU	1	0	ncRNA		rRNA
11301	CCACACUGGGACUGAGACACGGC	1	no match			
11302	GGGUCGAUGAAGAACGCAGCAAACUG	1	0	ncRNA		rRNA
11303	AGUGUAUGGAGCGUCAAUAUUCUAGU	1	0	ncRNA		rRNA
11304	AGCGUCGAACGAUUUCUGCCCACC	1	0	transposon	LTR	Quasimodo
11305	AGCUCGAACAAGCCCAGCAAGAGAAU	1	no match			
11306	AGAACCCUGACCUACGAGUCCGUCCAUCA	1	1	coding gene		CG6255-RA|CG6255
11307	ACCCUAGAACGCGCCCGAUCUCGUC	1	no match			
11308	GGCUCAACUGUGAGGGAUCCUCAA	1	no match			
11309	UGUUUCAUCGUUAGACGGCUCGGGCUCU	1	no match			
11310	UGUUUCUUCGUUAGACGGCUCGGGC	1	1	AT-chX		AT-chX-B
11311	GGAUCAAGGGCAACGAGUUCCUCUGC	1	0	coding gene		CG40947-RA|CG40947
11312	AAUCGAUGAGAUCAGGGCCUGGAGA	1	0	transposon	LINE	G
11313	CCCUACUGAUGAUGUGUUGUUGC	1	no match			
11314	ACCCUACUGAUGAUGUGUCUACGACC	1	no match			
11315	CAGAUCGACAAAGUACGAUGGGGA	1	1	no annotation		
11316	UAUUGCUUGUGGAACGUCGUUGC	1	0	no annotation		
11317	GCAACGAUACGUAGCCGACCUGAGAGGG	1	no match			
11318	AUGGCUCUUUCAGUCGCUUCCCGGGA	1	1	ncRNA		rRNA
11319	GCAUUUCAAAAACGCGAGUGGCAUCC	1	0	transposon	LTR	invader3
11320	UGUUUCAUCGUUAGACGGCUCUGG	1	1	AT-chX		AT-chX-B
11321	UAUUUCAUCGUUAGACGGCUCGGC	1	2	AT-chX		AT-chX-B
11322	CUCGACCGUGGCAACUCGAACCGAGGC	1	1	repeat		trf
11323	UAUUGGAGAUAGAGGCCCUUGGA	1	0	coding gene		CG7138-RA|r2d2
11324	GUCUCGAGAAUACCGAUCCAGUGG	1	0	no annotation		
11325	GGUCCUGAGAGAUGGGCCGAGGGA	1	no match			
11326	CAACCAGAGAUCUAGAGGGCCCAAGAAA	1	0	no annotation		
11327	UGCUCAUUCCGAACAUAAAUUGACAA	1	0	transposon	LINE	Doc
11328	GCUGAUGAUACCGCGAUACUCAGUCGC	1	0	transposon	LINE	Doc
11329	AAAUUCCGUAAAACGAAGCCCAAAAGUU	1	no match			
11330	CCCUUGAACCCAGGAGCUCUAGACC	1	no match			
11331	UGGGGAUGACCAGCAGGAGGUGGAGA	1	0	coding gene		CG15262-RA|CG15262
11332	GAGCCCAUGACCCUUGACAACAUCGU	1	no match			
11333	UUGACGGAAGGGCACCACCAGGAGU	1	0	ncRNA		rRNA
11334	GGGUUCCGGGGGGGAGUAUGGUUGCA	1	no match			
11335	CAACGAGUUCCUCUGCCGCGUGCCUAU	1	0	no annotation		
11336	CUGCCGAACGUAGACUGGUCGCGAGU	1	1	transposon	LTR	invader1
11337	UCCGACAGAAGCCCCGCUACCUCC	1	0	transposon	Low_complexity	F-element
11338	CUAAACGUAUUAACAGACACUAAA	1	0	transposon	DNA	transib2
11339	CGCAAAAGUUGACCGGACUUGAU	1	1	ncRNA		rRNA
11340	CAGGCAAGGUGUUGUAAGUGGUUGA	1	0	ncRNA		rRNA
11341	CGCGGAAUGAAGCAGUUACGUUUGGA	1	0	coding gene		CG18449-RA|CG18449
11342	AGCGUUGGAUUGUUCACCCAUGCAAGG	1	0	ncRNA		rRNA
11343	AAUAGACCUCUGUUCUGCUUUCAUUG	1	2	ncRNA		rRNA
11344	UCCAAGAACAACAACAACCAGAGCAA	1	0	transposon	Unknown	RR50960
11345	CUAAUACAUAUCGAUCAUUACGGCCCA	1	0	transposon	LTR	flea
11346	CACACAACAAAACUAUCACCCAGCUGU	1	1	transposon	Low_complexity	F-element
11347	AGAAACCGAACUUCAUACAGAUGCC	1	0	transposon	LTR	flea
11348	UGCGGAACAGGGCGAGGACUCCAAUG	1	0	coding gene		CG3980-RB|CG3980
11349	GCGUAACAUGGCCACACUGUCUCCUCCCGA	1	no match			
11350	UAACCAGAGAUCUAGAGGGCCCAA	1	1	no annotation		
11351	GCGAUCCACUGAAGCUUAUCCUUU	1	0	ncRNA		rRNA
11352	GCCUCUCAAAACGAAACCCGAUCAGA	1	0	transposon	LTR	gypsy
11353	ACUCGCCGAAUCCCGGGGUCGAGGGAGC	1	no match			
11354	AAUGCUAACAAGGCGCCGCCCGUGA	1	1	transposon	LINE	G6
11355	AAACGUGGAUGCUCGCUGUUGGCCUUU	1	no match			
11356	ACCACCCUGAUCGCGCCCGAUCUCGU	1	no match			
11357	UUGGGAACACCGCGUGUUGUUGGCC	1	0	ncRNA		5SrRNA
11358	CAAGAUAGGAAGGAAGAGUUAAAU	1	1	transposon	DNA	transib4
11359	CGUACUCACGCACUUUCCAUCGAUG	1	0	transposon	LINE	Circe
11360	CACAGCAACAUUGUCUGCAAGGGAC	1	0	transposon	LTR	Tirant
11361	CACGCAGUUACCGACCGUUGUGUCAU	1	0	transposon	LINE	Quasimodo
11362	CCGAUACAUAGAGCAACUUACACCAAC	1	1	transposon	Low_complexity	F-element
11363	AUUGUAGACGGUCUUACGGGA	1	0	transposon	DNA	S
11364	ACAUUCAUAACCGAGCGACUUUUUAA	1	2	transposon	LTR	GATE
11365	AACAUUCAUAACCGAGCGACUGUUCAA	1	1	transposon	LTR	GATE
11366	CCAUGGUUGCAACGGGUAACGGGGAAU	1	0	ncRNA		rRNA
11367	CUGAGGCGCUCGGUGUCGGUGAAGG	1	no match			
11368	GAUUCCUUGAGGAGCUGGAACCUGGU	1	0	transposon	LINE	G2

11369	UGAUAGACCUAUGCAGGGUUCCGAAUG	1	0	no annotation		
11370	AGCCUGAGAGGUGAUGCGUAGCCGA	1	no match			
11371	UCCUGAAAGUAGUUGGUGGGCACGUGG	1	0	Su(Ste)		Su(Ste)
11372	GAGUUCGAGACUCUUUUCCCGCUCC	1	no match			
11373	CUAAACCAUUCGUAGACGACCUGCU	1	no match			
11374	GGAUACGACGACGUAGACCAAG	1	no match			
11375	CUUGGCGUUAAGAGGCGAUGAAGGAC	1	no match			
11376	CACGACCCCCAGCUACGUGGCCUUC	1	no match			
11377	AAACGAGAAUAUACACAGGGGCUUA	1	1	transposon	LINE	Doc3
11378	ACCGGAGAAGACGUUGAAUGGAUC	1	0	no annotation		
11379	UGCACUAAUAGUAAGCCUGACGUCCGA	1	no match			
11380	UCUCUAAAUCUAUGGGUGGUGGUGCAUG	1	1	ncRNA		rRNA
11381	UUCCGAUAACGAACGAGACUCAAAU	1	0	ncRNA		rRNA
11382	UCCGAAGAUAACGAGGAGCAGAGCUGAU	1	1	transposon	LTR	297
11383	UUCGAGAACAAACGACACUGUUCGGC	1	1	transposon	LINE	G2
11384	CAACUGUGAGGUCAACUGUGAGGUCAA	1	no match			
11385	CAGCUGCCCGAAGUGCGGCGCGCCGGCCA	1	no match			
11386	GUAGUCGAGAUCAGCGUGCCGUGUUA	1	no match			
11387	UCAGCUACGGCAAUGCAGCUCGAGCCG	1	0	transposon	LTR	diver2
11388	UCCGAAGAUAACGAGGAGCAGAGGC	1	2	transposon	LTR	297
11389	GAUUCCGGAGAGGGAGCCUGGGA	1	1	ncRNA		rRNA
11390	UAGAAAGUGACCCUGUUGAUAAAUCGA	1	0	transposon	LTR	Idefix
11391	CAGGAAUGUCUAUCCGAAAGACGAGU	1	0	transposon	LTR	G4
11392	GCCAGGCCCGUAUAACGUUAAUGA	1	0	ncRNA		rRNA
11393	GGAACCAACCCAAAGAGUACUCACCGA	1	0	transposon	LINE	Doc2
11394	CCGUGUGGCGAGGAGUGCGGUUCUUU	1	no match			
11395	GCCUGACCGAGUAGUGUAGUGGGUG	1	no match			
11396	UAAUUGCCCGGGAUCUCGGUAGUGA	1	0	transposon	LTR	diver2
11397	ACAUGCCCUGGACCAACUAUACCACCA	1	1	coding gene		CG33106-RA|mask
11398	UUGUAGGUGUACUGACCCAGUCGGA	1	0	transposon	LINE	baggins
11399	GCCCGAUGUCCGCGAGAUUUGCCGC	1	no match			
11400	CAUGCAAUUAAAACAUGAACCUUAUG	1	0	ncRNA		rRNA
11401	CACGAUUGAGAACGGCUACCUGGC	1	0	coding gene		CG14446-RA|CG14446
11402	AUGCAUAGAUAAAGGAGACGCACCGUU	1	1	transposon	LTR	flea
11403	AUGGGCUAGAAGAGCAUGACAUAU	1	0	ncRNA		rRNA
11404	UUACUGACGAAUUUAGUUACAAGCA	1	0	no annotation		
11405	CAACGAGUCCUCUGCCGCGUGCCUA	1	no match			
11406	AGCAGGACGGUGGCCAUGGAAGU	1	2	ncRNA		rRNA
11407	AAAACCUGAUGAUGAUCCUGAGUCGC	1	1	transposon	LINE	Helena
11408	UACAAGCAAGACCACCACACCCGA	1	0	transposon	LTR	gypsy12
11409	AGCGAAUGAUUAGAGGCCUUAGG	1	0	ncRNA		rRNA
11410	AAAUAAUCAAAGACAAUCUCAGCCCGA	1	0	transposon	Low_complexity	F-element
11411	UGUUUCAUCGUUAGACGGCUCGGGAA	1	2	AT-chX		AT-chX-B
11412	ACGCCCGGUAGCACGAGUCCAUGCGC	1	0	transposon	LTR	ninja-Dsim-like
11413	ACUCGCCGAAUUCCGGGGCCGAGGGAGCG	1	no match			
11414	UAGACUAUGAAGAGACAUUUGCUCCU	1	0	transposon	LINE	Cr1a
11415	AGCUCUUCUAUACAACUCCGUCUUA	1	0	transposon	LINE	Doc
11416	ACCCUGAUGGAAGACCGAAACAGUUCU	1	0	ncRNA		rRNA
11417	UACGACGAGAACGAGAUCAGCGA	1	1	transposon	LTR	frogger
11418	AGCAAACCAAUCCCACAAACCCG	1	no match			
11419	ACUUACCAGGUCCGAACAUAA	1	0	ncRNA		rRNA
11420	UAGGAACGAAAGAGGCGAGGACA	1	0	coding gene		CG40271-RA|CG40271
11421	UUCGAUCAGAAUUGAUUUCGGCCCG	1	0	transposon	LINE	Cr1a
11422	CGAUUGAAUUAUUUAGUGAGGUCUCCG	1	0	ncRNA		rRNA
11423	AAAACCCUGAAGAUGAUCCUGAGUCGC	1	2	transposon	LINE	Helena
11424	GAGGUGUCCCAAGGCCAGCUCAGUGC	1	0	ncRNA		rRNA
11425	GCGUAUAACAACUACUUCUCGGUGAA	1	0	transposon	LTR	accord
11426	GCAUUACGAGACUCAAUGGCACGUCACAA	1	no match			
11427	ACGAGCGCACGUGUUAGGACCCGUAAG	1	no match			
11428	UAAACUCACAAUUAGGUGGGUCCCG	1	0	transposon	LINE	baggins
11429	CCCAAUUGUGUUUCGUCCUAAAGA	1	0	transposon	Low_complexity	F-element
11430	GAACCUUGAAGACUGAAGUGGGGAAG	1	no match			
11431	GUGCGAAGAGCGAUGAUACCGAAGA	1	0	transposon	LTR	Dm88
11432	AUCGAUAUUAUCCCAGCACUACGAAG	1	0	no annotation		
11433	AACUCCAGCAGAGACCGUACCCGAAAC	1	no match			
11434	CAAGACAGAAAAAUACCUGCGCAGGUGA	1	0	transposon	LINE	RR49783
11435	GCCGAACGUAGUCUCUCAAGAGCGAG	1	1	transposon	LTR	invader1
11436	GAGGACGAGGAUGUUAUGCCGCCGACG	1	0	coding gene		CG33935-RB|mei-217
11437	CAAUCCGAUACAUUUCACGCCGACAUC	1	0	transposon	LINE	INE-1
11438	GCAGCCGAGCUCAUUUUCCCUUG	1	1	transposon	LTR	copia
11439	UUGACUAUUAUUUGGAGAACCACA	1	0	transposon	LTR	opus
11440	GUAGGCAAUAUCCCAAGGUGCAAGA	1	0	transposon	LTR	gypsy3
11441	GCUUCCCGGGAAACCAAAGCUUUUGGG	1	0	ncRNA		rRNA
11442	UCCUGAUGACUCUUCGACAUAUAUA	1	0	transposon	LTR	Dm88
11443	UGUUUCAUCGUUAGACGGCUCGGGCAA	1	2	AT-chX		AT-chX-B
11444	AUAUGGCAAAUGUGAGAUGCAGUG	1	0	ncRNA		rRNA
11445	AGGGGCGAAAGACUAAUCGAACCAUCU	1	1	ncRNA		rRNA
11446	AUAGCUAAUUAACACAAUCCCGGG	1	0	ncRNA		rRNA
11447	ACAUGAAGCGAACCCGGGGAACUGAAA	1	no match			
11448	CCAUGGACCCCGUGGAGAAGGCUCUGC	1	0	coding gene		CG4264-RA|Hsc70-4
11449	UCAACUUUAACCCAAGAGACGUGACGA	1	0	transposon	DNA	transib4
11450	CGCCGAAUCCCGGGGCCGAGGGAGCG	1	no match			
11451	AACCAAGAUUAUCAACCCGUGUUA	1	1	transposon	LINE	Cr1a

11452	ACUGGGAGAAUAAACCCGAUGGACCCAAUG	1	no match			
11453	GGAUAGUCCCGAAGAAACCUAAAACC	1	2	transposon	LTR	opus
11454	UGCCGGUUCGAGUCCGGCCCCGCGCACC	1	no match			
11455	AAGGGAACGGGCUUGGCGGAAUCA	1	no match			
11456	GAGUAACCGAUCGUUCAAGUAUUGCCA	1	0	transposon	LTR	micropia
11457	CACCAAGAUUAUCAACUCGUGUUA	1	0	transposon	LINE	Cr1a
11458	AACCAUUUACAAAAUGUAUUCCGACCA	1	1	transposon	LINE	Doc
11459	UACUUGUUCCCCGGAUAGUUUAGUUUAC	1	no match			
11460	CAGUACGAUCCGUCUGCAAACUUCC	1	0	transposon	LTR	opus
11461	UCAACGCAUCAGAGGUGAUUUUUGA	1	0	transposon	LTR	rooA
11462	UGGGUGCGAGAGGCCCUGGGUUC	1	2	ncRNA		tRNA
11463	GAAGAUAUGAAACCGACUCAGGCAGU	1	0	transposon	LTR	Max
11464	UCAAGAAUAACAGGACCAUCCAA	1	0	no annotation		
11465	UGCGCUUGUGACGGCCCGUGAAAA	1	no match			
11466	AGAGAAUACCAAGGCGCUUGAGAGA	1	no match			
11467	CGCCCCUGGACGACCCACGUCCCUAUG	1	no match			
11468	CAGACUCCGACGUUGGCACCAGCAGCGCCUC	1	no match			
11469	CCGAAAGAAAGGGCAGACUAAUCCGU	1	0	transposon	LTR	ninja-Dsim-like
11470	CGCAGCGAUUUGCCAGAGAUCAU	1	2	coding gene		CG3338-RA|CG3338
11471	ACGACGAGAACGAGAUCAGCGCGA	1	0	transposon	LTR	frogger
11472	GAGGUUCGAAGGCGAUCAGAUACC	1	0	ncRNA		rRNA
11473	AGGCGUAGUCGAUGGAUAACCCAGUUG	1	no match			
11474	AGUUGGUGAGCGAUUUGUCUGGUU	1	no match			
11475	CGGCUCAUCUCAUCCUGGGGGCUGU	1	no match			
11476	CGAACCGAUUAAAAUUAAAGUGCA	1	no match			
11477	UCCAACGAGAGAGCGGAGAGCCAGCGU	1	0	transposon	Unknown	RR50960
11478	UCCAGGCCCAUCUGGUUGAACGUGU	1	0	Su(Ste)		Su(Ste)
11479	GAGGCGUUUGGAAGGCGGGCGA	1	0	coding gene		CG4434-RA|CG4434
11480	AAACCCAGAUUUGAGACGAAAGACCGAA	1	0	coding gene		CG17924-RA|Acp95EF
11481	UGUUAAAUUACAGAAGGAAUUCCUA	1	0	transposon	LTR	invader4
11482	CGUCUCAUAUGCGCACCCCACACACUU	1	no match			
11483	UUGAACGUGUCCUGAAAGUAGUUGGUG	1	0	Su(Ste)		Su(Ste)
11484	UACAAUGAUCUAUCGGGUUUUUUCU	1	no match			
11485	AUCUGAAGAGGUCGAUCCCAGCGUCA	1	0	coding gene		CG2093-RA|CG2093
11486	AGUUGGUGGAGCGAUUUGUCUGGUU	1	2	ncRNA		rRNA
11487	UGUGAAAAGAUCCCGUUUGAAUAUGA	1	0	transposon	DNA	S
11488	AACGCCUGGGACAUGCCAAGUUGGUGCCC	1	2	coding gene		CG31538-RA|CG31538
11489	GUGGCAUGUUUCCGGAUUCCCGC	1	0	ncRNA		alphagamma-element
11490	ACAACAUCCAGCGCUACGACUCC	1	0	transposon	LINE	Porto1
11491	AAUGGCACCAUGGACGAAUAUUGCAAC	1	0	transposon	LTR	HMS-Beagle
11492	AAGCCAGUUACCGAAGCUGCGGAUU	1	no match			
11493	CUCGCCGAAUCCCGGGCCGAGGAGUC	1	no match			
11494	ACCGAAGAAAACGAAGCCUCCGC	1	1	transposon	Low_complexity	F-element
11495	UGAAAAACGAGACGGAUGCGAUACC	1	0	coding gene		CG1763-RA|nod
11496	GCCUCAACUGUGAGGGGAUCCUCAA	1	no match			
11497	CAAUCGACGUGACCCAAAUGAAGCAUA	1	0	coding gene		CG11334-RA|CG11334
11498	GAGCGCACGUGUUAGGACCCGUAA	1	no match			
11499	ACAUACAUGAUGGCGGCCAACUAAA	1	0	transposon	LTR	frogger
11500	UGGUUCGAUUCUAGGAGAUGGCACCA	1	no match			
11501	AGACGACGAAAUCUACAGACUCAGCGU	1	1	transposon	LINE	G2
11502	AAAGCCAUGUGACGACGAGGCCCGUGC	1	0	coding gene		CG5762-RA|CG5762
11503	GGCACAAUUACCAGAAGAUGUUGACGC	1	0	transposon	LTR	roo
11504	GGAACCGGGCGGGAAGUUGGGGGGACC	1	no match			
11505	GGUGCCAGGCCCGUAUAACGUUAAUGAU	1	0	ncRNA		rRNA
11506	AACCCGUGCAUACUUUCGACUCCGU	1	0	transposon	LINE	accord2
11507	ACCGAAUUAAUCGCGAGGCUGC	1	0	transposon	LTR	mdg3
11508	UCUCAUCGUCGUAGAACAACCCCGA	1	1	Su(Ste)		Su(Ste)
11509	UGUCGUUGACGUCUAACCGAGGGCCGU	1	no match			
11510	UUGGGAACACCGCGUGUUGUUGGUCU	1	1	ncRNA		5SrRNA
11511	GACGUAGAAAAACUUGGAGACGGC	1	0	transposon	LTR	diver2
11512	UACCAAACGAAGUCCUACCAUAUGA	1	0	transposon	LTR	flea
11513	ACUACGACGACUCCGACUCUGACAGC	1	0	no annotation		
11514	UGAAGCGAACCCGGUGAACUGAA	1	no match			
11515	GCAGUCGUGGCCGAGCGGUUAAGGCUCU	1	no match			
11516	UUCCCGUGGAUAACGUGCCCGUGU	1	0	coding gene		CG31025-RA|CG31025
11517	AAAAGAAACGAAGACAAGAACAGCUCGA	1	0	transposon	LTR	invader6
11518	CGGGGCGUUCUAUAUAGUUAUGUAUG	1	1	ncRNA		rRNA
11519	UACAUCGAAAAGCGAGUUGACGUGUG	1	0	transposon	LTR	Max
11520	CAACAAAUCAUGAAAGAGCGCGA	1	0	transposon	LINE	F
11521	AAGCCGGAGGAAGGUGUGGAUGACG	1	no match			
11522	AUUCACACUCAAGAUGAUCGUGGCCA	1	1	transposon	LTR	297
11523	ACGCUGCCGGAGAGGGUGAUGGCCCCG	1	no match			
11524	AACUUCUUAGAGGGACAAGUGGC	1	no match			
11525	GAGUACUUCACCCAGACACUGGACG	1	1	Su(Ste)		Su(Ste)
11526	UGUGCUAGAAGACGAUUUUCGGGGU	1	1	transposon	LINE	Cr1a
11527	UGAGAUAUGCUAGCAGACCGCUCC	1	0	no annotation		
11528	GAGUCCAAGAGAUUUGCCCCCGAGGU	1	1	coding gene		CG18743-RA|Hsp70Ab
11529	UGAAGAUGAAAAUACUGUUAACCGA	1	0	no annotation		
11530	ACCACAAUCAUCACCAGACGGCCC	1	0	coding gene		CG10719-RA|brat
11531	AAGUUCAAUAUCCCAAAUUCGGCGUGA	1	0	no annotation		
11532	UUCAAGGUUACCCAGGAUGCUGUG	1	no match			
11533	GCAUUCAUAACGAGAUUGCGCAUUGCA	1	0	no annotation		
11534	GGCACGAGACCGAUAGUCAACAAGUACC	1	no match			

11535	UAACUAAGAACCGUGAGAUAUCGCU	1	0	no annotation		
11536	ACCCUGAACGCGCCCGAUCUCGCC	1	no match			
11537	GCUGAGAAGACGGUCGAACUUGACUU	1	no match			
11538	UCGGCCCUAACCCUUCUGUUGUGU	1	2	transposon	LINE	baggins
11539	CUUAGACAUCCAAUCUGUUGGCCA	1	1	transposon	LINE	F
11540	CACCCUGAACGCGCCGAUCUCGU	1	no match			
11541	AAAAGGGCAAAUGCUGACUUGAUCU	1	0	ncRNA		rRNA
11542	CAAGCUAGACUAUGUAUCCCCCGGGUA	1	no match			
11543	ACAACGCCAAGCACCCGAUUGUGAUAA	1	1	transposon	LTR	3S18
11544	GAGGCACUGAAACCAUCCGGGCGAG	1	0	coding gene		CG10616-RA|CG10616
11545	UUGGGAACACCGCGUGUUGUUGCCCU	1	1	ncRNA		5SrRNA
11546	UGCAAUCGAAAACGGGGAAAACUUCAUC	1	no match			
11547	GAGAAUCCGAGAACUUUGUUAGAAAA	1	0	transposon	LTR	HMS-Beagle
11548	AAGCCCGAAACAACGACUCGGUCGACA	1	1	transposon	LTR	gtwin
11549	ACUGAAUAACGAAUGGAUGAC	1	no match			
11550	UGCACCGAAAAUAUUUGCGAUGAAA	1	2	repeat		trf
11551	ACCCGGCCCGGACACGGACAGGAUUGA	1	no match			
11552	CACGCUGAGAACAUAAGAAAUCUAAA	1	0	transposon	LINE	HeT-A
11553	UGUUAUCGUAAGGCGACAUGGGUGCA	1	no match			
11554	UUUGCGAGAAAUUCGGUCACCGAGU	1	1	transposon	LTR	Dm88
11555	AGUCGGUCCUGAGAGAUGGGCGAGCGCC	1	no match			
11556	CGCUCACAGACUACCUAAGAGGCGC	1	1	transposon	LINE	Doc4
11557	UCGCCGCCCCGGGGCCGAGGGAG	1	no match			
11558	CCUUCCAGCAAGGACGAUUGAUGCUAC	1	no match			
11559	UGACGCGAUGUGAUUUCUGCCCAGUGCU	1	2	ncRNA		rRNA
11560	UAGAGCGUCCGACGAAUACCCUGGGGGG	1	no match			
11561	CAAACCAUCCCAACCAAACCCGUCAA	1	no match			
11562	UCAUCGAGAUAUUAAGGACUGGGGA	1	0	transposon	LTR	blood
11563	ACCCUGAACGCGCCCCGGAUCGUCGU	1	no match			
11564	CCAAGGUGAAGAGUCUCUAGU	1	0	ncRNA		rRNA
11565	UUGUAGAUCAAUCAUAUCACCGUGU	1	0	transposon	LINE	Cr1a
11566	AAAGCACCGUUUCCCGUCCGAUCAAC	1	no match			
11567	CCGAACUUGAGACCACAGCCGCU	1	0	no annotation		
11568	GGGGACUGAGGAGCGCAUUUCA	1	1	transposon	LTR	flea
11569	CACCAGGAGUGGAGCCUGCGGCUUAAU	1	0	ncRNA		rRNA
11570	AACCAAGCUAGAGUACAGAUCGA	1	0	transposon	LINE	jockey
11571	ACGGGAAAACUUACCAGGUCCGUAAC	1	no match			
11572	CGGUGACGGUUCCAAAGUUUAGC	1	0	coding gene		CG11807-RA|CG11807
11573	GAGCGUAGUCGAUGGGAAACGGGUUAA	1	no match			
11574	GCCCGAUGAACCUGAAUAUCCGUUAUG	1	0	ncRNA		rRNA
11575	AAAGAGAGUAAUUUUGGCCGUCAA	1	0	transposon	DNA	1360
11576	AAGAGGGCGUGAAACCGUUAAGGU	1	no match			
11577	ACCCCGAAAAUAAUUAAUAAACCCGG	1	0	transposon	LTR	invader3
11578	UCGAAACUCACUAACCUGCAAUUUGA	1	0	transposon	LINE	Rt1b
11579	AUCAAUCGCUGAUGAGACACCGGG	1	0	transposon	LTR	accord
11580	CCAAAGCUUUGGGCUCCGGGGGA	1	no match			
11581	UAUCCUGAGAAUCGAUUCGCAAGAAG	1	0	transposon	LINE	Cr1a
11582	UCCGAUGUACUAUCUGCCAAUCUC	1	0	transposon	LINE	Cr1a
11583	GGCGGUGGCGGCGGCGGCGGGGGC	1	2	transposon	LINE	R1
11584	UUUCUUGAAGAUGGCCCACAUAAAA	1	no match			
11585	GGAUCUCGACUUGUGACGCCAAAG	1	0	transposon	LINE	Doc
11586	AAAAACCUGAAGAUUAUCCUGAGUCGC	1	2	transposon	LINE	Helena
11587	GUCGCGAGAUCUACCAACACUAACCA	1	no match			
11588	GCCCCGUACUCUGUGGUGACCAACA	1	0	transposon	LINE	baggins
11589	UUGACCUGUAUUCGGAUAAUGGAAC	1	1	transposon	LTR	3S18
11590	UUAACAACGACAGGGGUACGUCAUU	1	0	transposon	LINE	Rt1b
11591	GCGGGACUUACGGGGAGAUGCUCAAGA	1	1	coding gene		CG31793-RA|CG31793
11592	AUACUUGUUCGAGCGAACAGUGGU	1	0	ncRNA		rRNA
11593	UUCUAGUCGUCGUGGGUUCGAGUCCCA	1	no match			
11594	GCUAGUUCCAAAAAAAUCAUUACCGA	1	1	transposon	DNA	NOF
11595	GGGUUCCGGGGAUCGAA	1	1	no annotation		
11596	ACCAACGGGAUACCUCAUACUCGAGUAA	1	0	transposon	LTR	micropia
11597	CGGAAUGGAAUCGAGCGCUGAGUAG	1	0	transposon	LINE	X
11598	GAGGAAUUUGAGCUGCGUUGACGC	1	0	transposon	LTR	ninja-Dsim-like
11599	CGAGUCUAGAGUUCCCAUCUGCAAGGGC	1	0	Su(Ste)		Su(Ste)
11600	GAGAUAGAUACAACUCACCCAGGGGGC	1	0	transposon	LINE	BS
11601	AACGGGAAAACUUACCAGGUCCG	1	1	ncRNA		rRNA
11602	CCAAAAGAUAAAACGUUUACGAGCA	1	no match			
11603	GAGAUAUGAAACCGACUCAGGCAG	1	1	transposon	LTR	Max
11604	GACCUCAUUCUCCGAGCCCGUCGU	1	0	no annotation		
11605	UAAUCCUUUAUAAUCCGAUCGUA	1	0	transposon	LTR	ninja-Dsim-like
11606	UGAGAGACAAAUUAGAUUUAAACA	1	0	transposon	LINE	G
11607	GUUGAAACUGAAGCACGAAUUGUC	1	0	transposon	LTR	Dm88
11608	CGAAAAUGCAGACCUAUUGUCAUCC	1	0	transposon	LTR	flea
11609	CGGCGGGGAUGGAGCGGUAUGA	1	0	no annotation		
11610	CCACCUGAACGCGCCCGAUCUCGUC	1	no match			
11611	AACUCUAAGCGUUGGAUCACUCGGUU	1	2	ncRNA		rRNA
11612	UUGAAUUAUUUAGUGAGGCCUCCGGA	1	0	ncRNA		rRNA
11613	UUCAGGGGGAUCUUUGAGUGAUGAAU	1	1	transposon	LINE	baggins
11614	CGCACACUCCAACCGCCUUCAGCA	1	0	transposon	DNA	1360
11615	CAAGCGAGUGGACUGGGCCAGGCA	1	0	transposon	LTR	ninja-Dsim-like
11616	ACGGAACCGCUAGCAGAUCGCGAAU	1	1	transposon	LTR	opus
11617	UCCAAAGAAGAUCAAUGGUGCGAGU	1	0	transposon	LINE	INE-1

11618	GAGCGUCCGACGAAUACCCUGGG	1	no match			
11619	GGGCAAAUAGACAAUUUGGAUGGC	1	0	transposon	LTR	HMS-Beagle
11620	GAAGAUUUCAUCGCGUUGACACA	1	0	transposon	LTR	micropia
11621	AGAUUCCGGAGAGGGAGCCUGAGAA	1	1	ncRNA		rRNA
11622	GGGCCCGGGUGGAGCCGCCGCAGGUGC	1	no match			
11623	UCAUGCGAGGCGAACCGGGUGAACUGA	1	no match			
11624	CAGACAGAGAUGGUGCAUUUUCAU	1	1	transposon	LTR	297
11625	GAUUGCGGACCGGAGAAGACGUUGA	1	0	no annotation		
11626	CAGGCCCAUCUGGUUGAACGUGUCCU	1	0	Su(Ste)		Su(Ste)
11627	CACGCAUUUACGCGAUUCACCGGACGU	1	0	transposon	LTR	diver
11628	UGAGCUGAGAGAGAUAUCGGUCC	1	0	transposon	LTR	opus
11629	GAAGCAAUCACCGAGAGCAUCAA	1	0	transposon	LINE	Rt1b
11630	AUGGGUCAGUCGGUCCUGAGAGAUGGGC	1	no match			
11631	CCCUACUGAUGAUGUGUUGUUGCC	1	no match			
11632	AUAAAUAAAAGGUGACCGGCAACGAC	1	0	no annotation		
11633	AGGUAGGGAGUUUGACUGGGGCGGU	1	0	ncRNA		rRNA
11634	AAAAGGUCCUCAACUGUGAGGCAA	1	no match			
11635	ACAACCUCCACGUUCGCUGAUGAUA	1	0	transposon	LINE	Doc
11636	UCCUAUGAGGCUACGGCAUGGGG	1	0	transposon	LINE	Cr1a
11637	GGUGAAAUUCUUGGACCGUCGU	1	0	ncRNA		rRNA
11638	AUGCCAAUCACGCCGUUUGAACCC	1	0	transposon	Simple_repeat	I
11639	GCAUUUAUGAGGUUGAGGUAGUACCCGA	1	1	transposon	LINE	G2
11640	GCUCUCGAAUGUAUGCCCGAUAUACG	1	0	transposon	LINE	R1-element
11641	CACGGGAAAACUUACCAGGUCCGAG	1	1	ncRNA		rRNA
11642	AAGAGGGACGCCGGGGGCAUUCGU	1	no match			
11643	GCGAGAAGCACGAGUUCUUCAUGG	1	0	coding gene		CG7228-RC|pes
11644	AACUUCGGGAUAAGGAUUGGCUCUGAA	1	0	ncRNA		rRNA
11645	CACAGAGUACGGGGCUUACUGAAGUCAU	1	1	transposon	LINE	baggins
11646	GGUAGGGAGUUUGACUGGGGCGGUAAA	1	1	ncRNA		rRNA
11647	AAGGGCAAAUGCUGACUUGAUCUCGGU	1	0	repeat		LSU-rRNA_Dme
11648	GGUCCAACUGUGAGGGGAUCCUCAA	1	no match			
11649	UCUCAUCGUCGUAGAACAAGCCCGAUUU	1	no match			
11650	CACGAUGAUCGAUGACGAACUACGAAG	1	0	transposon	LTR	ninja-Dsim-like
11651	GCAUCCUCAACUGUGAUGGGAUCCUCAA	1	no match			
11652	UCCAAUGAAGACGUUCAUUUCGGA	1	0	transposon	LTR	gypsy8
11653	AACGGGAUGGAAGUGACGCAGUU	1	no match			
11654	GAGGUCUCCGGACGUGAUCACUGUGU	1	1	ncRNA		rRNA
11655	AGGAUCUUCAACCUGUGAGGGGGAUCCUCAA	1	no match			
11656	GACUAGCUGGCCGAGUAUCUGAUGU	1	1	Su(Ste)		Su(Ste)
11657	UAGAUUGGACAGCCUCUGUAGUU	1	0	transposon	Low_complexity	F-element
11658	UACGGGUAGUACAACUUACAAUUGUGG	1	0	ncRNA		rRNA
11659	AGAUGUCCUGGGCUGCACGCGCGCU	1	0	ncRNA		rRNA
11660	ACGAGAUGUCCAGGGCGUGCACA	1	0	transposon	LTR	ninja-Dsim-like
11661	CGUGCUCAUCCUGCUGGCCGGACGCCAC	1	0	coding gene		CG11522-RA|RpL6
11662	GCCCGGCUACCUCUCGGUAGAGCAUGA	1	no match			
11663	GCAACAGGCAUACUGGGACUGGGCAGG	1	1	coding gene		CG10126-RA|CG10126
11664	ACGUACCUUGAGCAUAUAUGCUGUGACCC	1	0	ncRNA		rRNA
11665	AAGGAGUCUAACACUUGCGCGAGUC	1	no match			
11666	GCCGAAUCCCGGGGGCCGAGGGAGCGAG	1	no match			
11667	GCUUUAAUCACGAGCAAUGCGGUGACC	1	1	no annotation		
11668	CUACCGCAAGGCCGAGAAUGUGGAGA	1	0	coding gene		CG9422-RA|CG9422
11669	UAGAUCAGUCGAACAUUGUACUCUG	1	1	repeat		trf
11670	CUGACUUGUAUUCGGACAACGGAAC	1	1	transposon	LTR	3S18
11671	ACCCCGCGCGCUCCCUCUCGCCC	1	no match			
11672	GAAAGAACAAAAAGAGGACUUCGAG	1	0	no annotation		
11673	CUGGAAUGGCAGUGGUUACUGGG	1	0	coding gene		CG13695-RB|gk
11674	ACACCUGACAGAUACCCGAUGCAAGAUC	1	1	transposon	LTR	invader3
11675	GCAGAGACCGUACCCGAAACCGACACA	1	no match			
11676	UGCGUACAGAUCGAGUCAGAGAAGGAG	1	no match			
11677	GACGGGAAAACUUACCAGGUCCGAA	1	1	ncRNA		rRNA
11678	GCGAGAUGAGCUACACGGGUCGUGGU	1	0	coding gene		CG14801-RA|CG14801
11679	UGCCUCAACAUUUGGUGACCUCGACG	1	0	no annotation		
11680	UUCAUCGCUGGAGGACGCCCGGGC	1	0	transposon	LINE	R1-element
11681	UGCACUAAUAUAAGCCAUGACGUCCGA	1	0	transposon	LTR	G4
11682	UGUAACUACCGAAGGCAUAAAGCCCAA	1	0	transposon	LTR	Idefix
11683	CAUUAACGUAAGGGGAGUACUUGG	1	0	transposon	LINE	G2
11684	AUGGAACGAAAUUGGCCUGAUUAGCGG	1	1	transposon	LTR	GATE
11685	UUAAAGGUUACCCCAGGAUGCUGU	1	1	repeat		trf
11686	UGAACAUCGAGAAUACUGCACAGC	1	0	transposon	LINE	Doc
11687	GGGCAAAUGACAAUUUGGAUGGC	1	no match			
11688	ACCAAGAUCAUCAACCCGUGUUUA	1	1	transposon	LINE	Cr1a
11689	AAAUACAUGCCGACGGGCGCUGACCCCC	1	no match			
11690	UAAUGACCAGCGCUAACGAUCCCU	1	0	repeat		SAR_DM
11691	AAACUGAAUAAAACGAAAUGGAUGACACA	1	no match			
11692	AACGAUCUCGAGGAAUACGGCCUUA	1	0	transposon	LTR	ninja-Dsim-like
11693	CCUGAACGCGCCGAUCUCGU	1	no match			
11694	AGCGACGUCGCUUUUUGAUCCUUCGA	1	0	ncRNA		rRNA
11695	GCGAGAAUAAACGAAAUACAGUGGGA	1	0	transposon	LTR	diver2
11696	AAGAAAACUAAUCCGAGUAUGCAGU	1	0	transposon	LINE	Doc
11697	AUGUCGAGCUCCCAGAACAACAACA	1	no match			
11698	UCCGUGAUAGUUUAAUGGUCAGAAUGG	1	no match			
11699	AGGAACAGCCCGAGGACUUCGUGAUUGC	1	no match			
11700	GGCGGCAAUGGCACAGCAUCCUGGGG	1	0	repeat		trf

11701	AGACCGAAAGCAUCGAGGUGGCGAUCC	1	0	coding gene		CG5094-RA|Sgt
11702	AAUAGGCUCCUCUCAGUCGCUUCCCGGGA	1	no match			
11703	AACAUAACGACACUAGUGCAGCCACAG	1	0	transposon	LINE	G2
11704	CAACGAGUUCCUCUGCCGCGUGCC	1	0	coding gene		CG40719-RA|CG40719
11705	ACCCCCGUCAGAAGAUGCACUUUAAG	1	0	coding gene		CG3836-RA|stwl
11706	UCAUCAUCAUCAAUCACCGGCCAG	1	1	transposon	LTR	springer
11707	CGUAACUUCGGGAUAAGGAUUGGCU	1	0	ncRNA		rRNA
11708	AGAUCCCCGAAUCCGGAUUGGCGGAGA	1	no match			
11709	CGUUCACUUCGCCGAGGACGUUCAGGCUC	1	0	coding gene		CG18628-RA|CG18628
11710	AUACUAAUUGCCCGUGAGGCUUGACC	1	no match			
11711	GGUGCCAGGCCCGUAUAACGUUAAUGA	1	0	ncRNA		rRNA
11712	GUCUGGUUAAUUCCGAUAACGAACGAGU	1	1	ncRNA		rRNA
11713	UCGAUUCCGGAGAGGGAGCCUGA	1	0	ncRNA		rRNA
11714	GAUCAUCAACGAGCCCACGGCGGCCGCC	1	no match			
11715	UCGAGAUGGUACCGGUUCCGUUGGACA	1	no match			
11716	GAUGCGGGGCGAACGACAGGUGGUGG	1	no match			
11717	UACGUCCCUGCCCUUUGUACACAC	1	0	ncRNA		rRNA
11718	GCCUUACGAAACGAAACCCGAUCAAAU	1	1	transposon	LTR	gypsy
11719	AGACCGAACAAAAUACAACGUCGACAC	1	0	transposon	LTR	McClintock
11720	ACCCAUUACAUCCUAUACGGUCUUUU	1	0	transposon	LTR	GATE
11721	ACCUGGAACUGACCAAGGAUGAGGAGA	1	0	coding gene		CG4836-RA|CG4836
11722	AACGGGGAAUCAGGGUUCGAUU	1	0	ncRNA		rRNA
11723	CACGACCCCCAGCUACGUGGCCUUCACCGU	1	no match			
11724	CAAUCCGAUACAUUUCACGCCGAC	1	0	transposon	LINE	INE-1
11725	GGAAAACCUGAAGAUGAUCCUGAG	1	0	transposon	LINE	Helena
11726	CCAGGACGUUGAUAGGUCAGGUGUGG	1	no match			
11727	GAAGUUUAGCUAUCCCGGGGGC	1	no match			
11728	ACCCUGACGCGCCCGAUCUCGU	1	no match			
11729	UUCUUGAGAAUUUGGACGCCGUU	1	0	transposon	LINE	Cr1a
11730	UCCUACGAGAUGGAAGCUUACAUCG	1	0	no annotation		
11731	GUUAAUUCCGAUAACGAACGAGACUCA	1	0	ncRNA		rRNA
11732	CAAAGGCAUGCCCGAUGUCCGCGAAA	1	1	transposon	LTR	opus
11733	UUCAAACUAAUUGGUUCUCCGCCC	1	no match			
11734	AAGCUCGAGAAUCUCUGAAACGGA	1	0	transposon	LTR	ninja-Dsim-like
11735	ACGAUGAUCGAUGACGAACUACGAAGG	1	0	transposon	LTR	ninja-Dsim-like
11736	UACUCGAGACACCACGGUGAAGUGG	1	no match			
11737	UGAAAAUCCGGGGGAGAGGGUGU	1	no match			
11738	GACUUAACGACCGAGAAGUUUCUUGC	1	0	transposon	LTR	GATE
11739	CCCGAAAAAAUCCCCCGGCUACGACC	1	0	transposon	LINE	FW
11740	AAAGUCUUUGGGUUCUGGGGGGA	1	no match			
11741	GUCACCUACAGGUCCCGAUACGAGCU	1	1	transposon	LINE	X
11742	CCAAAAACUCGUAAGCACUCCAGCGC	1	2	transposon	LTR	Stalker4
11743	CCCUCAUCCUACAGACCAAUCAG	1	0	transposon	LINE	Doc2
11744	UCUCGUCGUCGUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
11745	AACCGGACAAGAUACCCCAGGUGG	1	0	coding gene		CG7907-RB|CG7907
11746	AACGAGCGCGACCCUUGUCCUUAGUUAC	1	no match			
11747	CUGCCAUCUUGUCGUU	1	1	no annotation		
11748	GAGUUCCUGAAACCGGUCGCAUGCAAGC	1	no match			
11749	UGCGGCCGUGUGGCGGCUUCCGAAAGCCU	1	0	transposon	LTR	GATE
11750	AGCUCCAACAGUUGGCGCUUCAGGCUGCUCC	1	0	no annotation		
11751	ACUGACCCAAAAAGUCAAUAUUAUUGCA	1	0	transposon	LTR	rooA
11752	GACCCAGGAUGACCGGUCGCUCUCUG	1	no match			
11753	AGCUUCAUCCAAGCGAACUGUGCCCC	1	1	transposon	LINE	R1-element
11754	AGCUUUAAUCACGAGCAAUGCGGUGA	1	1	no annotation		
11755	CGAGAGUGGACGGGAGCGGCGGGGGC	1	no match			
11756	GACGAAACGCAACAACAAAAUGGAC	1	0	transposon	LINE	G2
11757	GACAAAACACAUCCCGGGAAAUGAGC	1	no match			
11758	UAGUGACGAACCCGGCACCGGUGACA	1	0	coding gene		CG30020-RA|CG30020
11759	AGGAAAACGACAACGACGCAAGA	1	0	coding gene		CG41053-RA|CG41053
11760	CAGUCCGAAACACACCCCCGAUAAA	1	2	no annotation		
11761	GACUCUAUUACAAUACUACGACGAGGU	1	0	transposon	LTR	gypsy6
11762	AAGAUUGACGGAAGGUCACCACC	1	no match			
11763	CAGGCCCGUAUAACGUUAAUGAU	1	0	ncRNA		rRNA
11764	CGUUUCAUGAGGAAUGGGCCCAG	1	0	transposon	LINE	G2
11765	CACUAUCGAAACGACGGAUAAGCCCC	1	0	transposon	LTR	gypsy6
11766	AUGAGACUACAAACGGAUUGCGGACCG	1	1	no annotation		
11767	AGCCCCCGCGUGAGACCGUACCCGAAAC	1	no match			
11768	AUCCUUAACUGUGAGGGGAUCCUCAA	1	no match			
11769	GGGAUAAUGAAAUAAGACCUCUGUU	1	0	ncRNA		rRNA
11770	ACUCGCCGAAUCCCCGGGGGCCGAGGGA	1	no match			
11771	GCAAACCCAAAAAGACAAUACACGAC	1	0	repeat		trf
11772	UGUGAGGGGGAUCCUCAA	1	2	coding gene		CG3339-RC|CG3339
11773	CGUCCAAAAAACCCGUGCUUCUGGC	1	no match			
11774	CAGACUAAGACGUUCUCCACGUACA	1	1	coding gene		CG31359-RA|Hsp70Bb
11775	AGCGGGAAUGUGCACUUUUGUCAU	1	0	coding gene		CG14224-RA|Ubqn
11776	UGUGCUAGAAGACGAUUUUCGGG	1	0	transposon	LINE	Cr1a
11777	UUAAAACAGCAGGACGGUGAUCAU	1	0	ncRNA		rRNA
11778	UCGAGAAAAUUGACGAAUUGCUG	1	no match			
11779	AAAAGGACAAUUUUUGGCCGUCAC	1	no match			
11780	GCCGAAAGACCAAGGGUUCCUGCGUA	1	no match			
11781	AGGAACCUGCGGAAGGAUCAUUA	1	2	ncRNA		rRNA
11782	UGUGUUUCCCGGCGUACUAUACAGCC	1	no match			
11783	UGUUUCAGAGCGUUGGCCAGUUU	1	0	transposon	Low_complexity	F-element

11784	ACAAGCAACGCAUCCAAUUACCGAA	1	0	coding gene		CG11076-RA|CG11076
11785	ACAUCAAUCGCUGAUGAGACACCGGGA	1	0	transposon	LTR	accord
11786	GACCAGGACGUGACCCAAAAACGAUGCC	1	0	coding gene		CG9320-RA|CG9320
11787	CAAAACAACAUCCCGGGAAAUGAAAAA	1	no match			
11788	AGGUGUCCCAAGGCCAGCUCAGGGCGG	1	1	ncRNA		rRNA
11789	UCCGAUUUCAAGUAAUAGACGAACU	1	0	repeat		RSP
11790	CGGCGGAGGCGUAUCUGGGCGAGA	1	0	coding gene		CG31359-RA|Hsp70Bb
11791	AGAUCCCCGAAUCCGGAGUGGCGGAGU	1	no match			
11792	GCGUAAGUUGCUAGAGGUCGAUACUGA	1	0	transposon	LINE	accord2
11793	AACGAGCUGGUGCGAACCAAGACCCUG	1	1	coding gene		CG7808-RC|RpS8
11794	GGUAGAGAUAGGCCGAUAGCGUACAUU	1	0	transposon	LTR	opus
11795	ACUCGCUGAAUCCCGGGGCCGAGGGA	1	no match			
11796	CAAAGAGACAGAACUCAGUAGCAGCC	1	0	transposon	LINE	Cr1a
11797	UGACAGUUCACGAUUGAUUCCCCAC	1	0	transposon	LTR	opus
11798	CAUCUGUGGAGGACAGCAUGGAA	1	0	transposon	LTR	ninja-Dsim-like
11799	AUAUCAGUUAUGGUUCCUUAGAUC	1	0	ncRNA		rRNA
11800	UUCGCAAAAGUUGACCGAACUUGAUUA	1	0	ncRNA		rRNA
11801	GGAGCCGGGCGUGGAAUGUGAGUGCC	1	no match			
11802	GAAGACGAAAACCAAACUGACAAUGGGA	1	0	transposon	LINE	Porto1
11803	ACCCGGGGCGUUCUAUAUAGUUAUGUAU	1	1	ncRNA		rRNA
11804	CGCAAACCCAAAAGACAAUACACGAC	1	no match			
11805	ACCCUGAACGCGGCCGAUCUCGU	1	no match			
11806	AUUGCCAGGUAGGGAGUUUGACUG	1	0	ncRNA		rRNA
11807	CAAGUCCUCGCGCGAAAGAAUACCAA	1	no match			
11808	UCCGAUUUCAAGAACCAGACAAACAGA	1	0	repeat		RSP
11809	AUGCAGAUCGUAUGGUCUUGUACCGACG	1	0	ncRNA		rRNA
11810	AUGCAUGUCUAAGUACACACGA	1	0	ncRNA		rRNA
11811	AACUCUAAGCGGUGGAUCACUCGGC	1	0	ncRNA		rRNA
11812	ACGAGUGUUAUUGUGUGCCGGU	1	1	ncRNA		rRNA
11813	CGGACAGAGACGGCGCAUUUUCCC	1	1	transposon	LTR	McClintock
11814	ACCCUGAACGCGUCCCGAUCUCGU	1	no match			
11815	CAACAUCUCGGCGCAGUGCGGAUC	1	0	no annotation		
11816	GCCGAAUUAAUCGCGAGGCUGCAAGAC	1	0	transposon	LTR	mdg3
11817	UCGGUCGGGCUGGGGCGCGAAGCGGGG	1	no match			
11818	UGACGCCUUGCGUGUUACGGUUG	1	0	ncRNA		rRNA
11819	AUCGAGAAGCGAACCCCGGCGAACUGA	1	no match			
11820	UGCUCUCGAAUGUAUGCUCGAUA	1	1	transposon	LINE	R1-element
11821	AUUGACUUUUGGAAACUAUAUGACCU	1	0	transposon	LINE	Rt1b
11822	CCCGAAAACAGAUGAGGUAGCG	1	0	transposon	LINE	F
11823	CAUAGUGAUCCGGUGGUUCUGUAU	1	no match			
11824	GAUGUCGGCUCUUCCUAUCAUUGUGA	1	0	ncRNA		rRNA
11825	CCCAAGAGAAAAAUACGUGAUGGACU	1	0	transposon	LTR	Idefix
11826	CACAACUUGACUCAUCCAACUGGA	1	0	no annotation		
11827	AUAAGCAAAAUGCAGAACACCGGAGA	1	0	transposon	LINE	Doc4
11828	ACGAACUUGAGACCGCCCCUGGA	1	1	Su(Ste)		Su(Ste)
11829	AUGUGCUAGAAGACGAUUUCGGACCGA	1	2	transposon	DNA	1360
11830	CGGCGAGCGAAAAGAAAACAGUUCAGC	1	0	ncRNA		rRNA
11831	UAACCAUGAGACCGAUAGUAAAC	1	0	ncRNA		rRNA
11832	CAAUCCGAUACAUUUCACGCCGACAU	1	0	transposon	LINE	INE-1
11833	GCUGCAAUUGUUUGCCGUGGGACUAC	1	1	transposon	LINE	baggins
11834	AGAACAUAUCAGGGGCCUUCGCAGA	1	2	transposon	LTR	HMS-Beagle
11835	AUCCAACUAAAACUGUUCAGCGCCGA	1	0	transposon	LTR	micropia
11836	AAGGGAAGAGCCCAGCGCCGAAUCCCC	1	no match			
11837	UCUCAUCGUCGUAGAACUAGCCCGAG	1	0	Su(Ste)		Su(Ste)
11838	AACCACGAGAGGUACAUUUUUGCAGCU	1	1	transposon	LTR	springer
11839	GAGCAAGAUGGACGGGCUAAAGCGGG	1	0	coding gene		CG5417-RB|Srp14
11840	UCGAAACCCACGCAUCUGUGGCAGGC	1	0	transposon	LINE	R1-element
11841	UACGGUUGUUUCGCAAAAGUUUACCGAA	1	1	ncRNA		rRNA
11842	GACUACACAUCGAGAGGUCCCCGGUUCA	1	2	ncRNA		tRNA
11843	CGAAGCUAAAACGCUGGAAGGCUCGU	1	0	transposon	LTR	opus
11844	CGGUGAUCGAUGACGAACUACGAA	1	1	transposon	LTR	ninja-Dsim-like
11845	GCUCUCGAAUGUAUGCCCGAUAUU	1	1	transposon	LINE	R1-element
11846	AACCGUGACGGGACGUUACGCUGACGGG	1	no match			
11847	CUCCCCCGCCCCACCCCGCGCGCGCCGCUC	1	no match			
11848	ACUCGCCGAAUCCCGGGCCGAGGGAGUC	1	no match			
11849	AUUCACCAAGCGUUGGAUUGUUC	1	0	ncRNA		rRNA
11850	GACGUAGCAGCCCGGGCGUCCUCCAGC	1	0	transposon	LINE	R1-element
11851	CCGAAUACGAAGAACAGGCAGAA	1	0	transposon	LTR	gypsy12
11852	GGGCUCCGGGGGAAGUAUGGUUGCAAA	1	0	ncRNA		rRNA
11853	GAUGGGGCCGCUUGGGAACACCGC	1	no match			
11854	UCUCACUCGUCGUAGAACAAGCCCG	1	no match			
11855	ACACUCAUCAACCAAAGGACGUAGGA	1	0	no annotation		
11856	UUGUAGAUCGACCAUAUCACCGUGU	1	0	transposon	LINE	Cr1a
11857	ACGCGCGGCAGGACGAAAAGACCCCG	1	no match			
11858	CAUCGAGCUCAAUUGCAUCCCUCGCA	1	0	no annotation		
11859	AGACCAGACGUAAAUUGCAAAUC	1	0	coding gene		CG11856-RA|Nup358
11860	UAAUGUCCCUUAAAUCAGCACUCGU	1	0	no annotation		
11861	CGAGUAAUAUGAGCGCCCAGAGU	1	0	transposon	LTR	opus
11862	CCAAGUCCUCGCGCGAAAGAAUAUC	1	no match			
11863	AUCCGGACAAACACACAAGACCCGAUC	1	0	transposon	LTR	opus
11864	CAUCCUGAACGCGCCCGAUCUCGU	1	no match			
11865	UUAUUAACGAAACCUGAGACCGACAU	1	1	transposon	LTR	297
11866	UCACCGGGAAGCACACUGAUGCCU	1	0	transposon	LINE	G2

11867	CGCUCACAGACUACCUAAAAGGCGCA	1	0	transposon	LINE	Doc4
11868	ACAAACAGUACACCUACUAGCCCAUG	1	0	transposon	LINE	baggins
11869	AGAACUGGCGCUGUGGGAUGAACC	1	0	ncRNA		rRNA
11870	CAAUCUCAGAUAAAAACCCGAACG	1	0	transposon	LTR	297
11871	GAAUCUGAUAUAGAGAGCUGCACGUG	1	0	transposon	Low_complexity	F-element
11872	UACUUAACACCCUGUACUCAGCAUC	1	0	transposon	LTR	gypsy
11873	ACUCGGCUCGUGCGUCGAUGAAGAACGC	1	2	ncRNA		rRNA
11874	AAAGGGACCCAAAUUGGAGGGCAAGUCU	1	no match			
11875	GGACGGCGGAGCGAGCGCACGGGGUC	1	no match			
11876	UGAGCAAUAACAGGUCUGUGAUGCC	1	0	ncRNA		rRNA
11877	UAGAUGGUAGCCCAGGCAUUUGUGU	1	0	transposon	LINE	Rt1b
11878	AUCGCAAGAUCGUUAUAUUGGUUGA	1	0	ncRNA		rRNA
11879	UGCUCUCGAAUGCUUGCCCGAU	1	0	transposon	LINE	R1-element
11880	AACUGGAAAAACUGACGCAUAAG	1	0	transposon	LINE	G2
11881	CGACCACGUUGUGCUGCCACCGAGA	1	0	transposon	LTR	ninja-Dsim-like
11882	GGAAUUCUGAGAAAUACUCCAGCGACC	1	1	transposon	LTR	gypsy3
11883	ACCAAAAUCGCCGGGAUGAGGCCUACGA	1	0	coding gene		CG5111-RA|CG5111
11884	CCCCAAAACACACAUAAGCUACUGAA	1	0	transposon	LINE	Doc
11885	CGCGGGUCAACGGCGGGAGUAACUAUGA	1	0	ncRNA		rRNA
11886	UGUUUCAUCGUUAGGCGGCUCGGG	1	1	AT-chX		AT-chX-B
11887	AUGUUAAACAAAACCGUUAAAUCAGUAA	1	0	transposon	LTR	Idefix
11888	AAGUCGCGGGGCAGCAUGGGGGUUGC	1	no match			
11889	GCCGACCCCGAUUUUAAUAAACCGA	1	0	transposon	LTR	roo
11890	GACGUGGCGACCCGCUGAAUUU	1	no match			
11891	CGCAACGCAUUCCGAGGAUCUUCUCU	1	0	transposon	LTR	ninja-Dsim-like
11892	ACUUCAUAACCGAGCGACUGUUC	1	2	transposon	LTR	GATE
11893	GGCGGGCGCGGGACAUGUGGCGUA	1	no match			
11894	GUGAGGCCUCCGGACGUGAUCACUGUGAC	1	1	ncRNA		rRNA
11895	GACAAUCGAACGAAGUCAUAGAACA	1	0	transposon	LTR	gypsy8
11896	CACAAAGUUAUUCUUCUACCGAUGG	1	0	transposon	LTR	Rt1c
11897	UCAAAAUGGAAGAGACCCUGCGUG	1	0	transposon	LTR	opus
11898	GGUCCCUGUGGAGGUUGCUGUAGUCC	1	0	coding gene		CG31740-RA|CG31740
11899	UGAUCCAUAAAAACUGUACACAGACG	1	no match			
11900	CUCGAAGAACGGCCAGACCCGCGAGC	1	0	coding gene		CG1873-RA|Ef1alpha100E
11901	AGACCCAGUGCCGACGCGUCCGUUGUC	1	0	transposon	LTR	gypsy7
11902	UCGGCCUGGAUCACGGACAUUUCGGA	1	0	transposon	LTR	ninja-Dsim-like
11903	AAAGCACUCGACGGUAACCGAUGGU	1	0	transposon	LTR	Quasimodo
11904	ACCGAUAUGGUUGGACCACAAUCUUU	1	0	ncRNA		7SLRNA-RA
11905	UCCUAAACUACGGGAGAUAUUUGGA	1	0	transposon	DNA	hopper
11906	CGCCCGUCCCCGCCCCUUGCCUCUCGGC	1	no match			
11907	ACCACCCUGAGCGCGCCCGAUCUCGU	1	no match			
11908	CAAACCAAUCCCAACCAGACCCGUCA	1	1	transposon	LINE	G2
11909	AGAGGUGAAAUUCUGGACCGGCGC	1	no match			
11910	ACUCAACACGAAGACCCAGUAUACCGU	1	0	transposon	LTR	Idefix
11911	AGCAACAUGACCCAGGUUGCCUACCAGA	1	0	transposon	LTR	McClintock
11912	ACUCGCCGAAUCCCGGGGCCGAGGGC	1	no match			
11913	GAACCGAAAAAUCACAAUUUGCC	1	1	transposon	LTR	McClintock
11914	CCUCCACUAUGGCCAUGCGGUCGCUG	1	no match			
11915	CGGAUGUCUAUGAGAUGAACGUAAU	1	0	transposon	LTR	ninja-Dsim-like
11916	GUUGCUUGAUAGUGCAGCACUAAGUGG	1	0	ncRNA		rRNA
11917	ACCCCCUGGGACGCCGAGUUCCUGUC	1	0	coding gene		CG12227-RA|skpF
11918	GCGAAUGGCAUCCGGAUGAGAAA	1	0	transposon	LTR	flea
11919	CCCACCUGAAUCUCCGCAACCCAGUA	1	0	transposon	LINE	G2
11920	UGAACGCGCCCGAUCUCGUCUGAUCUC	1	no match			
11921	CAACCAGAGAGCCUACCAACGUGG	1	0	transposon	LTR	ninja-Dsim-like
11922	UUCCUUAGAUCGUUAACAGUUACUU	1	0	ncRNA		rRNA
11923	CUCGCAAGAAAACUAAUCCGAGUAUGC	1	0	transposon	LINE	Doc
11924	UCGUGAUGUUCUUGGCCUUGCCCGU	1	0	no annotation		
11925	CCGAGAGAAGUACGACCGGGUGCUGAA	1	1	coding gene		CG7556-RA|CG7556
11926	AGUCUAACACGUGCGCGAGUCGGGGC	1	no match			
11927	AAGAGGUACGGCCGGGGGCAUUCG	1	no match			
11928	CCCAGGACUGGAUCGCAUCGACGCACA	1	2	transposon	LINE	G2
11929	UUCACAUGCAAACAUCGACCAGGAG	1	0	transposon	LTR	flea
11930	CUCCAACUGAAGGAGUACGAAUGAUCG	1	0	transposon	LINE	R1-element
11931	AGGUGGGGGAGGUUUGACUGGGCGGGUA	1	no match			
11932	CUGGCGAAUAAAAGUAAACGAGCAAAA	1	0	transposon	Low_complexity	F-element
11933	GCAAACGUGAUGUUACGGUGCCCAA	1	2	ncRNA		rRNA
11934	AAAGUAAAAUGUAUCCGGAGCGA	1	2	transposon	LTR	297
11935	UGUUCAUCGUUAGACGGCUCGGGCGU	1	no match			
11936	AAAGCCCCUGGAGAUACAACUAGCCAU	1	0	coding gene		CG32300-RB|oxt
11937	GACGAGAUAAAGCGAAAACCGGCAA	1	0	transposon	LINE	G6
11938	AUCGAAUGGAGUGCCAGAACCCGUUAG	1	1	transposon	LINE	R1-element
11939	ACGAGCCCGAAUUCGACUACCUCAA	1	0	coding gene		CG6235-RA|tws
11940	ACAUUACAUAACCGAGCGACUUUCAA	1	no match			
11941	CAAGAUAUGACCAAAGCAAAGACGC	1	1	transposon	LINE	Cr1a
11942	AGGUCUCCGGACGUGAUCACUGU	1	0	ncRNA		rRNA
11943	AAGAACAACAGUGCAGACGGCGC	1	0	transposon	DNA	H
11944	CCAGAAUCUCCGCAAAGCCUGAGGA	1	0	transposon	LTR	diver2
11945	UGCUCUCGAAUGUAUGCCCGUAUAUA	1	no match			
11946	GACCGGCUCCGGGACGGCUGGGAAGGCC	1	no match			
11947	UCAUAAUAAGUUCCCCGACGUGCCGG	1	0	Su(Ste)		Su(Ste)
11948	UUGGGUUCCGGGGGGAGUAUGGUUG	1	2	ncRNA		rRNA
11949	AGAACCGACGUGCUAUCAUUCCGUUCC	1	0	coding gene		CG7840-RA|CG7840

11950	AGCCCGUGGACGGUGUGAGGCCGGUAG	1	no match			
11951	UCCAGUAUUAUGACGAGGUUGAGA	1	0	transposon	LTR	gypsy3
11952	AAGAUAUUGAGCGACUUUCCCCGUAG	1	1	transposon	LTR	3S18
11953	CAAUUCAACGAGGACCGCUAUGAGA	1	0	transposon	LTR	flea
11954	AAGAUGUCCGGACAGCUGGGCCCGUAAA	1	2	Su(Ste)		Su(Ste)
11955	CCGAGAGGUCCGGGUAAACCGCUG	1	0	ncRNA		rRNA
11956	GUCCCGCAACGAGCGCAACCCUCGCCA	1	no match			
11957	UGUUUCAUCGUUAGACCGCUCGGGCG	1	1	AT-chX		AT-chX-B
11958	GGUGGGCACGUGGCAGAGGAACUCGU	1	0	Su(Ste)		Su(Ste)
11959	CAUACGGUUACUGGAGUCGACGGAAA	1	1	transposon	LINE	R1-element
11960	AGCACGGGAGACACACGGGGGGUGCU	1	no match			
11961	CGGAGAGGGAGCCUGAGAAACA	1	1	ncRNA		rRNA
11962	GGCACAAUUACCAGAAGAUGUUGACGCA	1	0	transposon	LTR	roo
11963	CGGAGCCGGACGACCUUGACCUCU	1	0	transposon	LTR	diver2
11964	ACGAGCGCACGUGUUAGGACCGAAA	1	no match			
11965	GAAACCAUCAGACGCCUCCAGAGGAC	1	1	transposon	LINE	G
11966	AUAAUGAGAUUGAGCAAUAACAGGUC	1	0	ncRNA		rRNA
11967	ACGUAGGGUGCGAGCGUUGUCCGGAUU	1	no match			
11968	CACAGAGUACGGGGGCUUACUGAAGUCAC	1	no match			
11969	UCGUCAUCGUCGUAGAACAGCCCGA	1	no match			
11970	CGAACACAUCCGGUCGAUCCAGCAAA	1	0	coding gene		CG15258-RA|CG15258
11971	CUGGAACUAAGAUUGGCUUGCCAAA	1	0	transposon	LINE	INE-1
11972	CAAUACCGUAUUGUGGGAGACCAG	1	0	transposon	LINE	accord2
11973	UGAGAUUGUGAAUGAUAAAGAUUGG	1	0	transposon	LTR	invader3
11974	AGGGCUGCAAUUCUUUCCCGUGGGAC	1	1	transposon	LINE	Fw2
11975	CCGAAUACGAAGAACAGGCAGAAUCCGA	1	0	transposon	LTR	gypsy12
11976	GCACUCAUAACGAGGUUGCGCAUU	1	1	transposon	LINE	Doc4
11977	UUACCUGAACGCCUUAGCUGCUGGUA	1	0	coding gene		CG10286-RA|CG10286
11978	CUGUUGGAUGAUCCGCCCGACUGGAGG	1	0	coding gene		CG30412-RA|CG30412
11979	AGCCUAGAUGAGUUGCCACAGUUU	1	no match			
11980	GCGCGUCGGGGGGGGGCCGGGGC	1	no match			
11981	UUCGAGAACAAACGCUACUGUUCGGCCG	1	0	transposon	LINE	G2
11982	UGUCCCUUAAAUCAGCACUCGUAAAG	1	0	no annotation		
11983	AUACCCAGGAGUACUAUCGGGCAGA	1	no match			
11984	ACGAUGACGAUCCCACGGCUGGAGC	1	0	transposon	LTR	ninja-Dsim-like
11985	GCCGAACUUGAGACCCCAGCCGCUCC	1	0	no annotation		
11986	AACUUAAAGGAAUUGACGGAAGGGCACC	1	0	ncRNA		rRNA
11987	UCAGGCCAAGAAGGAGAUGGAGCCGAA	1	0	coding gene		CG10859-RA|CG10859
11988	AAUAGGCUCUCUCAGUCGCUUCCCGG	1	no match			
11989	AAGAGGGACGGCCGGGAGCAUUCGU	1	no match			
11990	CCGAAGAAAGCGAAGAUGCGGCC	1	0	coding gene		CG9696-RA|dom
11991	UCCUUCAACUGUGAGAGGAUCCUCAA	1	no match			
11992	CAUGAUUACAGAUAACAGGAUAAAGUC	1	0	transposon	LINE	Cr1a
11993	CAAAGGGCCUCGAAGAUUAUUAACGA	1	0	coding gene		CG10326-RA|CG10326
11994	CUAGCUCAGUCGGUAGAGCAUG	1	0	ncRNA		tRNA
11995	AAAAAAACUGAAGAUGAUCCUAGAAGCCGCA	1	no match			
11996	CAACCAGAGAUCUAGAGGGCCCUAUUA	1	no match			
11997	AAACGAGAACUUUGAAGGCCGA	1	no match			
11998	ACGAGAGGAACUACUACGCCGCC	1	0	coding gene		CG5642-RA|CG5642
11999	ACGAGAGAAGACUUUCACCCGAUGAGA	1	1	coding gene		CG12717-RA|CG12717
12000	AAACCUGAGGGGAUCCUCUAA	1	no match			
12001	UGUCCGGGGCUGCACGCGCGCUAC	1	1	ncRNA		rRNA
12002	CCAGACGUAAUACUCCACUGCCCGGACU	1	1	transposon	LTR	HMS-Beagle
12003	GAACUGGCGCUGCGGGAUGAACC	1	1	ncRNA		rRNA
12004	UACUCGAUUGCAUAAACUCACAUU	1	0	transposon	Unknown	RR50960
12005	ACUCACCGAAUCCCGGGGCCGAGGGAGC	1	no match			
12006	GCGGGAUGAACCGAACGCCGGGUUAAGG	1	no match			
12007	CACUGCUGUCACCGCUGCCGCCCCGU	1	no match			
12008	CGCAAACCCAAAUCGACAAUGCACG	1	0	repeat		trf
12009	CCUGGGAAACGGGGCGGGGCUGGAA	1	no match			
12010	ACUGAUACGAUUGCACCACCUUC	1	0	transposon	LINE	Doc3
12011	GAAGAACGAGGUGAGCAGUGGUGAU	1	0	coding gene		CG5434-RA|Srp72
12012	GCGUCGGGCGUGGUUAGUACUUAGAU	1	1	ncRNA		5SrRNA
12013	CAAAAUCUCAGAUAAAAACCCGAACGU	1	0	transposon	LTR	Burdock
12014	UGCAAGAGCAUGGACGAGUUCAACGU	1	0	coding gene		CG5407-RA|Sur-8
12015	AUCUAUGGGUGGUGGUGCAUGGCC	1	0	ncRNA		rRNA
12016	CUCCGGAUGUGCUGACCCCUGUC	1	no match			
12017	ACCUGAUUGAUUGCGUAGACGACGACC	1	0	transposon	LINE	Quasimodo
12018	UUGACGGAAGGGCACCACCAGGAG	1	0	ncRNA		rRNA
12019	CCCCGGAUAGUUUAGUUACGUAG	1	0	ncRNA		rRNA
12020	GGCCGAUGAAACCCUGGUUCGAUUC	1	no match			
12021	ACGGGAAACAUACCAGGUCCGAA	1	no match			
12022	UCGAUAUCAUUCGUACGAGCUUGAAACC	1	1	transposon	LTR	invader6
12023	CCUGCGCCGUAGGCAGGUCC	1	2	coding gene		CG15651-RA|CG15651
12024	AUUCCGGAGAGGGAGCCUGAGAAA	1	0	ncRNA		rRNA
12025	UAGAUCAUUCCUAGGUUACAGGUUCCU	1	0	transposon	LTR	springer
12026	ACCGAACCGAUUAAAAAUUAAAGUGCA	1	0	no annotation		
12027	UUUAUGUUAGUUACUUGUUCCCCGG	1	0	ncRNA		rRNA
12028	AGAGGUUAAGCCCGAUGAACCUGAAU	1	0	ncRNA		rRNA
12029	GCUAGAACGUAGUCUCUCGAGAGCGA	1	no match			
12030	GCCAUCUCAGUUCGGAUUGUACUCU	1	no match			
12031	CACUAUAACAACGGACCAGGGAAC	1	0	transposon	LINE	Doc4
12032	ACAACCAGAGAUCUAGAGGGCCCG	1	1	no annotation		

12033	AUGCGCCGAAGAUAUGGGUUUGCAGU	1	no match			
12034	AAAUCCGGACAAACACACAAGACCCGU	1	1	transposon	LTR	opus
12035	CUCGAAUCAACAAAUUGCGCCGAUAU	1	0	coding gene		CG41545-RA|CG41545
12036	AACUUGAGACCGCCCCUGGACGA	1	0	Su(Ste)		Su(Ste)
12037	CUGGUAAUUGCCGUUGUUACCGUUGU	1	no match			
12038	CCCGAAAGCAGACGAGGUAUGUACC	1	no match			
12039	AAACCGUGAAGAUGAUCCUGAGUCGUC	1	no match			
12040	ACCGCCCUGAACGCGCCCGAUCUCGUC	1	no match			
12041	UAUUCAAUCAAGUGACGCUGAAUCCGA	1	0	transposon	LTR	Stalker4
12042	GCACCCGUAGCUCAGCUGGAUAGAGCA	1	no match			
12043	AAACCGAUAUGGUUGGACCACAAUCUUUU	1	1	ncRNA		7SLRNA-RA
12044	ACUUCAAUACAGCGAGAUGGCCCA	1	2	transposon	LTR	Quasimodo
12045	GCUCUCGAUUGUAUGCCCGAUAUA	1	1	transposon	LINE	R1-element
12046	ACUUCGAACGCACUUGCGGCCCCGGGU	1	no match			
12047	AACUUCGGGAUAAGGGUUGGCUCU	1	1	ncRNA		rRNA
12048	CAAAGACGGAUCACUUGCCGAUUUAA	1	0	transposon	LTR	opus
12049	GAGAUAUUGAGCGACUUUCCCCGUAGU	1	1	transposon	LTR	3S18
12050	AACAAACAGAAUUCCCGAAUGAGAG	1	1	transposon	LINE	Cr1a
12051	AAGAAGAAGAAGGAUCCCUGUGCC	1	1	repeat		trf
12052	GCACGAAAUGAACCCGAAAGCACU	1	0	transposon	LINE	Doc3
12053	CGAGAGGUCCGGGUAAACCGCU	1	0	ncRNA		rRNA
12054	UACGGAGCAUUGGACGAGCAGCGC	1	0	coding gene		CG3443-RB|pcx
12055	GACCAUGAGAUUACACUGGAGGAUG	1	0	transposon	LINE	Cr1a
12056	ACCACCCUGAACGCGCACGAUCUCGUC	1	no match			
12057	CAGUACGUGAAACUGCUUAGAGGUU	1	0	ncRNA		rRNA
12058	CAGCUAGACUAUGUAUCCCCGGG	1	1	transposon	LINE	Doc
12059	UGGUGAACGACGGCGACAAGCC	1	no match			
12060	AACCGAAUAAACGAAAUGGAUGACA	1	1	transposon	LTR	invader4
12061	AAACCAAAGCUUUUGGGCUCCGGGGG	1	0	ncRNA		rRNA
12062	GCAUGGUGAACGAGGCUGAAAUUA	1	no match			
12063	ACGGUAAGACGUCGUGGAGGACCGAAC	1	no match			
12064	GCGAGGAGGAGGACUAGCUGGCCGA	1	0	Su(Ste)		Su(Ste)
12065	AAGCAAAAAUUGAGAUUAAAUGACA	1	0	transposon	LINE	RR49783
12066	CGCUUGGGAACACCGCGUGUUGUUGG	1	0	ncRNA		5SrRNA
12067	UCGCAAAGGUUGACCGAACUUGA	1	1	ncRNA		rRNA
12068	CCCCUGGACGACCCACGAUUUAACAUU	1	no match			
12069	ACCAAACUCCGAAUACCUGCGAG	1	no match			
12070	GCUCCGGGGGAAGUAUGGUUGCA	1	0	ncRNA		rRNA
12071	CUACGUCCGUUGGAUUAUGCCUG	1	0	ncRNA		rRNA
12072	UGACUCUGUGCUGCUUAGGGAAACCAC	1	0	transposon	LTR	DM88
12073	AUCCCCACGGACUUGAUCGCAUCGA	1	no match			
12074	ACGGGAAACUUACCAGGUCCGAAC	1	no match			
12075	AGGCGCGGGUAACCCGUUGAACCCC	1	no match			
12076	CUGUUAAGAAACCACUCGGGCCGCC	1	1	transposon	LINE	Doc4
12077	ACCCUGAACCGCGCCCGAUCUCGU	1	no match			
12078	AAAACUACAAGCUGAGAUAGACCAC	1	0	transposon	LTR	mdg1
12079	AGCUGCCCGUGAGAUAUUGCAACCCAU	1	0	coding gene		CG18662-RA|CG18662
12080	GUGGAGCCGGGCGUGGAAUGCGAGUGCC	1	no match			
12081	GCUAAAUACUCGUGAUCGACCGAUAGU	1	no match			
12082	CCAGAAAGCACCGUUUCCCGUCCGAUCAA	1	no match			
12083	AUACACGUUAUCGAGCAGGAACUA	1	1	transposon	LTR	blood
12084	UAAGGGUCGGGUAGUGAGGGCCUU	1	no match			
12085	UAUGGGACAUGUGCUUAUUAGGCUA	1	no match			
12086	UCGAGAUGGAGAGGCUUCAGAGUGAG	1	0	transposon	LTR	ninja-Dsim-like
12087	ACACCCAAUCCCAAACAUGGACGAA	1	0	transposon	LTR	McClintock
12088	GAAGGAGGAGAACCCGAUGCCAUCAGC	1	0	coding gene		CG18250-RA|Dg
12089	GUGGGAGAUAUUAGACCUCGGUUUGGU	1	0	repeat		LSU-rRNA_Dme
12090	UGCACCGAAAGUAUAUGAGAAGAACA	1	no match			
12091	AAACUUCGGGGAUAAGGAUUGGCUCUG	1	1	ncRNA		rRNA
12092	ACCCUACUGUUGAUGUGUUGUU	1	no match			
12093	CGGUCUAUUCGCCAGGUCCCGACGGAU	1	1	transposon	LINE	Cr1a
12094	GAAGCCGGGAGAUGGAGGUUGC	1	no match			
12095	UUGUAAGGACCUGAGAACUUGUACG	1	0	transposon	LINE	Doc4
12096	AGAAGAAGAGAUGCCCCAGUGAGUUU	1	no match			
12097	UGACCAUUUACAAAAUGUAUUCCGA	1	0	transposon	LINE	Doc
12098	CGCACACACGAGGAGAGCAGAAUGGAGAGCA	1	no match			
12099	GCUUAAAGUAGUUGAUGGUUUAAU	1	0	transposon	LINE	Cr1a
12100	AUAACAAGAUGCAUACGAUGUGGC	1	0	transposon	DNA	1360
12101	UGAUUAGACAUUCCCGUUUAUCGCGAC	1	0	transposon	LTR	RR50207
12102	CGCCGACUUAUUCCGAGAUGCCAAUG	1	no match			
12103	CCGAGAUGGAUAUACCUCUGGUGUACCA	1	no match			
12104	CAGUCCGUGGACGGUGUGAGGCCGGU	1	no match			
12105	AACUGAAUAAACGAAAUGGAUG	1	0	transposon	LTR	invader4
12106	GGCGAUCUCUUCACGGGCGGUGAUCA	1	no match			
12107	AUACUAAGCAGGGACGUUAAUGAA	1	0	coding gene		pseudogene
12108	GUGGCGUCCGGUGAGCUCUCGCUGGCCC	1	no match			
12109	UGCUCUCGAAUCUAUGCCCGAUAUAC	1	1	transposon	LINE	R1-element
12110	ACUGGCACGGACUUGGGGAAUC	1	0	ncRNA		rRNA
12111	AAGCUGAGUGAUCUACCCAUGGCCAG	1	no match			
12112	CAACCAGAGUUCCCAGUGCUUAGU	1	0	transposon	LTR	Dm88
12113	GAGGUUAAGCCCGAUGAACCUGAAU	1	0	ncRNA		rRNA
12114	AACGUGAGCUGGGUUUAGACCGUC	1	0	ncRNA		rRNA
12115	GGACAUGUGGCGUACGGAAGACCCGC	1	no match			

12116	UGCAACAAGAGCGUGUCCCGGGCAGC	1	1	coding gene		CG10229-RA|katanin-60
12117	CAAUAAGUGACCACCAUGCUAAUGUG	1	0	transposon	LTR	17.6
12118	CAAACCGAGAGAAGACUUUCACCCGAU	1	0	coding gene		CG12717-RA|CG12717
12119	GCCGAACUUGAGACCCCAGCCGCUC	1	0	no annotation		
12120	UGCCGGAGCGAGGAGGAGGACUAU	1	1	Su(Ste)		Su(Ste)
12121	CACAUCGGUAUGGCGACUUCUUU	1	0	coding gene		CG5642-RA|CG5642
12122	CAUCAGGCCAGGCCACCGUUCCGAA	1	0	transposon	LTR	opus
12123	ACUGUGAGGGGUCCUCAA	1	2	coding gene		CG8442-RA|Glu-RI
12124	ACCCCUGAACGCGCCCCGAUCUCGU	1	no match			
12125	CACACUCUGAUUCCGAUUCGAGUUCAA	1	2	transposon	DNA	transib2
12126	AUUUCCGAUACGUGUUCAACGCCGA	1	0	transposon	LTR	GATE
12127	AGCCGAAGUCAUUGUGCGCCAGAGUAA	1	0	transposon	Simple_repeat	I
12128	AGAUCCUACAACCGUGUGAGGGGAUCCUCAA	1	no match			
12129	ACUCGCCGAAUCCGGGGCCGUAGGGA	1	no match			
12130	AACAAAGAAGAGCAGAGCACCAUGAU	1	0	coding gene		CG5252-RA|Ranbp9
12131	UACCGGAGCACCAUCGACCGAUCCGGAU	1	no match			
12132	CUGGCGAAUAAAAGUAAACGAGCA	1	0	transposon	Low_complexity	F-element
12133	ACGCAUGUGUCCUAAGGCGAGCUC	1	no match			
12134	ACCCAUGAGCGAGUCAUUCCGACA	1	0	transposon	LTR	blood
12135	ACGCUUCAUCCAAGAGAACUGUGCCCG	1	2	transposon	LINE	R1-element
12136	CUCCCCGCCCCACCCCGCGCGCGCCGC	1	no match			
12137	AUGGGACUCAAAGAAACAAUCCAGUGGU	1	no match			
12138	GCACCGUUUCCCGUCCGAUCAA	1	no match			
12139	GCUUGCUGGGCGGAUUAGUGGCGAA	1	no match			
12140	GGUGCCCAAAUUAACAACUCAUGCAG	1	0	ncRNA		rRNA
12141	AGCGAUGAUACCGAAGAACGGGAAAU	1	0	transposon	LTR	Dm88
12142	CAACGUUUAAAGAUGGAUGAACACG	1	0	no annotation		
12143	UCAACUGUGGGGAUCCUCAA	1	no match			
12144	UCACCGGAUGAAUAACGAGAACCU	1	0	transposon	LTR	gypsy
12145	GUGCCUAUACAUUGCUAUGUUGACGCG	1	0	transposon	LTR	DM88
12146	AAUACUGUCCCUUAAAUCAGCACUCGU	1	0	no annotation		
12147	AAUGGUUGCAACGGGUAACGGGGAA	1	1	ncRNA		rRNA
12148	CAGACAUCUACUUCCCCCGCUCUUA	1	0	transposon	LTR	invader1
12149	GGAAACCCUGAUGGAAGACCGAAACA	1	0	ncRNA		rRNA
12150	AUUCGUAGACGACCUGGGCGGGUUU	1	no match			
12151	CGGGAAACUUACCAGUCCGAA	1	no match			
12152	AUACCACCCUGAACGCGCCCGAUCU	1	no match			
12153	GAACUUGAGAUCGGCCCUGGGACGACC	1	no match			
12154	CCGAAGAUGACCAAGUUGGCUACCGAA	1	0	transposon	LTR	gypsy6
12155	CCCGAAAAAAUCCCCGGCUGCGAC	1	1	transposon	LINE	F
12156	CAUACCUUGAUUGGCAGUCCCGAUGA	1	0	transposon	LINE	Rt1b
12157	CAUCGAUCUGAACAAUGACACUCC	1	no match			
12158	ACGAACAAUAAAACAUACGCCCGGU	1	0	transposon	LTR	GATE
12159	AAAGAGACAAUUGACGGUACCGCAU	1	1	transposon	LTR	Dm88
12160	GCGAGGAGGACGGACUAGCUGGCCGA	1	no match			
12161	AUGACCGGGCGAGGAGGACGGACUAGCU	1	no match			
12162	ACAACCGAGGAGGACCUUGAUCUGCACG	1	0	coding gene		CG30417-RA|CG30417
12163	AAUCAUGCCGACGGGCGCUGACCCCC	1	no match			
12164	AUUCCUGGAGCAGCGAUAUAGGACU	1	0	transposon	LTR	diver2
12165	CACGAAACUUGAACGGGAUUAAGA	1	1	no annotation		
12166	CAGAUCAUUUCUGGGGCUAGCAAG	1	0	transposon	LTR	gypsy4
12167	UGAGCAACGGUUAGACCCUGAGAUAA	1	0	transposon	LTR	micropia
12168	CUGGCAACAUGUCGAGCUCCCAGAA	1	no match			
12169	GACUACCAGGAGACUUUUGAGUUCA	1	0	coding gene		CG7176-RA|Idh
12170	CAUGCAAGGGAACGUGAGCUGGGUUU	1	0	ncRNA		rRNA
12171	ACAAACAGAAUUCCCGAAUGAGAGUA	1	0	transposon	LINE	Cr1a
12172	AUGCUAGAACGCGGUGAUUUCUUUGCU	1	no match			
12173	CGAAGAUGGUGAACUAUGCCUGGGC	1	no match			
12174	UCCGCGAUAGCGCAGGGUCGUCGCAC	1	0	transposon	LINE	R1-element
12175	CAAUCGAGUGUACAAUAUUUAGGAUU	1	0	transposon	DNA	H
12176	ACGGGCGAAGUGCCGGAUGUGAUGAUU	1	0	coding gene		CG4625-RA|Dhap-at
12177	CCGGGGGGCUGGCGGUGCGGCGAC	1	no match			
12178	UCCGCAGUGCGUUGAUAUGGCCAG	1	0	Su(Ste)		Su(Ste)
12179	UGAGACCACAAACGGAUUGCGGACC	1	1	no annotation		
12180	UGCUCUCGAAUGUAGUGCCCCGAUAUAC	1	no match			
12181	UAGACAGAAAAUAAGGGGGACACGC	1	no match			
12182	GCGAAAGCAGGACAUAGUGAUCCGGUG	1	no match			
12183	ACCCUUGAUAAACUGUUGACUUUCGG	1	1	transposon	LINE	Rt1b
12184	UCGAAAUCCAAUUCCUCCAGUUCU	1	0	transposon	LTR	Quasimodo
12185	AACCCCUGAACGUGCCCCGAUCUCGUC	1	no match			
12186	UAUGACCAGCUUGGACCGAACACU	1	0	Su(Ste)		Su(Ste)
12187	ACGGCCAUACCACCCUGAACGCGCCCGAU	1	no match			
12188	CGAGAUAGUACGAAUCGGGUACGAA	1	0	transposon	LINE	I
12189	CAGCUAGACUAUGUAUCCCCCGGGUAA	1	no match			
12190	GCCUAAGCGAUGUGUGGUGCAAGUCA	1	0	Su(Ste)		Su(Ste)
12191	UACGGCCCGGACUGCUCCACGCCCAAAA	1	0	transposon	LTR	ninja-Dsim-like
12192	GUCCCAAUUCAAGAAGUCGGUGUACC	1	1	transposon	LINE	R1-element
12193	GAUCAGGGGAAAGCCGAGAAACGAA	1	0	transposon	LINE	Cr1a
12194	CCGAAUUAAUCGCGAGGCUGCA	1	0	transposon	LTR	mdg3
12195	GCACACAGAAGGAGAGCAGAUGAGAGCA	1	no match			
12196	UACACAAGAUGACGACAUUAAUGCA	1	0	transposon	LINE	G2
12197	GCAUUCAAUACAGCGAGAUGGCCUA	1	0	transposon	LINE	Quasimodo
12198	ACGUUACUGACAGAAUAAGCACCGG	1	no match			

12199	CACCGGGCGGACUUUGCUGCGGACCCUG	1	1	coding gene		CG17935-RA|Mst84Dd
12200	ACAACAACAACACCACCAGCACCCGC	1	0	transposon	LTR	roo
12201	AGAGGGACGGCCGGGGCAUUCG	1	no match			
12202	AGAGCAACUACGAUGGAAGCAAACCG	1	0	transposon	LTR	rooA
12203	AGACAACACAUAUAAAACCCGCCA	1	0	transposon	LINE	G
12204	AUGUCUAUGAGAUGACCGUAAUGAGU	1	1	transposon	LTR	ninja-Dsim-like
12205	GUAAAGGUGGAAAUGAAGAAAGUACA	1	no match			
12206	AAGACAAUGAGAGGUACUUUGACCAA	1	0	transposon	DNA	1360
12207	AGAGAGAACUCGGGUGAAGGAACUAGGC	1	no match			
12208	CAACAAGACCCUUGGAAAUCUAUCGA	1	0	coding gene		CG14938-RA|crol
12209	CAACGUAGGGGGUGGUGCAGUA	1	no match			
12210	UGCUCGCCGAAUCCCGGGGCCGAGGG	1	no match			
12211	CCAUGGUUGCAACGGGUAACGGGGA	1	0	ncRNA		rRNA
12212	AGCGGCGAGGCGGUCGAGGUACUGGU	1	0	coding gene		CG10365-RA|CG10365
12213	UUCAACCAGAUGGGCUUGGAGUAUUU	1	0	repeat		trf
12214	CAGAUCAACAAAGUACGAUGGCGA	1	1	no annotation		
12215	ACUGACUGUAAUUCUUUAAAGGCC	1	0	transposon	LTR	micropia
12216	CAUCUUCUUAACGCAGCUGCCGAACU	1	0	Su(Ste)		Su(Ste)
12217	AACACGUGCGCGAGUCGGGGGC	1	no match			
12218	AUCCUCAACUGUGAGGGAUCCUCUAA	1	no match			
12219	GCAAACCCAAAAAGACAAUACACGACA	1	0	repeat		trf
12220	AUGUUUCAUCGUUAGACGGCUCGGGCG	1	1	AT-chX		AT-chX-B
12221	AAACGGUGUUAUUGUGGGCCGGUAC	1	no match			
12222	AUAAACGUUAACCCGCUACUCGCCGC	1	0	transposon	LTR	opus
12223	AGGCGCGGGUAGCCCGUUGAACCCC	1	no match			
12224	CGCACCGCCAUCUCGGACGCAUCC	1	1	coding gene		CG7235-RA|Hsp60C
12225	UCUGGUUGAACGUGUCCUGAAAGU	1	0	Su(Ste)		Su(Ste)
12226	GCCUGAAGGAAUGAGGGUUUUCCCCG	1	0	transposon	LINE	R1-element
12227	UAUAUUAAAGUUGUUGCGGUUAAAACG	1	0	ncRNA		rRNA
12228	UGCUCUCGAACGUAUGCCCGAUAUA	1	1	transposon	LINE	R1-element
12229	CCCUGUGGACCCUGUGGACCUUGUUGCG	1	0	coding gene		CG17945-RA|Mst84Dc
12230	CGCAGAGUACGGGGCCUACUGAA	1	1	transposon	LINE	baggins
12231	AUUCCGAUAACGAACGAGACUCUGGC	1	no match			
12232	ACCAGAUGGGCUUGGAGUAUUUCAG	1	0	repeat		trf
12233	UGGAAUCCGAGGUGCCGUAGAU	1	no match			
12234	GCGAACGUUGCGAUACCAGUCCUUU	1	1	transposon	LINE	INE-1
12235	AAGGACAUUUUGGAAGAGUUCCAG	1	0	transposon	Simple_repeat	I
12236	GCAUUCAAUACAGCGAGAUGGCCCUAA	1	2	transposon	LTR	Quasimodo
12237	GUGUAACAACUCACCUGCCGAAGCUA	1	1	ncRNA		rRNA
12238	UGUGAGAGGCCCCGGGUUCGAUCCCC	1	0	ncRNA		tRNA
12239	ACAUUCAUAACCGAGCGGCUGUUCAA	1	1	transposon	LTR	GATE
12240	AGAGCGUCCGACGAAUACCCUGGG	1	no match			
12241	CCAAUUACAUUAGACGAGACGGCCC	1	0	transposon	LTR	opus
12242	UCUCUUUAGUAAUAACUCGAUAUUU	1	no match			
12243	ACGAGCGCACGGUGUUAGGACCCGAAA	1	no match			
12244	ACACAGAGGAGAGCAGAUGAAGCA	1	no match			
12245	CCGUUGGAUUAUGCCUGAACGCC	1	0	ncRNA		rRNA
12246	UACGCAGACAGAGCAAAACCCAUGA	1	0	transposon	LTR	297
12247	GACGCGAUGUGAUUUCUGCCCAG	1	2	ncRNA		rRNA
12248	GAUUCCAAGAAGAUUAACAGUGACA	1	0	no annotation		
12249	UGUUCCCCGGAUAGUUUAGUUACGU	1	0	ncRNA		rRNA
12250	UGCAACUCGAGCGACGCCGUCUCU	1	0	transposon	LTR	Quasimodo
12251	CCUAUUGGAUAGUCCCGAAGAAACC	1	no match			
12252	UGGCUCCGGGGGAAGUAUGGUUGC	1	1	ncRNA		rRNA
12253	CAAACCGAUAUGCUAUCUUGG	1	0	transposon	LINE	Doc
12254	CCCGUAAACCGCCAUCAGCAACACACC	1	0	transposon	Low_complexity	F-element
12255	UGAGUAGAGCUCCGAGACCCCAG	1	0	repeat		trf
12256	GUAUACCUAUACAUUACCGCAAA	1	1	ncRNA		rRNA
12257	UGCGGUGGUGCAGUCCGACAUGAAGC	1	no match			
12258	ACUUGAACGAUCGGUUACUCGUCAAC	1	0	transposon	LTR	micropia
12259	UCAGAUAAAAACGUGUUAGUAAAUAUA	1	0	no annotation		
12260	UUUGUCAAUAAACAAGAUUCCCGAAA	1	0	transposon	LTR	micropia
12261	UCAACGUGAGGGGAUCCUCAA	1	no match			
12262	CAACUCGUUACAAUCGUGUCUACGAC	1	0	transposon	LTR	opus
12263	GACACUAGACAAUAACGACCGGAUA	1	0	transposon	LTR	RR50207
12264	UCAAAUAUUAUUUAAUUUAAUAUUCGC	1	0	repeat		AT_rich
12265	GAACUUGAGACCCCAGCCGCUCCUC	1	0	no annotation		
12266	CUGAAACAAGAAGAGGACGACGAUC	1	0	transposon	Low_complexity	F-element
12267	UACGACGAGAACGAGAUCAGCGAGU	1	2	transposon	LTR	frogger
12268	CCCCCUGCGUAUCGAUAUAGCCGA	1	0	coding gene		CG1340-RA|CG1340
12269	ACCGACGAGGAUCAGUUGAAAGUCGAA	1	0	no annotation		
12270	AUUGAGAGGCGCCGUAGUUACCACCC	1	1	no annotation		
12271	UCUAGAGACAUAGACAGCGCAAGU	1	0	transposon	LINE	INE-1
12272	AAGAACAAAAAGAGGACUUCGAGUAG	1	0	no annotation		
12273	AGAACGCUGAACGAACCCGAAAUCAAC	1	0	transposon	LTR	Idefix
12274	GAUUCCGUGGGUGGUGGUGCAUGGCCG	1	no match			
12275	UGUGGUCGACUCGGAUGCAGUGCCGCGC	1	1	coding gene		CG4799-RA|Pen
12276	CAAUCCAGAAUGUUCUCCUUGAGU	1	1	transposon	LINE	G2
12277	ACAACCGGACGGGCCUAGAGUUAUU	1	1	coding gene		CG40635-RA|CG40635
12278	UAACAAGAUGCAUACGAUGUGG	1	0	transposon	DNA	1360
12279	UGUAUCUAGGACUACCAUGGUUGCAA	1	1	ncRNA		rRNA
12280	AGAUCAAGAAUCCCGUGCCAAGACUGUC	1	1	transposon	LINE	Circe
12281	UAAUCGAGCGCCUCACCUAUCACAU	1	0	coding gene		CG7793-RA|Sos

12282	AGCGACAUGCUCGAGAUGACAACC	1	0	transposon	LTR	micropia
12283	GCCAAUUACAUUAGACGAGAGGCCC	1	2	transposon	LTR	opus
12284	CAAAGCGUUUUGGGCUCCGGGGGA	1	no match			
12285	GCGGGAGUAGCUCAGUUGGUAGAGCGC	1	no match			
12286	AACUCAACUGUGAGGGGAUCCUCAA	1	no match			
12287	GCCAAUGAAAAACUACACCCAGACGGA	1	no match			
12288	AACCCUGAACGCGCCCGAGUCUCGUC	1	no match			
12289	GCGGAAAGAGAGCAAUUUUGGCC	1	0	transposon	LINE	Cr1a
12290	AACAUCGAGAAUACUGCACAGCCAU	1	0	transposon	LINE	Doc
12291	ACAAACCGACCCCGAGCUGUUUGCCGCC	1	no match			
12292	AACGGAUGACAAUGUGGCACACGC	1	no match			
12293	ACUCGUCGAAUCCCGGGGCCGAGGGAGC	1	no match			
12294	UUAGGACCAAAGAUGGUGAACU	1	no match			
12295	CCGAAGAACAAUUUUACCUACAAGACCG	1	0	transposon	LINE	G6
12296	CAGACCCAUCAACGUCAACCAGACGG	1	0	transposon	LINE	INE-1
12297	GAUGUUGACCAGUGGAUAAAUGA	1	0	coding gene		CG40084-RA|CG40084
12298	CGGUCUGAUUGCCUCAGGGCUAGGA	1	0	transposon	LINE	F
12299	CAAGCUAGACUAUGUGUCCCCGGGUAA	1	1	transposon	LINE	Doc
12300	AGCGAGAUAAAGCGAAAACCGUCUAAA	1	no match			
12301	GAGGGCAACCUACGGGGCAUUAUAC	1	1	transposon	LINE	R1-element
12302	GCCCGGAUAGCUCAGUCGGUAGAG	1	0	ncRNA		tRNA
12303	CACCAAGAUUAUCAACCCGUGUUUU	1	0	transposon	LINE	Cr1a
12304	CUCCGGAGUGAGGCUUGGCCAUUGC	1	0	ncRNA		snRNA
12305	UUCCUGGGAUCUGCCAUAUGAUAGGCG	1	0	transposon	LINE	Rt1b
12306	AAAACCCGAGAUGAUCCUGAGUCGC	1	no match			
12307	UGUUUCAUCGUUAGGCGGCUCGG	1	1	AT-chX		AT-chX-B
12308	UGCUCCGAAAACAAGGCCCCGUCGCUAA	1	no match			
12309	UCAGCCCGAACCCUUCUGGUGUGU	1	1	transposon	LINE	baggins
12310	UCCGGUCCCGGGACAUGGCGUGGUGGGU	1	no match			
12311	UUAGAUGGGGGACCGCUUGGGAACAA	1	1	ncRNA		5SrRNA
12312	CAUCGCUGAGGCCCACUGGCAGGGUG	1	0	Su(Ste)		Su(Ste)
12313	AAAAACCAAAAUAUCCUGAUUCAAC	1	0	transposon	LINE	Doc
12314	GGGGGGGCGGCGGGACCGGGGG	1	no match			
12315	GAGAUUCAAACAACGAACGCCGCGGGUG	1	0	coding gene		CG33935-RB|mei-217
12316	ACCCGAAAGCAGACGAGGUAACGUACC	1	1	transposon	Low_complexity	F-element
12317	GCCACACGCAUCCUAGAAUCGACCGA	1	0	transposon	LTR	flea
12318	AACCAAGCUUUUGGGCUCCGGGGG	1	2	ncRNA		rRNA
12319	AAUCGCUCGAUGUCGUAAGGUGACCG	1	0	coding gene		CG40609-RA|CG40609
12320	CGCUUUGCGUGUGAGAGGCCCCGGGU	1	1	ncRNA		tRNA
12321	CAAGCUAGACUACGUAUCCCCGGGUAA	1	1	transposon	LINE	Doc
12322	UCGAGAUUUGUGUUAGCCACUGAGA	1	1	transposon	LTR	G4
12323	GGGAAAGGCGAAAAGAACCCCGGAG	1	no match			
12324	AAGAAGGCCAAGACGACAACCGG	1	0	coding gene		CG7590-RA|scyl
12325	GAAUGCCGCAGCGAGGAGGAGGACUA	1	1	Su(Ste)		Su(Ste)
12326	GAGGAAAUCAUUGAAGACCUUCGCAGA	1	0	transposon	LINE	G6
12327	CGACGCGAGAGGUGAAAUUCUUGGACC	1	0	ncRNA		rRNA
12328	AGUAGCCUCGAGGAAAUCCAGACGA	1	no match			
12329	CUAGUUAUAACGCCAAGUCUGGACCGA	1	1	transposon	LTR	G4
12330	CCCGAAAAAUCCCCGGCUACGACCU	1	2	transposon	LINE	F
12331	CGGGAUAAGGAUUGGCUCUGA	1	0	ncRNA		rRNA
12332	ACCAAAGUCUUUGGGUUCCGGGGGGAG	1	no match			
12333	AAAACCUGAAGAUGAUCCUCAGUCGC	1	1	transposon	LINE	Helena
12334	CCGGCUAGCUCAGUCGGUAGAGCAUG	1	0	ncRNA		tRNA
12335	GUACUUCGGGAUAAGGAUUGGCU	1	2	ncRNA		rRNA
12336	ACGGUGAAUACGUUCCCGGGCCUUGUAC	1	no match			
12337	AAAAACCUGAAGAUGAUCCUGAGUCUC	1	2	transposon	LINE	Helena
12338	UUCAAGGUUACCCCCAGGAUGCUGUGCC	1	no match			
12339	UGCAUCCCAAUUUAGGCGCCGCAA	1	0	transposon	LINE	Cr1a
12340	AACUCGCCGAAUCCCGGGGCCGAGGG	1	no match			
12341	GGGAUCAACUGGGAGGGGAUCCUCAA	1	no match			
12342	CGUACAACAUAUUUCGACGAACGGUGAC	1	no match			
12343	UCGAUGGCGACUACCUCCGAUGGC	1	0	transposon	LTR	Max
12344	GGGCCACCCCUUCCACGGCGCGACCGC	1	no match			
12345	ACGUAAACAACCCCAGGCUGACUCCG	1	0	transposon	LTR	Max
12346	UGGAACACAUAGAGGCUUCGAGGA	1	0	transposon	LTR	HMS-Beagle
12347	UGCUCUCGAAUGUAUGCCCGUAUAUAC	1	no match			
12348	UUCCCUAUGAUGAUGUGUUGUU	1	no match			
12349	GCACAGAGGAGAGCAGAUGAGAGCA	1	no match			
12350	UCCAAGCGAAUCGAGGAUGCAAAU	1	0	transposon	DNA	transib2
12351	UAUUUCAUCGUUAGACGGCUCGGG	1	1	AT-chX		AT-chX-B
12352	CAACUAAGGUGGAAAUGAAGAAAGA	1	no match			
12353	GCCCCGAAAACACGAGAGCCGAGCGCUGGU	1	1	transposon	LTR	opus
12354	AUACGAAACCAAACCGUAAAGA	1	1	no annotation		
12355	CCGAACUUGAGACCGCCCCUGGACAA	1	0	Su(Ste)		Su(Ste)
12356	AAGGAGUGUGUAACAACUCACCUGCC	1	0	ncRNA		rRNA
12357	ACACAUGCAAGUCGAGCGGGCCCU	1	no match			
12358	AUUGUUCCAAACACUUACUUGUCCGC	1	0	transposon	LTR	opus
12359	CGGAGAAGCCGGCGGGAGCCCCGGGGAGA	1	no match			
12360	CGUACUAAAACAAAAACCUCCGCGUGC	1	0	transposon	LINE	Rt1b
12361	CGGCACGAGACCGAUAGUCAACAAGUACC	1	no match			
12362	UACAUCGAAAAGCGAGUCGACGUA	1	1	transposon	LTR	invader4
12363	CAAGCUAGACUAUGUAUCCCCGGGUC	1	1	transposon	LINE	Doc
12364	UAGUAACAUAUCUCGGCGUUCACCUA	1	0	transposon	LINE	Doc2

12365	CAGCAGCCUGUAAUUCCAGCUCC	1	no match			
12366	CCGCAGGUACGGACCAAUGGCACAA	1	0	repeat		LSU-rRNA_Dme
12367	AACUGACGGCGCCACUGUAAAUGUGUA	1	2	transposon	LINE	R1-element
12368	CACCAUAGGAGGACCGUCGUAUACA	1	0	transposon	LINE	baggins
12369	GCCCAAGCACGAGUCCCACUACGCCUC	1	1	coding gene		CG6298-RA|Jon74E
12370	CAGACAACUAGACCGAUUUAUUCCA	1	0	transposon	DNA	transib2
12371	ACGAGCGCACGUGUUAGGACCCCGAAA	1	no match			
12372	ACAUAACGGAACCGCUAGCAGAUCGCGA	1	0	transposon	LTR	opus
12373	AAUGCUGGACGAGGGUAUGGAGUU	1	0	coding gene		CG13340-RA|CG13340
12374	AAAGCACCGUUUCCCGUCCGAUCAACUG	1	no match			
12375	UGACUUGGGUAUAGGGGCGAA	1	no match			
12376	ACAUCAUAACCGAGCGACUGUUCAA	1	no match			
12377	GUUCGAUUCCGGAGAGGGAGCCUGAGAA	1	0	ncRNA		rRNA
12378	AACGAUGGCGGCGCUGUCUCACCCG	1	no match			
12379	GCAUUCGAGACCCAACGACACACAU	1	0	transposon	LINE	Rt1b
12380	AGGAAUUGACGGAAGGGCACCACCAGG	1	0	ncRNA		rRNA
12381	AACUGUGAGGGGAGUCCUCAA	1	no match			
12382	AAGAUAAUAUUCCGAUACAUUUUCC	1	0	transposon	LINE	Circe
12383	UCUCCACGAACGCACAGCUGAAGA	1	0	transposon	LTR	ninja-Dsim-like
12384	GCGGCGAGGACUUCGAUAACCGGCUGGU	1	0	coding gene		CG8937-RA|Hsc70-1
12385	ACUCGAAUGAGUGCUUGAUAUGACG	1	1	Su(Ste)		Su(Ste)
12386	UGCCCCAAGAAGCGCGCCGUGAUCAAU	1	0	coding gene		CG10045-RA|GstD1
12387	UUCAGAAUACCCGGGCUUGAACUUCGG	1	0	repeat		trf
12388	UGGGGAUAACAUGCUGAUACCGCCCAA	1	no match			
12389	UAGGAUGUUGGUGAAAUGGCUUUAA	1	no match			
12390	UGCUAGAAGACGAAUUUACACGAGGAAU	1	1	transposon	DNA	1360
12391	GCUAGAGAGUUGCCGGAGCUGGAGU	1	0	transposon	LTR	ninja-Dsim-like
12392	UAAACUCACAAUUAGGUGGGUCCCGGGA	1	0	transposon	LINE	baggins
12393	UUACAGACUUAGCCCCCGAAGAGA	1	no match			
12394	UGAUUAAAUGGAACGUGUAUCAUUUCC	1	0	repeat		LSU-rRNA_Dme
12395	UACGAACAGAAUACGAGUUGAUGGA	1	0	no annotation		
12396	CUCCCCACGGAACACCAGAGCAAUCCCACC	1	no match			
12397	UUCACCAAGCGUUGGAUUGUUC	1	0	ncRNA		rRNA
12398	CAGCACUACCACAGCUGUAAAGGCCA	1	0	transposon	LTR	Stalker2
12399	UGGACCGCGAGGAUGCCCAGAACCAGG	1	0	coding gene		CG5020-RA|CLIP-190
12400	AGGAACAAGCCGUUACCACCCCGUAG	1	0	transposon	LINE	INE-1
12401	CAAACUGUAAUUCAAGAGGACACAAA	1	0	transposon	Unknown	RR50960
12402	UCCGGGAACAUGCCACCUCCGUUGAU	1	1	coding gene		CG1763-RA|nod
12403	UGUUUCAUCGUUAGACGGCUCGGGCC	1	1	AT-chX		AT-chX-B
12404	CCUGCAUGAAGAUGAUUCCGAAGAU	1	0	transposon	LINE	Circe
12405	UAUUACAAUAAGAAAUAGCACGCGA	1	0	transposon	LINE	TART-A
12406	UGGUCACCAGACGUUGGGGUUGAUGCA	1	0	no annotation		
12407	ACCGAUGGCGAAGGCAGCUCCCUGGGCC	1	no match			
12408	ACGUAGGUCCCGAGCGUUGUCCGGAUU	1	no match			
12409	UGACACAAUGUGAUUUCUGCCCAGUGC	1	0	ncRNA		rRNA
12410	AAGAGCAUAAAGACGACGAAAGUGACAC	1	2	transposon	LINE	Doc3
12411	ACACACCCAUUGCAGACAAAGGAU	1	0	coding gene		CG12051-RA|Act42A
12412	CUGAAGUGGAGAAGGGUUUCGU	1	0	ncRNA		rRNA
12413	UUGAGGCACCAGUCUCCUCGGAG	1	1	ncRNA		tRNA
12414	GUUGCUUUUUGAUCCUUCGAUGUCGG	1	1	ncRNA		rRNA
12415	AACUUCGGGAUAAGGAUUGGCUCUGAAG	1	0	ncRNA		rRNA
12416	AUGGCUCUCUCAGUCGUUUCUCGCGAC	1	no match			
12417	CUGACAUUACAUCAAAAUUUUUUGA	1	no match			
12418	GCGCAUCUUGGGAUAGGCCUUAAUCUG	1	2	no annotation		
12419	UGGAACAUAACGCCAAGCCGUGAC	1	0	transposon	LTR	Quasimodo
12420	GAUCCUCAACUGUGAGGGGAUCCUCUAA	1	no match			
12421	AGCCGUACCUGUUGGUUUGUCCCAUUAU	1	0	ncRNA		rRNA
12422	GCACACAGAGGAGACAGACUCGAGAGCA	1	no match			
12423	AGUACAAGAAGGAUCUGCGCUCCA	1	0	coding gene		CG18743-RA|Hsp70Ab
12424	CGAACUUGAGAUCGCCCCUGGACGACCC	1	0	coding gene		CG42398-RA|SteXh:CG42398
12425	GUGGCAUGUUUCCGGAUUCCCGACA	1	1	ncRNA		alphagamma-element
12426	GGUGACGGCCAAAUUACUCUCUUU	1	1	transposon	DNA	1360
12427	AUACAUCGAAAAGCGAGUUGACGUGU	1	0	transposon	LTR	gypsy
12428	UCUGUAGUUAGCUUGUCCCUCGA	1	2	transposon	LINE	R1
12429	AUCCUUAACAAGAACGACGUUGUCCA	1	1	transposon	LTR	diver2
12430	CGCACUAACACCAUCGCACACCCGA	1	0	transposon	LINE	G6
12431	CCGACAAUUAUAGUUUCCUGUUCGCC	1	2	transposon	LTR	Idefix
12432	GCUGCGUUAAGAAAAUGUCCGGACA	1	0	Su(Ste)		Su(Ste)
12433	ACUCGCCGAAUCCCGGGGCCCAGGGA	1	no match			
12434	GUUCCAGUCCCGUACGUAACUCCA	1	no match			
12435	AUAAUACACAUCGAUCAUUACGGACCA	1	0	transposon	LTR	diver2
12436	CACAGAGUACGGGGCUUACUAAU	1	1	transposon	LINE	baggins
12437	AACUGGCGAUGCGGGAUGAACCGA	1	no match			
12438	AGACUGGCGCUGCGGGAUGAACC	1	no match			
12439	AGCAACAACUCGAUUCAGCUAUGCACC	1	0	coding gene		CG8982-RA|Acp26Aa
12440	CGGUCCCGGGACAUGGCGUGGUGGGUA	1	no match			
12441	AGGGACGGCCGGGGGCAUUCGUAUUGC	1	no match			
12442	ACAUGUCAUCCAGACUCAACACGA	1	0	transposon	LTR	Idefix
12443	GUUCAUGUUAAGGACGAGAAUCUACC	1	0	transposon	LTR	diver2
12444	ACCAGAUCAUCAACCCGUGUUA	1	2	transposon	LINE	Cr1a
12445	GCCGAACUUGACAUCGCCCCUGGACA	1	1	repeat		trf
12446	CAAUUCUGCAACUCGAGCGACGCCGUC	1	0	transposon	LTR	Quasimodo
12447	GGAUCCUCAACUGUGAGGGUAUCCUCAA	1	no match			

12448	UAGGGAGCGAGAGAGCGGAGAGCGC	1	0	transposon	DNA	1360
12449	GCCCUCGUCUCGAUGACACAGCUGGUGG	1	2	coding gene		CG5539-RA|CG5539
12450	UUGCCACACCAAGACAUGUUGUACCC	1	0	transposon	LINE	F
12451	AGAUGGGGGGACCGCUUGGGAACAC	1	no match			
12452	CAAGGAGACCUGAGACUGAGUGUU	1	0	coding gene		CG10637-RA|Nak
12453	GAAACUUAAAGGAAUUGACGGAAGAAA	1	no match			
12454	ACGUAACUAUACACUUCAUAAAGACC	1	0	transposon	LINE	Helena
12455	CUGUAACUCGGGCUGUAACUCGGG	1	no match			
12456	UGGCUCUCUCAGUCGCUUCCCGG	1	0	ncRNA		rRNA
12457	ACUCGCCGAAUCCCGGGGGCCGAGGGGA	1	no match			
12458	GGGCGACUGUGAGGGGAUCCUCAA	1	no match			
12459	AACGACAAAAGUAACGGCGAACAUG	1	0	transposon	DNA	H
12460	UUAAGCGUUCGUUGACGUUGAUGGC	1	0	transposon	LTR	HMS-Beagle
12461	AACCAAAUGAAAGACAUGAUGCGAC	1	0	transposon	LTR	Quasimodo
12462	GAACCGUCAUUCCAGCGCAAGUUGCCCC	1	1	transposon	LTR	ninja-Dsim-like
12463	ACUCGAAUCAACAAAUUGCGCCGG	1	1	coding gene		CG41545-RA|CG41545
12464	CUUACCAGGUCCGAACAUAAGUGUGU	1	0	ncRNA		rRNA
12465	GGAUCCUCAACUGUGAGGCGAUCCUCAA	1	no match			
12466	CAACGAACGGUUAAAGUGCGAUCA	1	0	transposon	LTR	HMS-Beagle
12467	GCCAGUUACAUUAGACGAGACGGCC	1	1	transposon	LTR	opus
12468	UCACCCUGAACGCGCCCGAUCUCGU	1	no match			
12469	UGAGAUAUGCUAGCAGACUUGAGA	1	1	transposon	LINE	G2
12470	GUUCCUCGGGAUCAAGGGCAACGAGUUA	1	1	coding gene		CG40635-RA|CG40635
12471	AGACUUAUAUCCGAGGCCCUGUAAU	1	2	ncRNA		rRNA
12472	UCGCCAGAGGAUGAGCCCGCGUAG	1	no match			
12473	ACGCUGACGAGUGGCUAACGGGU	1	no match			
12474	CUGACAUCUACUCGACUGAACGACA	1	2	transposon	LTR	HMS-Beagle
12475	UGCUGAUGAAGAUCCGUCUUGGCCG	1	0	coding gene		CG3561-RA|KH1
12476	CUGCGACGAGCAAGGUGGCCGCCGGGA	1	0	coding gene		CG8136-RA|CG8136
12477	AACGAAAGGGCACAAGCAACCAGA	1	0	transposon	LTR	3S18
12478	CAUGCAGAUCGUAUGGUCUUGUACCGA	1	0	ncRNA		rRNA
12479	AACGCCAAGCACACCCAUUGGGGAUCU	1	0	transposon	LINE	Doc
12480	CGCAAGCGUCACUUCCAGGCUCCCUCC	1	0	coding gene		CG6846-RA|RpL26
12481	AGACGGCGAUCGAGUGCGGAUAUUACG	1	0	coding gene		CG11262-RA|CG11262
12482	UGUUUCAUCGUUAGACGGCUCGGUC	1	1	AT-chX		AT-chX-B
12483	AAAUGAGAACUUUGAAGACUGAA	1	no match			
12484	UAUGUGCGAGGAGCCCGUGUUCGACU	1	0	coding gene		CG5497-RA|mRpS28
12485	UGUUCAUCGUUAGACGGCUCGGCGU	1	no match			
12486	UAGUAAAUCACACACCUCUAUCGAAGA	1	0	no annotation		
12487	AAGAGAUGGGGAAGCUCCGUUUCAAA	1	no match			
12488	UGCUAUCCGAAUAGACCGCGGCGUAA	1	0	coding gene		CG12861-RA|CG12861
12489	CUUUAAGGGCUGAGUACAAACA	1	0	transposon	DNA	1360
12490	UAAGACGCCUGCAGAGAACGCACCCG	1	0	transposon	LINE	G2
12491	GAUUCCGGAGAGGGAGCCUGAGAGACG	1	1	ncRNA		rRNA
12492	AACUACAACACAUCGAGCGACCGU	1	0	transposon	LINE	Doc3
12493	GUCAGAAAAGUUACCAUAGGGAUA	1	0	ncRNA		rRNA
12494	AGCACUAAUAUAAGCCCUGACGUCCGA	1	1	transposon	LTR	G4
12495	GGUCCUGAGAGAUUGGCGAGCGCCGUUC	1	no match			
12496	AACCUGAAGAUGAUCCUGAGUCGCA	1	0	transposon	LINE	Helena
12497	CAGAGCGUGGCCUGAUUGCUAUGCACCGA	1	0	repeat		trf
12498	AAAUACCAUUGCAAGCUUUACACGC	1	0	transposon	LINE	F
12499	GCUAAUCCCACUGCCAUGCUGCUCUGUC	1	2	coding gene		CG6439-RA|CG6439
12500	AGAUUGAGAGCUCUUUCUUGAUUUU	1	no match			
12501	GAAAUCGGAAAAUCACACCUUUUGACC	1	0	repeat		RSP
12502	AGAAACGUGAACUCGCCCAGUGGUC	1	no match			
12503	UCCGAUAACGAACGAGACUCAAAUAU	1	0	ncRNA		rRNA
12504	AAAUUUAUAUCAGCCCGUAUCCAGCCAU	1	0	coding gene		CG8701-RA|CG8701
12505	CGAGAUGACAACCUUGGACCUGACCC	1	1	transposon	LTR	micropia
12506	GACAAGAUAAACCAGCACUCAAGCA	1	1	transposon	LINE	X
12507	UUCCGGCACAAUGCGCAUGAAUG	1	0	no annotation		
12508	CAGAAGCAUAAAGAGUGCACCUGGC	1	0	transposon	DNA	S
12509	CUGAACGCGCCCGAUCUCGUCUGAUCU	1	no match			
12510	ACGAGCCACGUGUUAGGACCCGAAA	1	no match			
12511	CUACCCGUGAGUGUCCCAAGACUUC	1	0	no annotation		
12512	AAGCUUAUCCUUUGCUUGAUGAUUCG	1	0	ncRNA		rRNA
12513	CAGUCCGAAACACAUCCCCCGAUUAAA	1	0	no annotation		
12514	GUCUCAUCGUCGUAGAACUAGCCCGA	1	1	Su(Ste)		Su(Ste)
12515	AAAGCCUAAAAAUGUUAUUUGGAA	1	no match			
12516	AAAUUCCGUAAACGAAGCCCAAAGUUCG	1	1	transposon	LTR	Rt1c
12517	AUCGAUCCUUUAGUCCCUCGGAAUU	1	no match			
12518	ACGGGAAAGCUUACCAGGUCCGAA	1	1	ncRNA		rRNA
12519	AAACCGAGUGAUUCAGAGUAUGCAUC	1	0	transposon	DNA	invader2
12520	GAGAGAAGACCCUCAGCCUACGAG	1	0	coding gene		CG41521-RA|CG41521
12521	CGCGGAAAGACGGAAAGACCCCAUG	1	no match			
12522	AGUUCCUCGAGAAGCCCGUCCAGAAGCC	1	0	coding gene		CG6066-RA|CG6066
12523	UAGCCGGGGCGAGGACGGAGGACUAGCUG	1	no match			
12524	GCAGUUAUUACUACGAGAAGAUGGC	1	1	transposon	Unknown	RR50960
12525	UGCUCUCGAAUGUGUGCCCCGAUAU	1	no match			
12526	UAUCUCGAAAGACUCGUUCCGGUCGA	1	1	transposon	LTR	G4
12527	UGAUCAAACAAGAAAUGGUACUCGGA	1	0	transposon	LTR	DM88
12528	GUGAUCACUGUGACGCCUUGCGUG	1	0	ncRNA		rRNA
12529	CGCCCCGGGGAGCCUGGCGCGC	1	no match			
12530	GGCGCGUAAAUUACCCACUCCCAGCUCG	1	0	ncRNA		rRNA

12531	AAGGAGUCUAACACGUGCGCGAGUCG	1	no match			
12532	GCCCCCAGUGGCCUAAUGGAUAAGGCAUC	1	2	ncRNA		tRNA
12533	AGCUACGAGAAUAUACCGCAAACC	1	0	transposon	LTR	gypsy12
12534	AAUAGCGACAACCAGAGUUCCCAGU	1	0	transposon	LTR	Dm88
12535	CAUCAUGAUAACGCACCGCCUAGU	1	0	transposon	LTR	opus
12536	UUCUGUGAACGAGUGAACUGGCAA	1	0	Su(Ste)		Su(Ste)
12537	UAAACCGUCAUCCAGCGCAAGUUCC	1	no match			
12538	GGCGCAAUGGCACAGCAUCCUGGGG	1	2	repeat		trf
12539	GGAAACUUUAACUCGAGCAGCGACUAC	1	0	coding gene		CG34407-RC|Not1
12540	AAAAGCCGAGCACUACUAAUUGCCC	1	no match			
12541	GCGAAGGUGGCGGAAUUGGCAUAG	1	no match			
12542	GGUGAAUACGUUCCCGGGCCUUGUACACA	1	no match			
12543	AACGGGGCUCAAACCAGGUACCGAA	1	no match			
12544	ACGAGUAAUAUGAGCGCCCAGAGUGAA	1	1	transposon	LTR	opus
12545	CUGUAACUCGG	1	0	transposon	LTR	gypsy9
12546	ACCAAGCGUUGGAUUGUUCACCCAUGCA	1	0	ncRNA		rRNA
12547	UAUACCAACAACAGAUAAUAUACCGAU	1	0	transposon	LINE	accord2
12548	UAAUAAACUUGACCAUGUUUCCCCCGGGA	1	0	transposon	Low_complexity	F-element
12549	ACGGCCCGAACCCUUCUGGCGUGU	1	2	transposon	LINE	baggins
12550	AGCCCAGAUCUACGCCAUCCGCCAGGCC	1	0	coding gene		CG4046-RA|RpS16
12551	AACUGAUAACGAAAUGGAUGACA	1	no match			
12552	AUGCCGAUGGUGCCGACCCUGGAGCU	1	0	coding gene		CG13340-RA|CG13340
12553	ACGCCCAGAGCAAACACAGUGUGGCG	1	0	coding gene		CG1999-RA|CG1999
12554	CCAGACGUGAUACUCCUCUACCCAGU	1	0	transposon	LTR	HMS-Beagle
12555	AAAACCAUUCGUAGACGACCUGCU	1	no match			
12556	UACCAAACGAAGUCCUACCAUAUGC	1	1	transposon	LTR	flea
12557	CGGUGAGGGGAUCCUCAA	1	2	coding gene		CG4584-RA|dUTPase
12558	CAAACCAAUCCCCAACCAAACCCGUCAAA	1	no match			
12559	UCAGAAAUUAUCCUGUCCCUCAUGA	1	0	transposon	LTR	Stalker2
12560	CAACGACUCUCAGCGGCAGGCCACC	1	2	coding gene		CG31359-RA|Hsp70Bb
12561	AAAACCUGUAGAUGAUCCUGAGU	1	1	transposon	LINE	Helena
12562	GGAGAAGCUGGCGGGAGCCCCGGGGA	1	no match			
12563	ACCCUCAACUGUGAGGGGAUCCUCAA	1	no match			
12564	ACCUGAUUGAUAGAGUAGACACGA	1	no match			
12565	AGAACAACAUCCUGCAACUUGAUCCGC	1	0	coding gene		CG1945-RA|faf
12566	GGCAUUGAGAGGCGCCGUAGUUACCACCC	1	1	no annotation		
12567	GAACUUGAGACCGAACCUGGACGACCC	1	1	no annotation		
12568	GCGGCGUCCGGUGAGCUCUCGCUGGCC	1	no match			
12569	UAACCAAUGAAGUCCUCCUGAGG	1	0	ncRNA		snRNA
12570	ACCAAGAUGAAGGAGACCGCCGAG	1	0	coding gene		CG4264-RA|Hsc70-4
12571	GCAAUCGUUUUAGACGAGUAUUCCGA	1	no match			
12572	UGGAGACGGUAGCAGGCCGAUGG	1	0	transposon	LTR	GATE
12573	UACUGCCGAAGCCUUCCCAGAGAGUU	1	1	coding gene		CG6289-RA|CG6289
12574	CUACAUAUGGUUGAGGGUUGU	1	0	ncRNA		rRNA
12575	CCAGUUACCGAAGCUGCGGAUUUGC	1	no match			
12576	AACUACGAAACCCAAACUCCCGGACA	1	1	no annotation		
12577	AACCAAAGCUUUGGGCUCCGGGG	1	no match			
12578	ACGGGAAAACUCACCAGGUCCGAA	1	1	ncRNA		rRNA
12579	UCGUUAGAUCCAGAAUAUUUAAA	1	no match			
12580	CCCUUCGAGGCGGGCGCGUAUUUCAA	1	0	coding gene		CG31648-RA|CG31648
12581	UCUUGGUGACAUAGUAACUUCUGA	1	0	transposon	LTR	297
12582	GUCGAACACCGGCUUCAGGAUCACGUCC	1	0	Su(Ste)		Su(Ste)
12583	CCACAUCUAAGGAAGGCAGCAGGCG	1	0	ncRNA		rRNA
12584	UCACGAAUAACAGGACCAUCCAAGA	1	1	no annotation		
12585	GCUUUAAUCACGAGCAAUGCGUUGAACG	1	1	no annotation		
12586	GGCUCCGGGGAAGUAUGGUUGCA	1	no match			
12587	CUUAGAUGUCCGGGGCUGCACGCGCGC	1	1	ncRNA		rRNA
12588	GUACGGAAGACCCGCUCCCCGGCGC	1	no match			
12589	UAAUUCGGCCCGAACCCUUCUGGUG	1	0	transposon	LINE	baggins
12590	CCUACAAACAGUACACCUACUAGCCCA	1	0	transposon	LINE	baggins
12591	AGCGAAACGAGCUCGAGGAAUACGG	1	0	transposon	LTR	ninja-Dsim-like
12592	UCCGCCCCCCGGCCCCGCGUCCUCCCU	1	no match			
12593	AAUAAAUAAAAGGUGACCGGCAACGA	1	0	AT-chX		AT-chX-A
12594	AUCGUGUGAACUGCAGGACACAUGAA	1	0	ncRNA		rRNA
12595	GCAGAAUUGAGUCUGUUUUCCCAGGA	1	0	transposon	LTR	GATE
12596	UCCCGUGGUAGGCAGUAGUGGGAU	1	0	ncRNA		7SLRNA-RA
12597	CUCGGCCUAUCGAUCCUUUUGGUUUAA	1	0	ncRNA		rRNA
12598	UACAUACCUGAGAAAUCUUCAAAGU	1	0	transposon	LINE	Cr1a
12599	UCGUAUACUGCUUCAACACGACCCA	1	0	transposon	LTR	Tabor
12600	UACAAUGAUCUGACGGGUUUUUCUG	1	no match			
12601	CGGGGCUAAGCCAGUUACCGAAGCUGCG	1	no match			
12602	ACACAGAGGAGAGCAGAUGAGAGA	1	no match			
12603	GCGGCCCCCUGGACGAAGACUGACGC	1	no match			
12604	CCUGAUGGAAGACCGAAACAGUUCUGA	1	0	ncRNA		rRNA
12605	AUGACACAUAGGGCUUCGAGG	1	no match			
12606	UUCGAGAAGCGAACCCGGCGAACUGAA	1	no match			
12607	AUAAUCAAAACAUCCGAUGACGAA	1	0	transposon	LTR	McClintock
12608	UGAGCAUAAAGACGACGAAAGUGAC	1	1	transposon	LINE	Doc3
12609	GACUGCCCUGGAGCGUUCGACCAACA	1	0	coding gene		CG4264-RA|Hsc70-4
12610	AAAAUAUAGCUCAGUGGUAGAGCAUUC	1	no match			
12611	GACCGAAACAGUUCUGACGUGCAAAUC	1	0	ncRNA		rRNA
12612	ACAGACUCGGAACGCCUCAUUGG	1	0	coding gene		CG31359-RA|Hsp70Bb
12613	CAAGACCCACGGGAUGACUUCUUU	1	0	transposon	LINE	BS

12614	GGCAUUGAGAGGCGCCGUAGUUACC	1	1	no annotation		
12615	GCCCGAUGCCGACGCUCAUCAGUACCCC	1	no match			
12616	AAGUAGUCUAACACGUGCGCGAGU	1	no match			
12617	UUGGAUUGACUCUAGACCGGAAGCUC	1	1	transposon	LINE	G
12618	CUCUGAGAAAGCAGACGCCUGAGU	1	1	transposon	LTR	Max
12619	UUCUUGCGAGGGGAUUCGUAGGUG	1	no match			
12620	ACGAUAAAGAGGCUACCUACUAGAGGA	1	0	no annotation		
12621	UAACAUCAGACUCAAUGCCCUUCAG	1	0	transposon	Low_complexity	F-element
12622	GUGAUGUCAACGUUUGAUCCCCCUGG	1	2	transposon	LTR	ninja-Dsim-like
12623	CCGAAGCCCGGCGAUCUAGGCAUGACC	1	no match			
12624	GGCUUUGAGAUGCGCCGUAGUUA	1	1	no annotation		
12625	UGGGCAAGAGCAAGAAGCAGGUGCGC	1	0	coding gene		CG11601-RA|CG11601
12626	CAGGAUCAAAUAAGUAGACUUCGAGGA	1	0	transposon	LTR	micropia
12627	UAGGAGUUCACCGCCACGACGACC	1	0	transposon	LTR	GATE
12628	CUGACCUGUAUUCGGAUAAUGGAACAA	1	0	transposon	LTR	3S18
12629	CUCGAAUGUAUGCCCGAUAUACGGAGC	1	0	transposon	LINE	R1-element
12630	UCCUAGACCGAGAUGUCCGGGUAAACCG	1	1	ncRNA		rRNA
12631	UGCUCUCGAAUGUAUGCCCGAUAGUGAACC	1	no match			
12632	ACAGCCUGAGAGGUGAUGCGUAGCC	1	no match			
12633	AGUACUCCAGGCCCAUCUGGUUGA	1	0	Su(Ste)		Su(Ste)
12634	CCCCUAUCCAAGCUACAGCACAGUU	1	0	transposon	Low_complexity	F-element
12635	GACAAUGUGAACAAGACGAGCGCCGC	1	1	transposon	LINE	Doc
12636	UGAUACGAGGGAGAAUUGUGAGUCGGC	1	0	transposon	LTR	Rt1c
12637	UAGCCACUGAUAUGCGCCCGUUUU	1	0	no annotation		
12638	CAACGACCGAGAUGCUAGCAGACAUA	1	1	no annotation		

Aub-associated small RNAs

No.	Sequence	Read	Genome mismatch	Annotation (category)	Annotation (detail1)	Annotation (detail2)
1	UCUCAUCGUCGUAGAACAAGCCCGA	43207	0	Su(Ste)		Su(Ste)
2	UGUUUCAUCGUUAGACGGCUCGGGC	13615	0	AT-chX		AT-chX-B
3	UCUCAUCGUCGUAGAACAAGCCCG	6212	0	Su(Ste)		Su(Ste)
4	CAGCACACAGAGGAGAGCAGAUGAGAGCA	3971	no match			
5	UGUUUCAUCGUUAGACGGCUCGGG	3013	0	AT-chX		AT-chX-B
6	UAGAUCUUCACCGUUGACUAGCCCCA	1532	0	Su(Ste)		Su(Ste)
7	UCUCAUCGUCGUAGAAGAAGCCCGA	1183	1	Su(Ste)		Su(Ste)
8	UUCUCAUCGUCGUAGAACAAGCCCGA	1138	0	Su(Ste)		Su(Ste)
9	UCUCAUCAUCGUAGAACAAGCCCGA	1134	0	Su(Ste)		Su(Ste)
10	UCUCAUCGUCGUAGAACUAGCCCGA	1113	0	Su(Ste)		Su(Ste)
11	GCACACAGAGGAGAGCAGAUGAGAGCA	898	no match			
12	AUCUCAUCGUCGUAGAACAAGCCCGA	768	1	Su(Ste)		Su(Ste)
13	UCAACUGUGAGGGGAUCCUCAA	693	no match			
14	UCUCAUCGUCGUAGAACAAGCCCGAG	616	0	Su(Ste)		Su(Ste)
15	CUCAACUGUGAGGGGAUCCUCAA	578	no match			
16	UCUCAUCUUCGUAGAACAAGCCCGA	436	1	Su(Ste)		Su(Ste)
17	AUCCUCAACUGUGAGGGAUCCUCAA	392	no match			
18	UCUCAUCGUCGUAGAACAAGCCCGAA	334	0	Su(Ste)		Su(Ste)
19	UAGAUCUUCACCGUUGACUAGCCCC	328	0	Su(Ste)		Su(Ste)
20	CAACUGUGAGGGGAUCCUCAA	315	no match			
21	UAGAUCUUCACCGUUGACUUGGCCCA	308	0	Su(Ste)		Su(Ste)
22	UCUCAUCGUCGUAGAACAAGCCCGAGG	278	0	Su(Ste)		Su(Ste)
23	UAGAUCUUCACCGUUGACUUGGCCC	276	0	Su(Ste)		Su(Ste)
24	AUCCUCAACUGUGAGGGGAUCCUCAA	266	no match			
25	UCUCAUCGUCGUAGAACAAGCCCAA	263	1	Su(Ste)		Su(Ste)
26	UCUCAUCGUCGUAGAACAAGCCCCGA	242	2	Su(Ste)		Su(Ste)
27	UCCGAACGUACCACGGUGCCCCAGU	237	0	transposon	Low_complexity	F-element
28	UCAUCUUCUUCUCGACGUUGACCA	227	0	transposon	LTR	ninja-Dsim-like
29	UUCUCAUCGUCGUAGAACAAGCCCG	204	0	Su(Ste)		Su(Ste)
30	CCUCAACUGUGAGGGGAUCCUCAA	197	no match			
31	AUGUUUCAUCGUUAGACGGCUCGGGC	190	1	AT-chX		AT-chX-B
32	UAACAAGAUGCAUACGAUGUGGC	186	0	transposon	DNA	1360
33	CGUUGACCAAUUUCCGAACAUACGC	185	0	transposon	LTR	DM88
34	UGUUUCAUCGUUAGACGGCUCGGCC	183	1	AT-chX		AT-chX-B
35	UCCUCAACUGUGAGGGAUCCUCAA	181	no match			
36	UGUUUCAUCGUUAGACGGCUCGGGU	172	1	AT-chX		AT-chX-B
37	CCUCAUCGUCGUAGAACAAGCCCGA	169	1	Su(Ste)		Su(Ste)
38	UAAAUGCCAGAAUUUUAUUGGGGUUC	154	0	transposon	LINE	baggins
39	CAAGAUAUGACCAAAGCAAAGACAC	150	0	transposon	LINE	Cr1a
40	UGUUUCAUCGUUAGACGGCUCGG	134	0	AT-chX		AT-chX-B
41	UAGAUCUUCACCGUUGACUAGCCC	132	0	Su(Ste)		Su(Ste)
42	UUUCUCAUCGUCGUAGAACAAGCCCGA	129	0	Su(Ste)		Su(Ste)
43	UCUCAUCGUCGUAGAACAAGCCCGU	128	1	Su(Ste)		Su(Ste)
44	UCCUCAACUGUGAGGGGAUCCUCAA	121	no match			
45	UAGAUCUUCACCGUUGACUUGCCCCA	121	0	Su(Ste)		Su(Ste)
46	AUCUCAUCGUCGUAGAACAAGCCCCGA	120	no match			
47	AAAACCUGAAGAUGAUCCUGAGUCGC	119	0	transposon	LINE	Helena
48	UGUGACUGAUCUAGAGUACAAUGU	118	0	transposon	LTR	DM88
49	AUCUCAUCGUCGUAGAACAAGCCCG	118	1	Su(Ste)		Su(Ste)
50	UCUCAUCGUCGUAGAAGAAGCCCG	117	1	Su(Ste)		Su(Ste)
51	GCCCCGUACUCUGUGGUGACCAACA	117	0	transposon	LINE	baggins
52	GGAUCCUCAACUGUGAGGGGAUCCUCAA	116	no match			
53	UAGAUCUUCACCGUUGACUUGGCC	109	0	Su(Ste)		Su(Ste)
54	UCUCAUCAUCGUAGAACAAGCCCGAU	107	0	Su(Ste)		Su(Ste)
55	UCGUCAUCGUCGUAGAACAAGCCCGGA	105	no match			
56	UAGAUCUUCACCGUUGACUUGCCCC	102	0	Su(Ste)		Su(Ste)
57	UAAGUAAGCCCGAAGAUUGUCCAAU	102	0	transposon	LTR	opus
58	UGUUUUCAUCGUUAGACGGCUCGGGC	102	2	no annotation		
59	UCUCAUCGUCUUAGAACAAGCCCGA	102	1	Su(Ste)		Su(Ste)
60	UUAUACCCAGGUGAUUACCACUAUU	101	0	Su(Ste)		Su(Ste)
61	CUCUCAUCGUCGUAGAACAACGCCCGA	97	no match			
62	CCUCAACUGUGAGGGAUCCUCAA	97	no match			
63	UGUUUCAUCGUUAGACGGCUCGGGCG	91	0	AT-chX		AT-chX-B
64	UACAAUCUCCGAGCAAGACCCCGA	90	0	transposon	LINE	Circe
65	UGUUUCAUCGUUAGACGACUCGGGC	88	1	AT-chX		AT-chX-B
66	UAGAUCUUCACCGUUGACUUGCCC	83	0	Su(Ste)		Su(Ste)
67	UCUCAUCGUCGUAGAACGAGCCCGA	80	1	Su(Ste)		Su(Ste)
68	UUCAAGGUUACCCCAGGAUGCUGU	80	0	repeat		trf
69	UCUCAUCGUCGUAGAACUAGCCCG	79	0	Su(Ste)		Su(Ste)
70	UGUUUCAUCUUUAGACGGCUCGGGC	79	1	AT-chX		AT-chX-B
71	UUAAUGCCGGAAGCGAUCCCACUGC	77	0	AT-chX		AT-chX-B
72	UCUCAUCGUCGUAGAACAAUCCCGA	77	1	Su(Ste)		Su(Ste)
73	UGAUACCGAGAAUGACAAUGGUUGA	77	0	transposon	LTR	gypsy3
74	GACACAGAGGAGAGCAGAUGAGAGCA	76	no match			
75	CUCAACUGUGAGGGAUCCUCAA	76	no match			
76	UCUCAUCGUCGUAGAACAACGCCCGA	73	no match			
77	CGCACACAGAGGAGAGCAGAUGAGAGCA	72	no match			
78	UCGUCAUCGUCGUAGAACAAGCCCGA	68	2	Su(Ste)		Su(Ste)
79	UCUCAUCGUCGUAGAACAACCCCGA	68	1	Su(Ste)		Su(Ste)
80	CGUUGACCAAUUUCCGAACAUACGCC	68	0	transposon	LTR	DM88
81	UCUCAUCGUCGUAUAACAAGCCCGA	66	1	Su(Ste)		Su(Ste)

82	UCUCAUCGUCGUAGAGCAAGCCCGA	66	1	Su(Ste)		Su(Ste)
83	UACAAUCUCCGAGCAAGACCCCGAUC	66	0	transposon	LINE	Circe
84	UCAUCUUCUUCUCGACGUUGACCAG	65	0	transposon	LTR	ninja-Dsim-like
85	UACUGACCCAGUCGGAUCUAUUGUA	65	0	transposon	LINE	baggins
86	UGGACUUUUUUCCGAUGAGCCAGU	64	0	transposon	LINE	G2
87	UCUCAUCAUCGUAGAACAAGCCCG	64	0	Su(Ste)		Su(Ste)
88	GACUAGCUGGCCGAGGAUCCGAUU	63	0	Su(Ste)		Su(Ste)
89	UCCGAGGAACCAAUUGAUCCAGCUG	62	0	Su(Ste)		Su(Ste)
90	UCUCAUCGUCGUAGAACAGGCCCGA	62	1	Su(Ste)		Su(Ste)
91	UCCGAUCUCCUGAGUCGGACUCCGAG	61	0	transposon	LINE	R1
92	UUAUACCCAGGUGAUUACCACUAU	60	0	Su(Ste)		Su(Ste)
93	CGUUGACCAAUUUCCGAACAUACG	60	0	transposon	LTR	DM88
94	UAAAGAAUCCUGACAAGAUCCCGGG	60	0	transposon	LTR	invader1
95	UCAAGGUAGUAACUAUGACGAUUUGA	60	0	transposon	LTR	flea
96	UCUCAUCGUCGUAGAACAAGCCCUAG	59	1	Su(Ste)		Su(Ste)
97	UCUCAUCGUCGUAGGACAAGCCCGA	58	1	Su(Ste)		Su(Ste)
98	AAUGCCGGAAGCGAUCCCACUGCUU	58	1	coding gene		CG41580-RA|CG41580
99	UCUCAUCGUCGUAGAACAAGCCCGG	58	1	Su(Ste)		Su(Ste)
100	CAAGAUAUGACCAAAGCAAAGACUA	57	1	transposon	DNA	1360
101	UCUCAUCCUCGUAGAACAAGCCCGA	56	1	Su(Ste)		Su(Ste)
102	UGUUUCAUCGUUAGACGGCUCGGC	55	1	AT-chX		AT-chX-B
103	UCUCAUCGUCGUAGAACAAGCCC	55	0	Su(Ste)		Su(Ste)
104	UCUCAUCGUCAUAGAACAAGCCCGA	53	1	Su(Ste)		Su(Ste)
105	UUUCAUCGUCGUAGAACAAGCCCGA	53	0	no annotation		
106	UCUCAUCGUCGUAGAACAAGCCCGACUGUAACUCGGGGUCAAU	53	no match			
107	CACACAGAGGAGAGCAGAUGAGAGCA	52	no match			
108	CACACACAGAGGAGAGCAGAUGAGAGCA	52	no match			
109	CAGCACACAGAGGAGAGCAGAUGAGAGA	52	no match			
110	UCUCAUCGUCGUAGAACAAGCCCA	51	1	Su(Ste)		Su(Ste)
111	GCACACAGAGGAGAGCAGAUGAGAGCAA	51	no match			
112	UCUCAUCGUCGCAGAACAAGCCCGA	50	1	Su(Ste)		Su(Ste)
113	AACUGUGAGGGGAUCCUCAA	50	no match			
114	UUUCUCAUCGUCGUAGAACAAGCCCG	49	0	Su(Ste)		Su(Ste)
115	GGCAACUGUGAGGGGAUCCUCAA	49	no match			
116	AUGUUUUCAUCGUUAGACGGCUCGGGC	49	no match			
117	UCUCAUCGUCGUAGAACAAGUCCGA	48	1	Su(Ste)		Su(Ste)
118	GAUCCAACUGUGAGGGGAUCCUCAA	48	no match			
119	GCUGCCGAGAUAUGCUGCGAGUUGCCAA	48	0	transposon	Simple_repeat	I
120	UCUCAUCGUCGUAGAACAAGCCCUA	47	1	Su(Ste)		Su(Ste)
121	GAUCCUCAACUGUGAGGGAUCCUCAA	47	no match			
122	GACUAGCUGGCCGAGUAUUUGAUAU	47	0	Su(Ste)		Su(Ste)
123	UCUCAUCGUCGUAGAACAAGCCCGUA	46	1	Su(Ste)		Su(Ste)
124	UCUCGUCGUCGUAGAACAAGCCCGA	46	1	Su(Ste)		Su(Ste)
125	CAGCACACAGAGGAGAGCAGAUGAGAGCCA	46	no match			
126	UCGCAUCGUCGUAGAACAAGCCCGA	46	0	Su(Ste)		Su(Ste)
127	UGUUUCAUCGUUAGACGGCUCGGGA	45	1	AT-chX		AT-chX-B
128	UUGGACCGAACACUUUGAACCAAGUA	45	0	Su(Ste)		Su(Ste)
129	UAGAUCUUCACCGUUGACUAGCCCCAC	45	0	Su(Ste)		Su(Ste)
130	UGCCGAGGAGGACCCAGCAGCCAU	44	0	transposon	LINE	Doc4
131	UCUCAUCGUCGUAGAACAAGCCGA	44	2	Su(Ste)		Su(Ste)
132	CUGUGAGGGGAUCCUCAA	44	2	coding gene		CG13800-RA|CG13800
133	UCUCAUCUUCGUAGAACAAGCCCG	44	1	Su(Ste)		Su(Ste)
134	UCUCAUCGUUGUAGAACAAGCCCGA	43	1	Su(Ste)		Su(Ste)
135	AUCCUCAACUGUGAGGGAUCUCAA	43	no match			
136	AAUCUCAUCGUCGUAGAACAAGCCCGA	43	2	Su(Ste)		Su(Ste)
137	UAAAGAAUCCUGACAAGAUCCCGGGC	43	0	transposon	LTR	invader1
138	UGUUUCAUCGUUAGACGGCUCGGGCC	43	1	AT-chX		AT-chX-B
139	CCGAUCUCCUGAGUCGGACUCCGAGC	42	0	transposon	LINE	R1
140	UGUUUCAUCGUUAGACGGCUCGGA	42	1	AT-chX		AT-chX-B
141	UCUCAUCGUCGUAGAACAAGCCCC	42	1	Su(Ste)		Su(Ste)
142	GUAACUUCGGGAUAAGGAUUGGCUCU	42	0	ncRNA		rRNA
143	GUCGCGAUCCCCAGUGAGUAUGUU	42	0	transposon	Low_complexity	F-element
144	GCCGAACGUAGUCUCUCGAGAGCGA	41	0	transposon	LTR	invader1
145	AAAACCUGAAGAUGAUCCUGAGUCGCA	41	0	transposon	LINE	Helena
146	UAUGCUCUCGAAUGUAUGCCCGAUA	39	0	transposon	LINE	R1-element
147	UUCGGCUUUCUGACGAGAACUCGAGC	39	0	no annotation		
148	UCCGAUCUCCUGAGUCGGACUCCGAGC	39	0	transposon	LINE	R1
149	UCCGAGGAACCAGUUGAUCCAGCUC	39	2	Su(Ste)		Su(Ste)
150	UAAAGAAUCCUGACAAGAUCCCGG	38	0	transposon	LTR	invader1
151	UUAUCGACUCGUGUACCUGGUUAAU	38	0	transposon	LTR	micropia
152	UCUCAUCUCGUAGAACAAGCCCGA	38	no match			
153	UCUCAUCGUCGUAGAACAAGCCCGAAA	37	1	Su(Ste)		Su(Ste)
154	AUCGUCUCGGGGAUGAAAUCUCAUCGUCGUAGAACAAGCCCGA	37	no match			
155	UCUCAUCGUCGUAGGACAAGCCCG	36	1	Su(Ste)		Su(Ste)
156	UUAAUGCCGGAAGCGAUCCCACUG	36	0	AT-chX		AT-chX-B
157	UCUAGGCUGAGCGGAGAACCAGAGU	36	0	transposon	Low_complexity	F-element
158	CUCAACUGUGAGGAUCCUCAA	36	no match			
159	UUUUUCAUCGUUAGACGGCUCGGGC	36	1	AT-chX		AT-chX-B
160	UAACAAGAUGCAUACGAUGUGGCU	36	0	transposon	DNA	1360
161	UCUCAUCGUCGUAGAACAGCCCGA	35	no match			
162	GAUCCUCAACUGUGAGGGGAUCCUCAA	35	no match			
163	UUAUAAGGAUGAUGAUACCGACUAAA	35	0	no annotation		
164	AACUUCGGGAUAAGGAUUGGCUCU	35	0	ncRNA		rRNA

165	UGUUUCAUCGUUAGACGGCUGGGGC	35	1	AT-chX		AT-chX-B
166	UCUCAUCGCCGUAGAACAAGCCCGA	35	1	Su(Ste)		Su(Ste)
167	CAGACACAGAGGAGAGCAGAUGAGAGCA	35	no match			
168	GUAAUUGGGCCCGAACCCUUCUGGUG	35	1	transposon	LINE	baggins
169	CUCAUCGUCGUAGAACAAGCCCGA	35	0	Su(Ste)		Su(Ste)
170	GUAGAUCUUCACCGUUGACUUGGCC	35	0	Su(Ste)		Su(Ste)
171	UUCUCAUCGUCGUAGAACUAGCCCGA	34	0	Su(Ste)		Su(Ste)
172	CCGAACGUACCACGGUGCCCCAGUGA	34	0	transposon	Low_complexity	F-element
173	AUCGUCAUCGUCGUAGAACAAGCCCGGA	34	no match			
174	UGUUUUCAUCGUUAGACGGCUCGGG	34	2	no annotation		
175	CCGAACGUACCACGGUGCCCCAGU	34	0	transposon	Low_complexity	F-element
176	CAACACAGAGGAGAGCAGAUGAGAGCA	34	no match			
177	UCUCAUCGUCGUAGAACAUGCCCGA	33	1	Su(Ste)		Su(Ste)
178	UGUUUCAUCGUUAGACGGCUAGGGC	33	1	AT-chX		AT-chX-B
179	AAGGUGGAAAUGAAGAAAGUACA	33	no match			
180	AUUCUCAUCGUCGUAGAACAAGCCCG	33	1	Su(Ste)		Su(Ste)
181	UCUGAUGACUCUUCGACAUAUACAUG	33	2	no annotation		
182	UAGAUCUUCACCGUUUACUUGCCCC	32	1	Su(Ste)		Su(Ste)
183	CAGCACACAGAGGAGAGCAGAUAGAGCA	32	no match			
184	UCCGAUCUCCUGAGUCGGACUCCGAGCU	32	0	transposon	LINE	R1
185	UUGACGGAGCUCACGUACCCCGUGG	32	no match			
186	UCUCAUCGUCGUAGAACAAGCCCCG	32	2	Su(Ste)		Su(Ste)
187	UACUCAUCGUCGUAGAACAAGCCCGA	32	1	Su(Ste)		Su(Ste)
188	UCUCAUCGUCGUAGAAAAAGCCCG	31	1	Su(Ste)		Su(Ste)
189	UAGAUCUUCACCGUUGACUAGCCCA	31	1	Su(Ste)		Su(Ste)
190	UCUCAUCGUCGUAGAACAAGCCUGA	31	1	Su(Ste)		Su(Ste)
191	UCCAACUGAAGGAGUACGAAUGAUC	31	0	transposon	LINE	R1-element
192	CCCGAAGCGGACCAGCCCCAAUUAUU	30	0	coding gene		CG6446-RA|Sema-1b
193	UGUUCCAUCGUUAGACGGCUCGGGC	30	1	AT-chX		AT-chX-B
194	CUCUCAUCGUCGUAGAACAAGCCCGA	30	1	Su(Ste)		Su(Ste)
195	UGUUUCAUCGUUAGACGGCUCGGGUU	29	2	AT-chX		AT-chX-B
196	UGUCUCAUCGUUAGACGGCUCGGGC	29	1	AT-chX		AT-chX-B
197	UGGACCUUUUCCCGAUGAGCCAGU	29	0	transposon	LINE	G
198	UCUGUUUCAUCGUUAGACGGCUCGGG	29	0	AT-chX		AT-chX-B
199	CAAGAUAUGACCAAAGCAAAGACACU	29	0	transposon	LINE	Cr1a
200	UGUUUCAUCGUUAUACGGCUCGGGC	29	1	AT-chX		AT-chX-B
201	UCCGAGGAACCAAUUGAUCCAGCUGC	29	0	Su(Ste)		Su(Ste)
202	UCCGAACGUACCACGGUGCCCCAG	28	0	transposon	Low_complexity	F-element
203	CAGCACACAGAGGAGAACAGAUGAGAGCA	28	no match			
204	UUAUAAGGAUGAUGAUACCGAUUAAA	28	0	no annotation		
205	GACUAGCUGGCCGAGGAUCCGAUUU	28	0	Su(Ste)		Su(Ste)
206	UAGAUCUUCACCGUUGACUAGCCCCCA	28	1	Su(Ste)		Su(Ste)
207	CAGCCACAGAGGAGAGCAGAUGAGAGCA	28	no match			
208	UCUCAUCGUCGUAGAACAGGCCCG	28	1	Su(Ste)		Su(Ste)
209	CAGCACACAGAGGAGAGCAGAUGGAGCA	28	no match			
210	AUCUCAUCGUCGUAGAACUAGCCCGA	27	1	Su(Ste)		Su(Ste)
211	UCUCAUCGUCGUGGAACAAGCCCGA	27	1	Su(Ste)		Su(Ste)
212	CAGCACACAGAGGAGACAGAUGAGAGCA	27	no match			
213	UCUCAUCGUCGUAGAAUAAGCCCGA	27	1	Su(Ste)		Su(Ste)
214	UAGAUCUUCACCGUUGACUUGGCCCAC	27	0	Su(Ste)		Su(Ste)
215	UCAUCGUCGUAGAACAAGCCCGA	27	0	Su(Ste)		Su(Ste)
216	UAGAUCUUCCCCGUUGACUUGGCC	27	1	Su(Ste)		Su(Ste)
217	UAAAUGCCAGAAUUUUAUUGGGG	27	0	transposon	LINE	baggins
218	UCUCACCUCGUAGAACAAGCCCGA	26	2	no annotation		
219	GUAUGACCCCUGGUACUGAAAUGCA	26	0	transposon	DNA	Bari1
220	AUGUUUCAUCGUUAGACGGCUCGGG	26	1	AT-chX		AT-chX-B
221	UCCGAGGAACCAAUUGAUCCAGCU	26	0	Su(Ste)		Su(Ste)
222	CAAGAUAUGACCUAAGCAAAGACAC	26	1	transposon	LINE	Cr1a
223	UCUCACCGUCGUAGAACAAGCCCGA	26	1	Su(Ste)		Su(Ste)
224	GUGUUUCAUCGUUAGACGGCUCGGGC	26	1	AT-chX		AT-chX-B
225	UUCUCGCUGAGAGCUGAGAGCCCGCAU	25	1	ncRNA		alphagamma-element
226	UCUCAUCGUCGUAGAACAAGCCCGAC	25	1	Su(Ste)		Su(Ste)
227	UACAAUCUCCGAGCAAGACCCCGAU	25	0	transposon	LINE	Circe
228	UGCUUCAUCGUUAGACGGCUCGGGC	25	1	AT-chX		AT-chX-B
229	CGAAAUGUUUCAUCGUUAGACGGCCUCGGGC	25	no match			
230	GGAUCAACUGUGAGGGGAUCCUCAA	25	no match			
231	UCUGUUUCAUCGUUAGACGGCUCGGGC	25	0	AT-chX		AT-chX-B
232	UCUCAUUGUCGUAGAACAAGCCCGA	24	1	Su(Ste)		Su(Ste)
233	CGUUUCAUCGUUAGACGGCUCGGGC	24	1	AT-chX		AT-chX-B
234	UAACAAGAUGCAUACGAUGUGGCUC	24	0	transposon	DNA	1360
235	AAAACCUGAAGAUGAUCCUGAGUC	24	0	transposon	LINE	Helena
236	UGGUUUCAUCGUUAGACGGCUCGGGCCG	24	2	no annotation		
237	UGCAUUCGCGUCGAGGCCCAGUAUA	24	2	transposon	LINE	R1-element
238	AAAACCUGAAGAUGAUCCUGAGUCG	24	0	transposon	LINE	Helena
239	UUAAUACCUCGAGUACGUCUGCUCU	23	0	transposon	LTR	Max
240	UGUGAGGGGAUCCUCAA	23	2	coding gene		CG13800-RA|CG13800
241	GCCGAACGUAGUCUCUCGAGAGCGAG	23	0	transposon	LTR	invader1
242	UAAGACCCAAUCUAUGUGCUUAACA	23	0	transposon	LTR	gtwin
243	AUCCUCAACUGGUGAGGGAUCCUCAA	23	no match			
244	CCGAACGUACCACGGUGCCCCAGUG	23	0	transposon	Low_complexity	F-element
245	UCUCAUCGUCGUAGAACAAGCC	23	0	Su(Ste)		Su(Ste)
246	CCUAACGAGGUAGAUGACUCCUGGAA	23	0	transposon	LINE	Cr1a
247	UGUUUCAUCGUUAGACGUCUCGGGC	22	1	AT-chX		AT-chX-B

248	AAAGGUGGAAAUGAAGAAAGUACA	22	no match			
249	UAUGCUCUCGAAUGUAUGCCCGAUAU	22	0	transposon	LINE	R1-element
250	UCUCAUCGUCGUAGAACAAGCUCG	22	1	Su(Ste)		Su(Ste)
251	UAUGCAAUAUUUGUUGCGCCCCGG	22	0	transposon	LTR	opus
252	UUGGACCGAACACUUUGAACCAAG	22	0	Su(Ste)		Su(Ste)
253	UGUUUCAUCGUUAGACGACUCGGG	22	1	AT-chX		AT-chX-B
254	UAAAUGCCAGAAUUUUAUUGGGGU	22	0	transposon	LINE	baggins
255	UCAAAUCCGAGAUACUCUAUUUUUG	22	0	transposon	LTR	invader2
256	CAAGAUAUGACCAAAGCAAAGACA	22	0	transposon	LINE	Cr1a
257	UCUCAUCGUCGUAGAACAAGCCCGGA	22	1	Su(Ste)		Su(Ste)
258	UUCCGAGGAACCAAUUGAUCCAGCUG	22	0	Su(Ste)		Su(Ste)
259	UUCUUUGAAGAUGGCCCACAUAAAGA	21	0	repeat		trf
260	UGUUUCAUCUUUAGACGGCUCGGG	21	1	AT-chX		AT-chX-B
261	GUCUCAUCGUCGUAGAACAAGCCCGA	21	1	Su(Ste)		Su(Ste)
262	UCUCAUCGUCGUAGAACAAAGCCCGA	21	no match			
263	UUCUCAUCAUCGUAGAACAAGCCCGA	21	0	Su(Ste)		Su(Ste)
264	UCUCAUCGUCGUAGAACAAGCUCGA	21	1	Su(Ste)		Su(Ste)
265	CGUAACUUCGGGAUAAGGAUUGGCUCU	20	0	ncRNA		rRNA
266	CAGGCACACAGAGGAGAGCGAGAUGAGAGCA	20	no match			
267	CAGCACAGAGGAGAGCAGAUGAGAGCA	20	no match			
268	GUAUGACCCCUGGUACUGAAAUGCAG	20	0	transposon	DNA	Bari1
269	CCUCAUCGUCGUAGAACAAGCCCG	20	1	Su(Ste)		Su(Ste)
270	CAUGGAUUGGCUUGUUUUGUUCA	20	no match			
271	ACUCAACUGUGAGGGGAUCCUCAA	20	no match			
272	UCCAAAGAGUCGUGUUGCUUGAUAGU	20	0	ncRNA		rRNA
273	UCAACUGUGAGGGAUCCUCAA	20	no match			
274	UGUUUCAUCGUUAGACGGCUUGGGC	19	1	AT-chX		AT-chX-B
275	UUAAUGCCGGAAGCGAUCCCACUGCU	19	0	coding gene		CG41580-RA|CG41580
276	UCUCAUCGUCGUAGACAAGCCCGA	19	no match			
277	UAAGAUCAACGAGAGAUGGCCAU	19	0	transposon	LTR	opus
278	CGAAAUCUCAUCGUCGUAGAACAAGCCCGA	19	no match			
279	UCUGCCCGAUGAGAAGAAUGGCCCGU	19	0	transposon	LINE	RR49783
280	CGAAAUGUUUCAUCGUUAGACGGCUCGGGC	19	no match			
281	GCCGUCGAGAGAUAUCGGAAGAGUU	19	0	ncRNA		rRNA
282	UUCAAGGUUACCCCAGGAUGCUGUGC	19	0	repeat		trf
283	AUGUUUCAUCGUUAGACGGCUCGGGCC	19	2	AT-chX		AT-chX-B
284	UCUCAUCGUCGUAGAAGAAGCCCGAG	19	1	Su(Ste)		Su(Ste)
285	UCUCAUCGUCGUAGAACAAGCCCGAU	19	1	Su(Ste)		Su(Ste)
286	UAACAAGAUGCAUACGAUGUGGCUCC	19	0	transposon	DNA	1360
287	UCCAAAGGACUCGAAAAACUCUUGA	19	2	transposon	LINE	G2
288	UGUUUCAUCGUUAGACGUCUCGGG	19	1	AT-chX		AT-chX-B
289	UCGGCUUUCUGACGAGAACUCGAG	19	0	no annotation		
290	GUAGAUCUUCACCGUUGACUUGGCCC	19	0	Su(Ste)		Su(Ste)
291	CAAGUUCUCGGCUGAUCAGUACCCA	18	0	transposon	LTR	Quasimodo
292	AAUCCUCAACUGUGAGGGAUCCUCAA	18	no match			
293	UUAAUACCUCGAGUACGUCUGCUC	18	0	transposon	LTR	Max
294	UUUUAAACAAUUGAAAGAUAAGACCCA	18	0	transposon	LINE	Cr1a
295	UGUUUCAUCGUUAGAUGGCUCGGGC	18	1	AT-chX		AT-chX-B
296	UCUCAUCGUCGUAGAACAAGCCCCA	18	1	Su(Ste)		Su(Ste)
297	GAUUGCCAUUCUCUGCGUAAGCGAC	18	0	transposon	Low_complexity	F-element
298	UCUCAUCGUCGUAGAUCAAGCCCGA	18	1	Su(Ste)		Su(Ste)
299	UCUCAUCGUCGUAGAACAAGCCAGA	18	1	Su(Ste)		Su(Ste)
300	UUUUAAACAAUUGAAAGAUAAGACCC	18	0	transposon	LINE	Cr1a
301	GUUUCAUCGUUAGACGGCUCGGGC	17	0	AT-chX		AT-chX-B
302	GCUGGCCGAGUAUUUGAUAUUCUU	17	0	Su(Ste)		Su(Ste)
303	AAAAACCUGAAGAUGAUCCUGAGUCGC	17	1	transposon	LINE	Helena
304	ACCACCCUGAUGUGCUUGACGUU	17	no match			
305	AGAUCUUCACCGUUGACUAGCCCCA	17	0	Su(Ste)		Su(Ste)
306	CUGUGAGGGAUCCUCAA	17	2	transposon	LTR	springer
307	GACUAGCUGGCCGAGUAUUUGAUA	17	0	Su(Ste)		Su(Ste)
308	UUCAUCGUCGUAGAACAAGCCCGA	17	0	no annotation		
309	UGCUCUCGAAUGUAUGCCCGAUAU	17	0	transposon	LINE	R1-element
310	UAAGAUUGGACCCAGUACACUGCCU	17	0	transposon	LINE	Doc
311	UACUGAGUCGUUUACCGAGGGCCGA	17	0	transposon	LTR	G4
312	CGUAACUUCGGGAUAAGGAUUGGCU	17	0	ncRNA		rRNA
313	CGGGAUGAGCGGAAUGGCCGAACA	17	0	coding gene		CG33555-RC|btsz
314	UAAAUGCCAGAAUUUUAUUGGGGUU	16	0	transposon	LINE	baggins
315	GAUCUUCACCGUUGACUAGCCCCA	16	0	Su(Ste)		Su(Ste)
316	CCGAGGAACCAAUUGAUCCAGCUGC	16	0	Su(Ste)		Su(Ste)
317	UUAUACCCAGGUGAUUACCACCUA	16	2	Su(Ste)		Su(Ste)
318	CCCUGAUGAAGAAGAGAUGCCCAGUGA	16	0	transposon	LTR	ninja-Dsim-like
319	ACUGUGAGGGGAUCCUCAA	16	no match			
320	CAGCACACAGAGGAGAGCAGAUGAGACA	16	no match			
321	UUCUUUGAAGAUGGCCCACAUAAAG	16	0	repeat		trf
322	UGUUUCAUCGCUAGACGGCUCGGGC	16	1	AT-chX		AT-chX-B
323	AUCUCAACUGUGAGGGGAUCCUCAA	16	no match			
324	UCUCAUCGUCCUAGAACAAGCCCGA	16	1	Su(Ste)		Su(Ste)
325	CGUUGACCAAUUUCCGAACAUAC	16	0	transposon	LTR	DM88
326	AAAAUCUCAUCGUCGUAGAACAAGCCCGA	16	no match			
327	UGUUUCAUCGUUAGACGGCUCGGGCA	16	1	AT-chX		AT-chX-B
328	UUUUCAUCGUUAGACGGCUCGGGC	16	1	AT-chX		AT-chX-B
329	CAGCACACAGAGGAGAGCAGAUGAGAGC	16	no match			
330	GCCCCGUACUCUGUGGUGACCAA	16	0	transposon	LINE	baggins

331	UAGAUCUUCACCGUUGACUAGCCCCC	16	1	Su(Ste)		Su(Ste)
332	CAAGAUAUGACCAAAGCAAAGACACUA	16	0	transposon	LINE	Cr1a
333	UGUCAUCGUCGUAGAACAAGCCCGA	16	1	Su(Ste)		Su(Ste)
334	GAUCUCCUGAGUCGGACUCCGAGC	16	0	transposon	LINE	R1
335	AAACCUGAAGAUGAUCCUGAGUCGC	16	0	transposon	LINE	Helena
336	UUGAUACGAAUGAGACCCGCAGUAA	16	0	repeat		trf
337	UAGAUCUUCACCGUUGACUAGCCCAA	16	1	Su(Ste)		Su(Ste)
338	AUCUCAUCGUCGUAGAAGAAGCCCGA	15	2	Su(Ste)		Su(Ste)
339	UCCGAACGUACCACGGUGCCCCAGUG	15	0	transposon	Low_complexity	F-element
340	UCGCAUCGUCGUAGAACAAGCCCG	15	0	Su(Ste)		Su(Ste)
341	CUCUCAUCGUCGUAGAACAACGCCCG	15	no match			
342	AGAAGCACCGUUUCCCGUCCGAUCAA	15	no match			
343	UAGCUGGCCGAGUAUUUGAUAUUCUU	15	0	Su(Ste)		Su(Ste)
344	UCUUUCAUCGUUAGACGGCUCGGGC	15	1	AT-chX		AT-chX-B
345	UCUCAUCGUUAGACGGCUCGGG	15	1	AT-chX		AT-chX-B
346	UUCGCAUCGUCGUAGAACAAGCCCGA	15	0	Su(Ste)		Su(Ste)
347	UGUUUCAUCUUAGACGGCUCGGGC	15	no match			
348	CUGUUUCAUCGUUAGACGGCUCGGG	15	0	AT-chX		AT-chX-B
349	CAGCACACAGAGGAUAGCAGAUGAGAGCA	15	no match			
350	UCUCAUCGUCGUGAACAAGCCCGA	15	no match			
351	UGUUUCAUUGUUAGACGGCUCGGGC	15	1	AT-chX		AT-chX-B
352	UCUCAUCGUCGUAGAAGAAGCCCCGA	15	no match			
353	UAGAAGAACAUUGAAGCGCUUGAC	15	0	Su(Ste)		Su(Ste)
354	UGUUCAUCGUUAGACGGCUCGGGC	15	2	AT-chX		AT-chX-B
355	AUCAACUGUGAGGGGAUCCUCAA	14	no match			
356	AAUCCUCAACUGUGAGGGGAUCCUCAA	14	no match			
357	UAGAUCUUCACCGUUGACUAGCCCCU	14	1	Su(Ste)		Su(Ste)
358	UUAAUGCCGGAAGCGAUCCCACU	14	0	AT-chX		AT-chX-B
359	UGGACUUUUUUCCGAUGAGCCAGUG	14	0	transposon	LINE	G2
360	GCACGGACCUGGAGAACAACCAGA	14	0	transposon	LTR	ninja-Dsim-like
361	UUGGACCGAACACUUUGAACCAAGUAU	14	0	Su(Ste)		Su(Ste)
362	GUAGAUCUUCACCGUUGACUUGCCC	14	0	Su(Ste)		Su(Ste)
363	UCUGACGAGAACUCGAGCCGCGUA	14	0	no annotation		
364	UGUUUCAUCGUUAGACGGCUCGGGUC	14	2	AT-chX		AT-chX-B
365	AUCUCAUCGUCGUAGAACAAGCCCCG	14	no match			
366	CUCAUCGUCGUAGAACAAGCCCGAGG	14	0	Su(Ste)		Su(Ste)
367	UCCUGUUGAUGUGAGGGCCCCAAGU	14	0	transposon	LINE	G2
368	AGCACACAGAGGAGAGCAGAUGAGAGCA	14	no match			
369	UAACAAGAUGCAUACGAUGUGGCUCCA	14	0	transposon	DNA	1360
370	UUGGACCGAACACUUUGAACCAAGU	14	0	Su(Ste)		Su(Ste)
371	UCUCAUCGUCGUAGAACUAGCCCGAG	14	0	Su(Ste)		Su(Ste)
372	UACUGUUUGUAGGUGUACUGACCC	14	0	transposon	LINE	baggins
373	UGUUUCAUCGUUAGACUGCUCGGGC	14	1	AT-chX		AT-chX-B
374	UCUCAUCGUCGUAGAACAAACCCGA	13	1	Su(Ste)		Su(Ste)
375	CAAAGAAGAUGCUGAGGUGCUCGAG	13	0	transposon	LTR	Dm88
376	UGUUCAUCGUUAGACGGCUCGGG	13	2	AT-chX		AT-chX-B
377	UGUGACUGAUCUAGAGUACAAUGUU	13	0	transposon	LTR	DM88
378	UAAGAUUGGACCCAGUACACUGCC	13	0	transposon	LINE	Doc
379	AUCUCAUCGUCGUAGAACAAAGCCCGA	13	no match			
380	UUCUGACGAGAACUCGAGCCGCGUAG	13	0	no annotation		
381	UUCUUGACAACCCGAUCGGACCUC	13	0	transposon	LTR	invader4
382	UCUCAUUCGUAGAACAAGCCCGA	13	no match			
383	GAAAGCACCGUUUCCCGUCCGAUCAA	13	no match			
384	UCUCAUCAUCGUAGAACAAGCCCGAUA	13	1	Su(Ste)		Su(Ste)
385	UGUUUCAUCGUUAGGCGGCUCGGGC	13	1	AT-chX		AT-chX-B
386	CCUCAACUGUGAGGAUCCUCAA	13	no match			
387	UUCUCAUCGUCGUAGAAGAAGCCCGA	13	1	Su(Ste)		Su(Ste)
388	UGAAGCUGAAAUUGUUUAGUUUGAU	13	0	no annotation		
389	GUGAGGGGAUCCUCAA	13	2	transposon	LINE	baggins
390	UAGAUCUUCACCGUUGACUUGCCCCGC	13	0	Su(Ste)		Su(Ste)
391	UUCUCAUCGUCGUAGAACUAGCCCG	13	0	Su(Ste)		Su(Ste)
392	UCUCAUCGUCGUAGAACAACGCCCG	13	no match			
393	AGAUCUUCACCGUUGACUUGGCCCAC	13	0	Su(Ste)		Su(Ste)
394	CAGCAACAGAGGAGAGCAGAUGAGAGCA	13	no match			
395	GGACUAGCUGGCCGAGUAUUUGAU	13	0	Su(Ste)		Su(Ste)
396	UCUGUUUCAUCGUUAGACGGCUCGG	13	0	AT-chX		AT-chX-B
397	GUAGAUCUUCACCGUUGACUAGCCCCA	13	0	Su(Ste)		Su(Ste)
398	AACUUCGGGAUAAGGAUUGGCUCUG	13	0	ncRNA		rRNA
399	UAACGACCAGACAUACGGCAUUAUC	12	1	transposon	LINE	R1-element
400	GUUUCAUCGUUAGACGGCUCGGG	12	0	AT-chX		AT-chX-B
401	UCUCAUCUUCGUAGAACUAGCCCGA	12	1	Su(Ste)		Su(Ste)
402	UCAAUGUUGAUCGAAGAAACGAAUU	12	0	transposon	LINE	G
403	UCUCAUCGUCGUAGAAGAAGCCCAA	12	2	Su(Ste)		Su(Ste)
404	UGUUACAUCGUUAGACGGCUCGGGC	12	1	AT-chX		AT-chX-B
405	ACUGCAUAUCGAGAGGUCCUCGGUU	12	2	ncRNA		tRNA
406	AUCGUCUCGGGGAUGAAAUGUUUCAUCGUUAGACGGCUCGGGC	12	no match			
407	UCUCAUCGUCGUAGCACAAGCCCGA	12	1	Su(Ste)		Su(Ste)
408	AAAGCACCGUUUCCCGUCCGAUCAA	12	no match			
409	GCUUCAUCCAAGCGAACUGUGCCCGA	12	0	transposon	LINE	R1-element
410	AAAAGGUGGAAAUGAAGAAAGUACA	12	no match			
411	UAUCCAAUCCGACGCAACCGAAAACA	12	no match			
412	UGUUUCAUCGUUAGACGGUUCGGGC	12	1	AT-chX		AT-chX-B
413	CAACUGUGAGGAUCCUCAA	12	2	no annotation		

414	GUAGAUCUUCACCGUUGACUAGCCC	12	0	Su(Ste)		Su(Ste)
415	UCUCAUCGUCGUUGAACAAGCCCGA	12	1	Su(Ste)		Su(Ste)
416	AACCGAUAUGGUUGGACCACAAUCUUU	12	0	ncRNA		7SLRNA-RA
417	UGUUUCAUCGUUAGACGGCUCGGGCU	12	1	AT-chX		AT-chX-B
418	UGUUUCGUCGUUAGACGGCUCGGGC	12	1	AT-chX		AT-chX-B
419	GGACAACUGUGAGGGGAUCCUCAA	12	no match			
420	AUAGAUCUUCACCGUUGACUAGCCCCA	12	1	Su(Ste)		Su(Ste)
421	UCUUAUCGUCGUAGAACAAGCCCGA	12	1	Su(Ste)		Su(Ste)
422	GCCCCGUACUCUGUGGUGACCAACAC	12	0	transposon	LINE	baggins
423	AUAGAUCUUCACCGUUGACUAGCCCCCA	11	2	Su(Ste)		Su(Ste)
424	UGUUUCAUCGUCAGACGGCUCGGGC	11	1	AT-chX		AT-chX-B
425	GGAUCCUCAACUGUGAGGGAUCCUCAA	11	no match			
426	UCCGAACGUAUCACGGUGCCCCAGU	11	0	transposon	LINE	F
427	AAAAACCUGAAGAUGAUCCUGAGUC	11	1	transposon	LINE	Helena
428	UUUUCAUCGUUAGACGGCUCGGG	11	1	AT-chX		AT-chX-B
429	UCUCAUCGUCGUAGAACAAGCCG	11	1	Su(Ste)		Su(Ste)
430	UAAGAAGAUGUCCGGACAGCUGG	11	0	Su(Ste)		Su(Ste)
431	AUCUCAUCGUCGUAGAACAAGCCCGAA	11	1	Su(Ste)		Su(Ste)
432	AAACCUGAAGAUGAUCCUGAGUCGCA	11	0	transposon	LINE	Helena
433	AUUCUCAUCGUCGUAGAACAAGCCCGA	11	1	Su(Ste)		Su(Ste)
434	CGUCUCGGGUAAGGAUCCUCAA	11	no match			
435	UGUUUCAUCGUUAGACGACUCGGGCG	11	1	AT-chX		AT-chX-B
436	UCUCAUCGUCUUAGAACAAGCCCG	11	1	Su(Ste)		Su(Ste)
437	UAAGAUUGGACCCAGUACACUGCCUU	11	0	transposon	LINE	Doc
438	UUAUACCCAGGUGAUUACCACUAUUAA	11	0	Su(Ste)		Su(Ste)
439	UCUCAUCGUCGUAGAACUAGCCCUA	11	1	Su(Ste)		Su(Ste)
440	UAGAUCUUCACCGUUGACUACGCCCCA	11	no match			
441	UCUCAUCGUCGUAGAAUAAGCCCG	11	1	Su(Ste)		Su(Ste)
442	UACAAUGAUCUGAUCGGGUUUUUCU	11	0	transposon	LINE	Cr1a
443	UAAAUGCCAGAAUUUUAUUGGGUUC	11	1	transposon	LINE	baggins
444	UCCGGGAGAAAACAAUGAUGGCCAA	11	0	coding gene		CG12717-RA|CG12717
445	AUGUUUUCAUCGUUAGACGGCUCGGG	11	no match			
446	CAGCACGACAGAAGGAGAGCAGAUGAGAGCCA	11	no match			
447	UGUUUCAUCGUUAGACGGCUCGGGAA	11	2	AT-chX		AT-chX-B
448	UCACAUCGUCGUAGAACAAGCCCGA	11	1	Su(Ste)		Su(Ste)
449	UUCAAGGUUACCCCAGGAUGCUGUG	11	0	repeat		trf
450	UGUACUGUUUGUAGGUGUACUGACCCA	11	0	transposon	LINE	baggins
451	UCUCAUAGUCGUAGAACAAGCCCGA	11	1	Su(Ste)		Su(Ste)
452	ACUGUGAGGAUCCUCAA	11	1	no annotation		
453	UCCAAAGAAUGAUGACCCUUGCAUU	11	0	transposon	LINE	RR49783
454	UCUCAUCGUCGUAGUACAAGCCCGA	11	1	Su(Ste)		Su(Ste)
455	UUCUGACGAGAACUCGAGCCGCGUA	11	0	no annotation		
456	AUCGUCUCGGGAGUGAAAUACUCUAUCGUCGUAGAACAAGCCCGA	11	no match			
457	UCCUGUUGAUGUGAGGGCCCCAAGUC	11	0	transposon	LINE	G2
458	UCUCAUCGUCGUAGAACAAGC	11	0	Su(Ste)		Su(Ste)
459	UGUUACUGAUCUAGAGUACAAUGU	11	1	transposon	LTR	DM88
460	ACUUCGGGAUAAGGAUUGGCUCUG	10	0	ncRNA		rRNA
461	UGUUUCAUCGUUAGACGGCUCGUGC	10	1	AT-chX		AT-chX-B
462	UGUUUCAUCGUUAGACGGCACGGGC	10	1	AT-chX		AT-chX-B
463	GGAUCCUCAACUGUGAGGCUCAA	10	no match			
464	AACUGUGAGGAUCCUCAA	10	1	no annotation		
465	AUCCUCAACUGUGAGGAUCCUCAA	10	no match			
466	UGUUUCAUCGUUGACGGCUCGGGC	10	no match			
467	UCAACUGUGAGGGAUCUCAA	10	no match			
468	CAACUGUGAGGGAUCCUCAA	10	no match			
469	AAACUGUGAGGGGAUCCUCAA	10	no match			
470	UCCAACUGAAGGAGUACGAAUGAUCG	10	0	transposon	LINE	R1-element
471	UUAUACCCAGGUGAUUACCACUAUUA	10	0	Su(Ste)		Su(Ste)
472	UACUGUUUGUAGGUGUACUGACCCA	10	0	transposon	LINE	baggins
473	GACGAGGUGGCCGAGUGGUUAAGG	10	0	ncRNA		tRNA
474	UUCCUCGGGAUCAAGGGCAACGAGU	10	0	coding gene		CG40635-RA|CG40635
475	GACUAGCUGGCCGAGUAUUUGAU	10	0	Su(Ste)		Su(Ste)
476	UUCAAGGUUACCCCAGGAUGCAGU	10	1	repeat		trf
477	AAAAACCUGAAGAUGAUCCUGAGUCGCA	10	1	transposon	LINE	Helena
478	UGUUUCAUCGUUAGACGGCCCGGGC	10	1	AT-chX		AT-chX-B
479	UAAACAAGAUGCAUACGAUGUGGC	10	2	transposon	DNA	1360
480	GAUCCUCAGCUGUGAGGGGAUCCUCAA	10	no match			
481	UCAACUGUGAGGGGAUCUCAA	10	no match			
482	UCUCAUCGUCGUAGAAACAAGCCCGA	10	no match			
483	GCCCCGUACUCUGUGGUGACCAAC	10	0	transposon	LINE	baggins
484	UCUCAUCGUCUAGAACAAGCCCGA	10	no match			
485	AUCGUCUCGGGGAUGAAAAUCUCAUCGUCGUAGAACAAGCCCGA	10	no match			
486	AUCGUCAUCGUCGUAGAACAAGCCCGA	10	no match			
487	CCACACAGAGGAGAGCAGAUGAGAGCA	10	no match			
488	UGUUUCAUCGUUAGACGGCUCUGG	10	1	AT-chX		AT-chX-B
489	UCUGACGAGAACUCGAGCCACGUAGCC	10	1	no annotation		
490	UGUUUCAUCGUUAGACGGCUCCGGC	10	1	AT-chX		AT-chX-B
491	UCUCAUCGUCGUAGAAAGCCCGA	10	no match			
492	UGAAUACCUCGGGUACGUCUGCUC	10	0	transposon	LTR	GATE
493	UGGACUUUUUUCCGAUGAGCCAGUGA	10	0	transposon	LINE	G2
494	UGUUUCAUCGUUAGACUGCUCGGG	10	1	AT-chX		AT-chX-B
495	UGUUUCAUCGUUAGACAGCUCGGGC	10	1	AT-chX		AT-chX-B
496	UACUCAUCGUCGUAGAACAAGCCCG	10	1	Su(Ste)		Su(Ste)

497	UCUCAUCGUCGUAGAACAAGCCCGC	10	1	Su(Ste)		Su(Ste)
498	UUUCUCAUCAUCGUAGAACAAGCCCGA	10	0	Su(Ste)		Su(Ste)
499	UCUAGGCUGAGCGGAGAACCAGAGUA	10	1	transposon	Low_complexity	F-element
500	UAGCAUCGUCGUAGAACAAGCCCGA	9	1	Su(Ste)		Su(Ste)
501	UCAACUGUGAGGAUCCUCAA	9	no match			
502	UUCUGCCCGAUGAGAAGAAUGGCCC	9	0	transposon	LINE	RR49783
503	UCUCAUCGUCGUAGAACCAGCCCGA	9	1	Su(Ste)		Su(Ste)
504	UGUUUCAUCGUCGUAGAACAAGCCCGA	9	1	no annotation		
505	CCAACUGUGAGGGGAUCCUCAA	9	no match			
506	GAUCUCCUGAGUCGGACUCCGAGCU	9	0	transposon	LINE	R1
507	CUCUCAUCGUCGUAGAACAACGCCCGUA	9	no match			
508	UGUUUCAUCGUUAGACGGCUCG	9	0	AT-chX		AT-chX-B
509	UCGUCAUCGUCGUAGAACAAGCCCG	9	2	Su(Ste)		Su(Ste)
510	UCUCAUCGUCGAAGAACAAGCCCGA	9	1	Su(Ste)		Su(Ste)
511	AAACCUGAAGAUGAUCCUGAGUC	9	0	transposon	LINE	Helena
512	CAGCACACAGAGGAGAGCAGAUGAGA	9	no match			
513	AGAGUGAGGGGAUCCUCAA	9	2	no annotation		
514	GACUAGCUGGCCGAGUAUUUGAUAUUC	9	0	Su(Ste)		Su(Ste)
515	UGUUUCAUCGUUGGACGGCUCGGGC	9	1	AT-chX		AT-chX-B
516	GAUCUUCACCGUUGACUUGGCCCA	9	0	Su(Ste)		Su(Ste)
517	UAAAAUGCCAGAAUUUUAUUGGGGUUC	9	2	transposon	LINE	baggins
518	GGCUCAACUGUGAGGGGAUCCUCAA	9	no match			
519	UAGAUCUUCACCGUUGACUUGCC	9	0	Su(Ste)		Su(Ste)
520	CCGAUCUCCUGAGUCGGACUCCGAG	9	0	transposon	LINE	R1
521	UUUAUACCUGAGCCAGUUUGUUUUG	9	0	transposon	Low_complexity	F-element
522	UCUCAUCGACGUAGAACAAGCCCGA	9	1	Su(Ste)		Su(Ste)
523	CAGAUCUUCACCGUUGACUAGCCCC	9	1	Su(Ste)		Su(Ste)
524	CACAGAGGAGAGCAGAUGAGAGCA	9	no match			
525	UUCUCAUCGUCGUAGAAGAAGCCCG	9	1	Su(Ste)		Su(Ste)
526	UUCCUCGGGAUCAAGGGCAACGAGUU	9	0	coding gene		CG40635-RA|CG40635
527	UUCUCAUCGUCGUAGAACAAGCCCAA	9	1	Su(Ste)		Su(Ste)
528	UUUCUCAUCGUUAGACGGCUCGGGC	9	2	AT-chX		AT-chX-B
529	UCUCGAUCGUCGUAGAACAAGCCCGACG	9	no match			
530	CAGAUACCCGUUACUCAGCUAGUGC	9	0	transposon	LINE	INE-1
531	UAUAAGGAUGAUGAUACCGACUAA	9	0	no annotation		
532	UUCGGCUUUCUGACGAGAACUCGAG	9	0	no annotation		
533	UCUCAUCGUGUAGAACAAGCCCGA	9	no match			
534	UUCACUCGUUCACAGAAAUAUGACUU	9	0	Su(Ste)		Su(Ste)
535	UCCGGGAGACACCAAUGCUGGCCAAGU	9	0	coding gene		pseudogene
536	CUUUAAACAAUUGAAAGAUAAGACCCA	9	1	transposon	LINE	Cr1a
537	AUCUCAUCUUCGUAGAACAAGCCCGA	9	2	Su(Ste)		Su(Ste)
538	AACCGAUAUGGUUGGACCACAAUCUU	9	0	ncRNA		7SLRNA-RA
539	UCUCAUCGUCGUAGAACAAGGCCCGA	9	no match			
540	UCUCAUCGCCGUAGAACAAGCCCG	9	1	Su(Ste)		Su(Ste)
541	UCUCAUCGUUAGACGGCUCGGGC	9	1	AT-chX		AT-chX-B
542	UACUGACCCAGUCGGAUCUAUUGUAC	9	0	transposon	LINE	baggins
543	UAACUUCGGGAUAAGGAUUGGCUCU	8	0	ncRNA		rRNA
544	CACCGUGGAGACGGCUUUAGGACCUG	8	0	transposon	LTR	aurora
545	GCCCCGUACUCUGUGGUGAUCAACA	8	1	transposon	LINE	baggins
546	CACGCGGGAGACCGGGGUUCAAUUC	8	0	ncRNA		tRNA
547	AUCGUCUCGGGGAUGAAAAUGUUUCAUCGUUAGACGGCUCGGGC	8	no match			
548	UCUCAUCGUCGAAGUACAAGCCCGA	8	2	Su(Ste)		Su(Ste)
549	UCAACUGUGAGGGGGAUCCUCAA	8	no match			
550	UUCUUUGAAGAUGGCCCACAUAAAGAG	8	0	repeat		trf
551	UUCCGAGGAACCAAUUGAUCCAGCU	8	0	Su(Ste)		Su(Ste)
552	GUGAUGUCAACGUUUGAUCCCCUG	8	0	transposon	LTR	ninja-Dsim-like
553	UUCUCAUCGUCGUAGAACUAGCCCAA	8	1	Su(Ste)		Su(Ste)
554	CAGCUGUGAGGGGAUCCUCAA	8	2	no annotation		
555	AAAAUGUUUCAUCGUUAGACGGCUCGGGC	8	no match			
556	AUCUCAUCAUCGUAGAACAAGCCCGA	8	1	Su(Ste)		Su(Ste)
557	ACUGACCCAGUCGGAUCUAUUGUAC	8	0	transposon	LINE	baggins
558	CAGCACACAGAGGAAAGCAGAUGAGAGCA	8	no match			
559	UCAUCGUCGUAGAACAAGCCCG	8	0	Su(Ste)		Su(Ste)
560	CCCAAGUACAAGCCCGGCGAAAGACC	8	0	repeat		trf
561	UGAGUGCUUGAUAUGACCAGCUUGGACC	8	0	Su(Ste)		Su(Ste)
562	CAGCACGCAGAGGAGAGCAGAUGAGAGCA	8	no match			
563	UGGACGGAGCGAUGUUCGGGCCCAG	8	0	repeat		trf
564	CAUCUCGCUUGAAAUGAUGAUGAC	8	0	no annotation		
565	AAGCACCGUUUCCCGUCCGAUCAA	8	no match			
566	UCUCAUCGUCGUAGAAGUAGCCCGA	8	1	Su(Ste)		Su(Ste)
567	CCCAACUGUGAGGGGAUCCUCAA	8	no match			
568	UGUUUUCAUCGUUUAGACGGCUCGGGC	8	no match			
569	UAAAUGCCAGAACUUUAUUGGGGU	8	0	transposon	LINE	baggins
570	CCCGUGUGUCAUUAAGGUCUUGAACC	8	0	transposon	LINE	baggins
571	UCUCAUCGUCGUAGAACAGCCCG	8	no match			
572	UAGAUCUUCACCGUUGACUUGCCCCC	8	0	Su(Ste)		Su(Ste)
573	UCGAUCUGGGCACCACCUACUCCUG	8	0	coding gene		CG18743-RA|Hsp70Ab
574	ACUUCGGGAUAAGGAUUGGCUCU	8	0	ncRNA		rRNA
575	UGGUUCAUCGUUAGACGGCUCGGG	8	1	AT-chX		AT-chX-B
576	UCUCAUCGUCGUAGAAGAACGCCCGA	8	no match			
577	UCCGCAUGUGAAUGUUGGCCCCAGA	8	0	no annotation		
578	AUGAUGACCGAAAUUUGGAAAAACGGA	8	1	repeat		ARS406_DM
579	CAACUGUGAGGGGGAUCCUCAA	8	no match			

580	UACCAGACUGGAUGCAACCCUUAU	8	0	transposon	LINE	Rt1b
581	UGUACUGUUUGUAGGUGUACUGACCC	8	0	transposon	LINE	baggins
582	UUAAACAAUUGAAAGAUAAGACCCA	8	0	transposon	LINE	Cr1a
583	UAUCAUCGUCGUAGAACAAGCCCGA	8	0	Su(Ste)		Su(Ste)
584	ACUCAUCGUCGUAGAACAAGCCCGA	8	1	Su(Ste)		Su(Ste)
585	UCUCAUCUCGUAGAACAAGCCCG	8	no match			
586	UCUCGAUCGUCGUAGAACAAGCCCGA	8	no match			
587	CCUAACGAGGUAGAUGACUCCUGG	8	0	transposon	LINE	Cr1a
588	UGCUCUCGAAUGUAUGCCCGAUAUA	8	0	transposon	LINE	R1-element
589	UAGACUAAUCGGACAGGUGACUCG	8	0	transposon	LTR	GATE
590	UGUUUCAUCGUUAGACGGCUCGGGG	8	1	AT-chX		AT-chX-B
591	CGUUUCAUCGUUAGACGGCUCGGG	8	1	AT-chX		AT-chX-B
592	UAAUGUCCCUUAAAUCAGCACUCGU	8	0	no annotation		
593	UGAACGUGUCCUGAAAGUAGUUGG	8	0	Su(Ste)		Su(Ste)
594	UCUGCCCGAUGAGAAGAAUGGCCCG	8	0	transposon	LINE	RR49783
595	UGUUUCAUCGUUAGACGGCUCUGGC	8	1	AT-chX		AT-chX-B
596	AUUGCGGACCGGAGAAGACGUUGAA	8	0	no annotation		
597	UCUCUUCGUCGUAGAACAAGCCCGA	8	1	Su(Ste)		Su(Ste)
598	UACAUACCUGGGAAGACUUCGGAAGU	8	0	transposon	LINE	Cr1a
599	UCGUCAUCGUCGUAGAACAAGCCCGG	8	no match			
600	UGUCUCAUCGUUAGACGGCUCGGG	8	1	AT-chX		AT-chX-B
601	UGUUUCAUCGUUAGACGGCUCGGUC	8	1	AT-chX		AT-chX-B
602	UCCGCUAGCUUAUUUCUCGGGAUGAA	8	2	transposon	LINE	baggins
603	CAGCACACAGAGGAGAGCAGAUGAGAA	8	no match			
604	UAGAUCUUUACCGUUGACUAGCCCCA	7	1	Su(Ste)		Su(Ste)
605	UAAAUGCCAGAAUUUUAUUGGGGUUC	7	2	transposon	LINE	baggins
606	UCUGACGAGAACUCGAGCCACGUAG	7	1	no annotation		
607	UAGAUCUUCACCGUUGACUAGCCCCACA	7	0	Su(Ste)		Su(Ste)
608	UUCAAGCCCGGGUAUUCUGAAGCCAA	7	0	repeat		trf
609	UCUCAUCGUCGUAGAACAAGCCCUAGG	7	1	Su(Ste)		Su(Ste)
610	UGUAUGACCCCUGGUACUGAAAUGCA	7	0	transposon	DNA	Bari1
611	CCCGUACUCUGUGGUGACCAACACA	7	0	transposon	LINE	baggins
612	CUAGAUCUUCACCGUUGACUUCGGCCCA	7	no match			
613	UCCGAGUAACCAAUUGAUCCAGCUG	7	1	Su(Ste)		Su(Ste)
614	ACUGCAUUCGCGUCGAGGCCCAG	7	0	transposon	LINE	R1-element
615	UGUUUCAUCGUUAGACGCUCGGGC	7	no match			
616	AACUGUGAGGGAUCCUCAA	7	no match			
617	AGAAAGCACCGUUUCCCGUCCGAUCAA	7	no match			
618	UAUUAUACCCAGGUGAUUACCACUAU	7	0	Su(Ste)		Su(Ste)
619	UGGAUUUUUUUCCGUAUGAGCCAGUG	7	no match			
620	UUCUAAUCGUCGUAGAACAAGCCCGA	7	1	Su(Ste)		Su(Ste)
621	CCCGUACUCUGUGGUGACCAACAC	7	0	transposon	LINE	baggins
622	AUCCUCAACUUGAGGGAUCCUCAA	7	no match			
623	UAUUUCAUCGUUAGACGGCUCGGGC	7	1	AT-chX		AT-chX-B
624	CAGCACACAGAGGGAGCAGAUGAGAGCA	7	no match			
625	CAUGACCUUCGAGGAGCUGCGCGGGCA	7	0	coding gene		CG4226-RC|GluRIIC
626	UUUCAUCGUUAGACGGCUCGGG	7	0	AT-chX		AT-chX-B
627	UAGAUCUUCACCGUUGACUACCCCCA	7	1	Su(Ste)		Su(Ste)
628	UAACAAGAUGCAUGCGAUGUGGC	7	1	transposon	DNA	1360
629	UCUCAUCGUCGUAGAUAAGCCCGA	7	no match			
630	UCUCAUCGUCGUAGAACAAGCCCGAAG	7	0	Su(Ste)		Su(Ste)
631	GUCGCUACCAGACGGGAUUACUCUG	7	0	transposon	LTR	ninja-Dsim-like
632	UUAUAAGGAUGAUGAUACCGAUUA	7	0	no annotation		
633	CUUAGACAUCCACUCUGUUGGCCA	7	0	transposon	LINE	F
634	GCACACAGAGGAGAGCAGAGAGAGCA	7	no match			
635	UCUCAUCGUCGUAGAACAAUCCCG	7	1	Su(Ste)		Su(Ste)
636	UCUCAUCUUCGUAGAACAAGCCCAA	7	2	Su(Ste)		Su(Ste)
637	UCUCAUCGUCGUAGAACAACCCCGAA	7	1	Su(Ste)		Su(Ste)
638	AGAUCUUCACCGUUGACUUGGCCCA	7	0	Su(Ste)		Su(Ste)
639	UUCUCAUCGUCGUAGAACAAGCCCCGA	7	2	Su(Ste)		Su(Ste)
640	UCCGAGGAACCAAUCGAUCCAGCUG	7	1	Su(Ste)		Su(Ste)
641	UGUUUCUAUCGUUAGACGGCUCGGGC	7	no match			
642	UCUCAUCGUCGUAGAACGAGCCCG	7	1	Su(Ste)		Su(Ste)
643	GAUCAACUGUGAGGGGAUCCUCAA	7	no match			
644	CAAGCCCGGGUAUUCUGAAGCCAAGA	7	0	repeat		trf
645	GGAUCUCAACUGUGAGGGGAUCCUCAA	7	no match			
646	UCUCAUCGUCGUAGAACAAGCCCGGG	7	1	Su(Ste)		Su(Ste)
647	UACAAUCUCCGAGCACGACCCCGAUC	7	0	transposon	LINE	Circe
648	UCUCAUCGUCGUAGAGCAAGCCCG	7	1	Su(Ste)		Su(Ste)
649	UAUGGCCCGAUCCAUAAAUCCAACU	7	0	no annotation		
650	UGCAUAUCAUUCACGCCACGCUCCGACC	7	0	Su(Ste)		Su(Ste)
651	GUUCUCAUCGUCGUAGAACAAGCCCGA	7	1	Su(Ste)		Su(Ste)
652	UCCACCGGGUUUUGGCACUUUGCCGC	7	0	transposon	LINE	R1
653	UCUCAUCGUCGUAGAACAAGCCCCGAGG	7	no match			
654	UCCGAGGAACCAAUUGAUUCAGCU	7	0	Su(Ste)		Su(Ste)
655	GGUACACCGACUUCUUGAAUUGGGAC	7	1	transposon	LINE	R1-element
656	AAAUCUCAUCGUCGUAGAACAAGCCCGA	7	no match			
657	UGUUUCAUCGUUAGACGGCUCAGGC	7	1	AT-chX		AT-chX-B
658	UGAGAUCCUCAACUGUGAGGUCAA	7	no match			
659	UCUGUUUCAUCGUUAGACUACUCGGGAA	7	no match			
660	UUGUUAAGGACGAGAAUCUACCCCC	7	0	transposon	LTR	diver2
661	CUGUUUCAUCGUUAGACGGCUCGGGC	7	0	AT-chX		AT-chX-B
662	ACAACUGUGAGGGGAUCCUCAA	6	no match			

663	UUAAUGCCGUAAGCGAUCCCACUGC	6	1	AT-chX		AT-chX-B
664	CAAGAUAUGACCAAAGCAAAGACAU	6	0	transposon	DNA	1360
665	UCUCAACGUCGUAGAACAAGCCCGA	6	1	Su(Ste)		Su(Ste)
666	AAAACCUAGAAGAUGAUCCUGAGUCGC	6	no match			
667	UGUUUUAUCGUUAGACGGCUCGGGC	6	1	AT-chX		AT-chX-B
668	CAGCACACAGAGGAGAGCAAUGAGAGCA	6	no match			
669	UGUUUCAUCCUUAGACGGCUCGGGC	6	1	AT-chX		AT-chX-B
670	UGGGAACACCGCGUGUUGUUGGCC	6	0	ncRNA		5SrRNA
671	CAGGCACACAGAGGAGAGCGAGUGAGAGA	6	no match			
672	UCAUCUUCUUCUCGACGUUGACCAGC	6	0	transposon	LTR	ninja-Dsim-like
673	UGUUUCAGAGCGUUGGCCAGUUUCC	6	0	transposon	Low_complexity	F-element
674	AAAUCCUCAACUGUGAGGGAUCCUCAA	6	no match			
675	UAAGAUCAACGAGAGAUGGCCAUU	6	0	transposon	LTR	opus
676	UCGCAUCGUCGUAGAACAAGCCCGAGG	6	0	Su(Ste)		Su(Ste)
677	UGUUACAUCGUUAGACGGCUCGGG	6	1	AT-chX		AT-chX-B
678	GAUCCUCAACUGUAAGGGGAUCCUCAA	6	no match			
679	CCCCGCGGAAAGACGGAAAGACCC	6	no match			
680	GACUAGCUGGCCGAGUAUUUGAUAUU	6	0	Su(Ste)		Su(Ste)
681	CGCAAGCCGGGCGUCAAGAUGGAC	6	0	coding gene		CG14476-RA|CG14476
682	CAGCACACAGAGGAGAGCAGAUGAGAC	6	no match			
683	CAGCACACAGCGGAGAGCAGAUGAGAGCA	6	no match			
684	UGUGGACCUUAUACCCGGGCGGGAC	6	1	no annotation		
685	UGUUUCAUGUUAGACGGCUCGGGC	6	no match			
686	CAGUACACAGAGGAGAGCAGAUGAGAGCA	6	no match			
687	UCUAAUCGUCGUAGAACAAGCCCGA	6	1	Su(Ste)		Su(Ste)
688	UACCGAGGGCCGAGUACAUAUCGAG	6	0	transposon	LTR	G4
689	UUUCUCAUCUUCGUAGAACAAGCCCG	6	1	Su(Ste)		Su(Ste)
690	UGAUCCUCAACUGUGAGGGGAUCCUCAA	6	no match			
691	UCUGUAUUCCGUACGUCUAGAUUGG	6	0	transposon	LINE	G2
692	UAAAUCUCAUCGUCGUAGAACAAGCCCGA	6	no match			
693	GCCACAGAGGAGAGCAGAUGAGAGCA	6	no match			
694	GGUCAACUGUGAGGGGAUCCUCAA	6	no match			
695	UGGACCGAACACUUUGAACCAAGUA	6	0	Su(Ste)		Su(Ste)
696	UCUCAUCGUUAGAACAAGCCCGA	6	no match			
697	ACAAGAUAUGACCAAAGCAAAGACAC	6	0	transposon	DNA	1360
698	UCUCAUCGUCGUAGAACAAGCCCGACUGUAACCUCGGGUCAAU	6	no match			
699	GAACACCGGCUUCAGGAUCACGUC	6	0	Su(Ste)		Su(Ste)
700	CUUCGGGAUAAGGAUUGGCUCUG	6	0	ncRNA		rRNA
701	CUCAUCGUCGUAGAACAAGCCCG	6	0	Su(Ste)		Su(Ste)
702	UCAAAUCCGAGAUACUCUAUUUUUGU	6	0	transposon	LTR	invader2
703	UCUCAUCGUCGUAGAACAAGCCCCGACUGUAACUCGGGGUCAAU	6	no match			
704	UCUCAUCGUCGUUGAACAAGCCCG	6	1	Su(Ste)		Su(Ste)
705	UGAUGACCGAAAUUUGGAAAAACG	6	0	repeat		ARS406_DM
706	UAGAUCUUCACCGUUGACUAUCCCCA	6	1	Su(Ste)		Su(Ste)
707	UGAAUUUGGCCAGUAUUUCAUCUC	6	0	transposon	LTR	DM88
708	UUUGUUGAUCAGUUUACACAUUAU	6	0	transposon	LINE	Cr1a
709	GAAGCAUUCCGAAAGUUCCGAGUACA	6	0	ncRNA		ncRNA
710	CAGCACACAGAGGAGAGCAGAUGAAGCA	6	no match			
711	UCUCAUCGUCGUAGGAACAAGCCCGA	6	no match			
712	UAUCUAGAGACCGAAGAUAGUUGA	6	0	transposon	LTR	opus
713	GCUACCGGGCGUGAGCAUCCACAAUGC	6	0	transposon	LTR	ninja-Dsim-like
714	CAAAGGUGGAAAUGAAUAAAGUACA	6	no match			
715	UCCAGGCCCAUCUGGUUGAACGUG	6	0	Su(Ste)		Su(Ste)
716	CAGCACCCAGAGGAGAGCAGAUGAGAGCA	6	no match			
717	UGUCCCAACGUACUGAUCCUGGCC	6	0	coding gene		CG31453-RA|CG31453
718	GGAUCCUCAACUGUGAGGGAAUCCUCAA	6	no match			
719	UCUCAUCGUCGUAGAUCAAGCCCG	6	1	Su(Ste)		Su(Ste)
720	CAAAUCUGAACGUACCACGGUGCCCCAGUG	6	no match			
721	UGUUUCAUCGUCAGACGGCUCGGG	6	1	AT-chX		AT-chX-B
722	UCGGCUUUCUGACGAGAACUCGAGC	6	0	no annotation		
723	CGUUGACCAAUUUCCGAACAUACGA	6	0	transposon	LTR	Dm88
724	UGUUUCAGAGGAGAGCAGAUGAGAGCA	6	no match			
725	CUCAACUGUGAGGGGAUCUUCAA	6	no match			
726	UUAAUGCCGGAAGCGAUCCCACUGU	6	1	AT-chX		AT-chX-B
727	UGUUUCAGAGCGUUGGCCAGUUU	6	0	transposon	Low_complexity	F-element
728	UUUCUCAUCGUCGUAGAAGAAGCCCGA	6	1	Su(Ste)		Su(Ste)
729	UCUCAUCGUCGUAGAACAAGUCCGAG	6	1	Su(Ste)		Su(Ste)
730	UGGACUUUUUUCCGAUGAGCCAG	6	0	transposon	LINE	G2
731	UUCACACGCACAAGGUCCCCGGUUC	6	0	ncRNA		tRNA
732	UCCGAGGAACCAAUCGAUCCAGCU	6	1	Su(Ste)		Su(Ste)
733	UGUUUCAUCGUUAGGCGGCUCGGG	6	1	AT-chX		AT-chX-B
734	UAAAUGCCAGAACUUUAUUGGGGUGC	6	1	transposon	LINE	baggins
735	UGUUUCAUAGUUAGACGGCUCGGGC	6	1	AT-chX		AT-chX-B
736	UGUUUCAUCGUUAGACGGCUCGGGGC	6	2	AT-chX		AT-chX-B
737	UAAUGUCCCUUAAAUCAGCACUCG	6	0	no annotation		
738	GAUGAACCCCAGGAAGCUCACGCU	6	0	transposon	LTR	springer
739	UACAAUCUCCGAGCACGACCCCGA	6	0	transposon	LINE	Circe
740	UAGAUCUUCACCGUUGACUUGCCCCAU	6	1	Su(Ste)		Su(Ste)
741	UCUCAUCGUCGUAGAACUAGCCCCGA	6	no match			
742	UCUCACCGUCGUAGAACAAGCCCG	6	1	Su(Ste)		Su(Ste)
743	UUAUAAGGAUGAUGAUACCGACUAAAG	6	0	no annotation		
744	CAUUCGGUGGCGAAGUUGGCUCAGA	6	0	transposon	LTR	297
745	UAAAGAAUCCUGACAAGAUCCCG	6	0	transposon	LTR	invader1

746	UGUUUCAUCGUUAGACGGCUCGA	6	1	AT-chX		AT-chX-B
747	UGGACUUUUUCCGAUGAGCCAGU	6	no match			
748	UCUCAUCGUCGUAGAACAAGCCCGACG	6	1	Su(Ste)		Su(Ste)
749	UAGAUCUUCACCGUUGACUUGGCCCC	6	1	Su(Ste)		Su(Ste)
750	UCAUCUUCUUCUAGACGUUGACCA	6	1	transposon	LTR	ninja-Dsim-like
751	UCUCAUCGUAGAACAAGCCCGA	6	2	Su(Ste)		Su(Ste)
752	UGUUUCCGGAUUCUCGACAGCCG	6	2	ncRNA		alphagamma-element
753	UCCUCAUCGUCGUAGAACAAGCCCGA	6	1	Su(Ste)		Su(Ste)
754	UCUCAUCGUCGUAGAACAAGCCCGAAU	5	1	Su(Ste)		Su(Ste)
755	UUUUAAAACAAUUGAAAGAUAAGACCC	5	1	transposon	LTR	copia
756	CCCAUCUGGUUGAACGUGUCCUGAA	5	0	Su(Ste)		Su(Ste)
757	CACACAGAGGAGAGCAGAUGAGAG	5	no match			
758	GGAUCCUCAACUGUGAGGGGACCUCAA	5	no match			
759	AUCAUCUUCUUCUCGACGUUGACCA	5	1	transposon	LTR	ninja-Dsim-like
760	UUUAUACCUGAGCCAGUUUGUUUU	5	0	transposon	Low_complexity	F-element
761	UCUCAUCGUCGUAUAACAAGCCCG	5	1	Su(Ste)		Su(Ste)
762	GGAUCCUCACUGUGAGGGGAUCCUCAA	5	no match			
763	UGUACUACCCCUUCAUGACUCAAAGAU	5	0	transposon	LINE	Fw2
764	UAACUGUGAGGGGAUCCUCAA	5	no match			
765	CAGCACACAGAGGAGAGCGAGAUGAGAGCA	5	no match			
766	UGUGAGGAGGAUCCUCAA	5	2	coding gene		CG3848-RC|trr
767	UGUUUCAUCGUUAGACGGCUCGGAC	5	1	AT-chX		AT-chX-B
768	UAGAUCUUCACCGUUGACUACCCC	5	1	Su(Ste)		Su(Ste)
769	UCUCAUCGUGUAGAACAAGUCCGA	5	no match			
770	UGGUUUCAUCGUUAGACGGCUCGGGC	5	1	no annotation		
771	UCUCAUCGUCGCAGAACAAGCCCG	5	1	Su(Ste)		Su(Ste)
772	CACCACACAGAGGAGAGCAGAUGAGAGCA	5	no match			
773	UCCGAUGUUUUGAAACUGCGUGACCC	5	0	transposon	LTR	flea
774	UUCGGCUUUCCGACGAGAACUCGAG	5	0	no annotation		
775	UCUCAUCAUCGUAGAACUAGCCCGA	5	1	Su(Ste)		Su(Ste)
776	UCAACUGUGAGGGGAUCCUCAAC	5	no match			
777	UGGUUCAUCGUUAGACGGCUCGGGC	5	1	AT-chX		AT-chX-B
778	UGCGAGGAGGCCCGUAUUACUGUCAC	5	0	transposon	LTR	ninja-Dsim-like
779	UAUAAUCGCCCCGAAGGUUGCCCUCUG	5	no match			
780	UCUCAUCGUCGUAGAACGAAGCCCGA	5	no match			
781	CAGCACACAGAGGAAGCAGAUGAGAGCA	5	no match			
782	UCUCAUCGUCGUAGAACAAGCCCCGAA	5	no match			
783	GGAUCCAACUGUGAUCAA	5	2	coding gene		CG32540-RB|CCKLR-17D3
784	UCUCAUCGUCGUAGAACUAAGCCCGA	5	no match			
785	UCUCAUCGUCGUAAAACAAGCCCG	5	1	Su(Ste)		Su(Ste)
786	UGUUUCAUCGUUAGACGGCUCGGGCCUGUAACUCCGGGUCAAU	5	no match			
787	UUCUCAUCGUCGUAGAACAAGCCC	5	0	Su(Ste)		Su(Ste)
788	UCUAGACUCGAAAAUUCGUUUUAUGU	5	0	Su(Ste)		Su(Ste)
789	CUCUCAUCGUCGUAGAACAAGCCCG	5	1	Su(Ste)		Su(Ste)
790	UCAUCUUCUUCUCGACGUUGACUA	5	1	transposon	LTR	ninja-Dsim-like
791	UCUCAUCGUCGUAGAACAACCCCG	5	1	Su(Ste)		Su(Ste)
792	UACAAUGAUCUGAUCGGGUUUUUCUG	5	0	transposon	LINE	Cr1a
793	UCCGAGGAACCAAUUGAUCCCGCUG	5	1	Su(Ste)		Su(Ste)
794	ACUCAUAUCCGAGGCCCUGUAAUUG	5	0	ncRNA		rRNA
795	CUCACUGUGAGGGGAUCCUCAA	5	no match			
796	UCCUUCUGAAGAGUUGUUGGCACGU	5	0	transposon	LTR	GATE
797	UCCCAUCGUCGUAGAACAAGCCCGA	5	1	Su(Ste)		Su(Ste)
798	UGACUCUACGUGACGGAACCGGAG	5	0	transposon	LINE	Rt1b
799	AGAUCCAACUGUGAGGGGAUCCUCAA	5	no match			
800	UAGAUGAGGUAUAAGGUUGGCCCA	5	0	transposon	Low_complexity	F-element
801	CAGCACACAGAGGAGAGAGAUGAGAGCA	5	no match			
802	CAGCACACAGACGGAGAGCAGAUGAGACGCA	5	no match			
803	UCCCAUCCUUAUGAUGACCGCUCGAG	5	0	transposon	LTR	HMS-Beagle
804	CAGCACCAGAGGAGAGCAGAUGAGAGCA	5	no match			
805	GUCUCAUCGUCGUAGAACAAGCCCG	5	1	Su(Ste)		Su(Ste)
806	UAAAUAAUCCUGACAAGAUCCCGGG	5	1	transposon	LTR	invader1
807	UAGAUCUUUACCGUUGACUUGGCCCA	5	0	no annotation		
808	UCUCAUCGUCGUAACAAGCCCGA	5	no match			
809	UGUUUCAUCGUUAGACGGCUGGGC	5	2	AT-chX		AT-chX-B
810	UCCAUCGUCGUAGAACAAGCCCGA	5	1	no annotation		
811	CCUGAUGAAGAGGAGAUGCCCAGUGA	5	1	transposon	LTR	ninja-Dsim-like
812	CAAGAUAUGACCAAAGCAAAUACAC	5	1	transposon	LINE	Cr1a
813	CAGAUCUUCACCGUUGACUUGCCC	5	1	Su(Ste)		Su(Ste)
814	UAGAUCUUCACCGUUGACUAGCCCCGA	5	2	Su(Ste)		Su(Ste)
815	AUAGAUCUUCACCGUUGACUUGCCCC	5	1	Su(Ste)		Su(Ste)
816	UUAAAGGGCCAAGAUUGAUUCCGUUUU	5	no match			
817	UCUCAUCGUCGUGGAAGAAGCCCG	5	2	Su(Ste)		Su(Ste)
818	UGUUUCAUCGUUAGACGGCUCGGCCCG	5	2	AT-chX		AT-chX-B
819	UCGGCUUUCCGGCGAGAACUCGAGC	5	0	no annotation		
820	UUUUCUCAUCGUCGUAGAACAAGCCCGA	5	0	Su(Ste)		Su(Ste)
821	GCACACGGAGGAGAGCAGAUGAGAGCA	5	no match			
822	GCCCCGUACUCUGUGGUGACCAACACAC	5	0	transposon	LINE	baggins
823	UCUUCACCGUUGACUUGGCCCACACA	5	0	Su(Ste)		Su(Ste)
824	UCAUCGUCGUAUAACAAGCCCGA	5	1	Su(Ste)		Su(Ste)
825	UUUCUGACGAGAACUCGAGCCGCGUAG	5	0	no annotation		
826	AAAACCUGAAGGUGAUCCUGAGUCGCA	5	1	transposon	LINE	Helena
827	AUCGUCUCGGGGAUGAACAGCACACAGAGGAGAGCAGAUGAGAGC	5	no match			
828	UCUCAUCUUCGUAGAAGAAGCCCGA	5	2	Su(Ste)		Su(Ste)

829	UAAGUAAGCCCGAAGAUUGUCCA	5	0	transposon	LTR	opus
830	UCUCAUCGUCGUAGAAGAAGCCCGAA	5	1	Su(Ste)		Su(Ste)
831	UCUCAUCAUCGUAGAACAAGCCCGAA	5	1	Su(Ste)		Su(Ste)
832	CAGCACACAGAGGAGAGCAGAUGAGGAGCA	5	no match			
833	AUCCUCAACUGUGAGAUCCUCAA	5	no match			
834	GCCCGGCUAGCUCAGUCGGUAGAGCA	5	0	ncRNA		tRNA
835	UCUUCACCGUUGACUUGCCCCGCA	5	0	Su(Ste)		Su(Ste)
836	GGAUCCUCAACUGUGAGG	5	2	coding gene		CG13506-RA|CG13506
837	UCUGACGAGAACUCGAGCCGCGUAG	5	0	no annotation		
838	AUAACUGUGAGGGGAUCCUCAA	5	no match			
839	CAGCACACAGAGGAGAGCAGAUGAGAGCACUGUAACUCGGGGUCA	5	no match			
840	CAAGAUAUGACCAAAGCAAAGACAA	5	1	transposon	LINE	Cr1a
841	GUAGAUCUUCACCGUUGACUUGCCCC	5	0	Su(Ste)		Su(Ste)
842	UGACCGUAGCACCGAACGAGCUCCU	5	0	transposon	LINE	Rt1b
843	AAACCUAGUUACUUACGGGCUCGACA	5	0	Su(Ste)		Su(Ste)
844	AACUUCGGGAUAAGGAUUGGCUC	5	0	ncRNA		rRNA
845	UAGAUCUCCACCGUUGACUAGCCCC	5	1	Su(Ste)		Su(Ste)
846	AGCAGUUUCCAGACGCAGCCCGGCC	5	0	transposon	LTR	ninja-Dsim-like
847	AUCUCAUCGUCGUAGAAGAAGCCCCGA	5	no match			
848	ACAACCAGAGAUCUAGAGGGCCCAA	5	0	no annotation		
849	AUCAUCGUCGUAGAACAAGCCCGA	5	0	Su(Ste)		Su(Ste)
850	UUUUCGCAUCGUCGUAGAACAAGCCCG	5	0	Su(Ste)		Su(Ste)
851	UGUACUACCCCUUCAUGACUCAAAG	5	0	transposon	LINE	Fw2
852	UCCUCAACUGUGAGGAUCCUCAA	5	no match			
853	UCUGACGAGAACUCGAGCCGCGUAGCC	5	0	no annotation		
854	CGAAAUCUCAUCGUCGUAGAACAGCCCGA	5	no match			
855	AUAGAUCUUCACCGUUGACUUGGCCCA	5	1	Su(Ste)		Su(Ste)
856	UCUCAUCGUCGUAGAACAAGCCCCUC	5	no match			
857	UCUCAUCAUCGUAACAAGCCCGA	5	no match			
858	UGCAUAUCAUUCACGCCAAGCUCCGA	5	0	Su(Ste)		Su(Ste)
859	UGUUUCAUCGUUAGACGGCUGGC	5	2	AT-chX		AT-chX-B
860	GGGAACUGUGAGGGGAUCCUCAA	5	no match			
861	CCUCAACUGUGAGGGGGAUCCUCAA	5	no match			
862	UCAAUCGCGGGUGCAAUCGAAUGUUGC	5	0	coding gene		pseudogene
863	UCGCUACCAGACGGGAUUACUCUG	5	0	transposon	LTR	ninja-Dsim-like
864	UAGAUCUUCACCGUUGACUUGCCCCG	5	0	Su(Ste)		Su(Ste)
865	UUCACGUCUGGCGAUUUUGACUCAUU	5	0	transposon	LTR	mdg3
866	UGAGCCGACGAGAUGCCCAGGGCGUG	5	0	transposon	LTR	ninja-Dsim-like
867	UCUCAUCGUCGUAGAACUAGCCCGAGG	5	0	Su(Ste)		Su(Ste)
868	UCGUCAUCGUCGUAGAAGAAGCCCGGA	5	no match			
869	UAAUGCCGGAAGCGAUCCCACUGC	5	0	AT-chX		AT-chX-B
870	GUCGCUACCAGACGGGAUUACUCU	5	0	transposon	LTR	ninja-Dsim-like
871	UCUCAUCGUCGUGGAACAAGCCCG	5	1	Su(Ste)		Su(Ste)
872	UAACAAGAUGCAUACGAUGUGG	5	0	transposon	DNA	1360
873	GCCAGGCCCGUAUAACGUUAAUGAUU	5	0	ncRNA		rRNA
874	CCUAACUGUGAGGGGAUCCUCAA	5	no match			
875	UAAGAAGAUGUCCGGACAGCUGGG	5	0	Su(Ste)		Su(Ste)
876	UCUCAUCGUCGUAGACCAAGCCCGA	5	1	Su(Ste)		Su(Ste)
877	UCUCAUCAUCGUAGAACAAGCCCGAG	5	1	Su(Ste)		Su(Ste)
878	GGAUCAACUGUGAGGGAUCCUCA	5	no match			
879	UGCAAGGUUACCCCAGGUGCUGU	5	no match			
880	AUCCUAACUGUGAGGGGAUCCUCAA	5	no match			
881	CAAUGAGAUAGCAUUUGGCGAGAGU	5	0	transposon	LTR	diver2
882	UAUGACCGAAAUUCGGAAAAACGGA	5	0	repeat		SAR_DM
883	UCUUCACCGUUGACUUGGCCCACAC	5	0	Su(Ste)		Su(Ste)
884	UGUUUCACCGUUAGACGGCUCGGG	5	1	AT-chX		AT-chX-B
885	UCUCAUCGUCGUAGAACAAGCCCGACUGUA	4	no match			
886	UCGACUACAUUCAGGACCCGUUCAA	4	0	coding gene		CG40635-RA|CG40635
887	UUAUAAGGAUGAUGAUACCGACUA	4	0	no annotation		
888	AUUUUCAUCGUUAGACGGCUCGGGC	4	2	AT-chX		AT-chX-B
889	UCCAACUGAAGGAGUACGAAUGAUCGC	4	0	transposon	LINE	R1-element
890	UAGACUAAUCGGACAGGUGACUCGG	4	0	transposon	LTR	GATE
891	UGUUUCAGAGCGUUGGCCAGUUUC	4	0	transposon	Low_complexity	F-element
892	UUCGCAUCGUCGUAGAACAAGCCCG	4	0	Su(Ste)		Su(Ste)
893	UAAUGACCGAAAUUAGGGAAUCAG	4	0	repeat		SAR_DM
894	UAGAGAGUGAUGCCGAGUAUAAAAG	4	0	repeat		trf
895	UCUCGUCGUCGUAGAACAAGCCCCGA	4	no match			
896	UCUCAUCGUCGUGAACAAGCCCG	4	no match			
897	UACAGUAUUGGUCCAAGAACGCUGCCU	4	0	transposon	LINE	G
898	UUCAUCUUCUUCUCGACGUUGACCA	4	0	transposon	LTR	ninja-Dsim-like
899	UCUGACGAGAACUCGAGCCGCGUAGC	4	0	no annotation		
900	CUCCUGUGAGGGGAUCCUCAA	4	no match			
901	UCAACGUCCCCAAGAACUCCCGGU	4	0	transposon	LINE	G2
902	UCUUUCAUCGUUAGACGGCUCGGG	4	1	AT-chX		AT-chX-B
903	AUAAACUGUGAGGGGAUCCUCAA	4	no match			
904	UGUUUCAUCGUUAACGGCUCGGGC	4	no match			
905	UCAACCGGACGGGCCUAGAGUUAUUC	4	0	coding gene		CG40635-RA|CG40635
906	UGUACUGUUUGUAGGUGUACUGGCCCAG	4	0	transposon	LINE	baggins
907	AAACUUCGGGAUAAGGAUUGGCUCU	4	1	ncRNA		rRNA
908	AGCCCCGUACUCUGUGGUGACCAACA	4	0	transposon	LINE	baggins
909	GAAGAUCAUUCCCUUCGCAACCCG	4	0	transposon	LTR	ninja-Dsim-like
910	GGAUCCUCAACUGUGAGGGGAUCUCAA	4	no match			
911	UUCAACUGAAGGAGUACGAAUGAUC	4	0	transposon	LINE	R1-element

912	UAGGUUAUGGGCCCAGUAAACAUU	4	0	ncRNA		alphagamma-element
913	UCUCAUCGUCGUGAAGAAGCCCGA	4	no match			
914	UAGGUUAUGGGCCCAGUAAACAUUU	4	0	ncRNA		alphagamma-element
915	UCAAUACGACCCGGUUUAUUAAAA	4	0	transposon	LTR	mdg1
916	UAAGUAAGCCCGAAGAUUGUCCAAUGA	4	0	transposon	LTR	opus
917	UAGAUCUUACCGUUGACUUGGCCC	4	no match			
918	UCUCAUCGUCGUAGAACAAGCCCCAA	4	1	Su(Ste)		Su(Ste)
919	GAUCUUCACCGUUGACUUGCCCCA	4	0	Su(Ste)		Su(Ste)
920	UUCUCAUCGUCGUAGAACAAGCCCGU	4	1	Su(Ste)		Su(Ste)
921	GGGAUCCUCAACUGUGAGGGAUCCUCAA	4	no match			
922	AUCUCAUCGUCGUAGAACAACGCCCGA	4	no match			
923	GACGCGAGAUGUGAAAUUCUUGGACC	4	1	ncRNA		rRNA
924	CGUUGACCAAUUUCCGAACAUACGCCA	4	0	transposon	LTR	DM88
925	GAUCUCGGAUCUUGUUGUUGACCAG	4	0	transposon	LINE	G3
926	UCUUCACCGUUGACUUGGCCCAC	4	0	Su(Ste)		Su(Ste)
927	UCCGAGGAACCCAUUGAUCCAGCUG	4	0	no annotation		
928	AUAGAUCUUCACCGUUGACUAGCCCC	4	1	Su(Ste)		Su(Ste)
929	GUUCAACCGGACGGGCCUAGAGUUA	4	0	coding gene		CG40635-RA|CG40635
930	AUCGUCUCGGAUGAAAUCUCAUCGUCGUAGAACAAGCCCGA	4	no match			
931	UCCCUGAUGAAGAAGAGAUGCCCAUUU	4	2	transposon	LTR	ninja-Dsim-like
932	AUCUCGUCGUCGUAGAACAAGCCCGA	4	2	Su(Ste)		Su(Ste)
933	GCAAAGGUGGAAAUGAAGAAAGUACA	4	no match			
934	AUCGGUCUCGGGAAUAGAAAAUGUUUCAUCGUUAGACGGCUCGGG	4	no match			
935	UCCGAGGAACCAAUUGAUUCAGCUGC	4	1	Su(Ste)		Su(Ste)
936	UCUCAUCGUCGUAGAACAAGCCUGAG	4	1	Su(Ste)		Su(Ste)
937	UCCAACUGUGAGGGGAUCCUCAA	4	no match			
938	UCUCAUCGUCGUUAGAACAAGCCCGA	4	no match			
939	UGUUUCAUCGUUAGACGGCUCGGGCCG	4	1	AT-chX		AT-chX-B
940	GAUCUCAACUGUGAGGGGAUCCUCAA	4	no match			
941	UGUACUGACCCAGUCGGAUCUAUU	4	0	transposon	LINE	baggins
942	UGUUUCAUCGUUAGACGGCACGGG	4	1	AT-chX		AT-chX-B
943	UAAGGUUGGCCCAAGAACGCUGCCU	4	0	transposon	Low_complexity	F-element
944	CAGCACACAGAGGGGAGCAGAUGAGAGCA	4	no match			
945	UGUUUCAUCGUUAGACCGACUCGGGC	4	no match			
946	UCGCUACCAGACGGGAUUACUCU	4	0	transposon	LTR	ninja-Dsim-like
947	GGAUCCUCAACUGUGAGGGGGUCCUCAA	4	no match			
948	UCAUCUUCUUCUCGACGCUGACCA	4	1	transposon	LTR	ninja-Dsim-like
949	UUCCUCGGAAUCAAGGGCAACGAGU	4	0	Su(Ste)		Su(Ste)
950	UGUUUAAUCGUUAGACGGCUCGGGC	4	1	AT-chX		AT-chX-B
951	UGUUUAAUCGUUAGACGGCUCGGG	4	1	AT-chX		AT-chX-B
952	UCAUCUUCUUCUCGACGUUGACCAGCC	4	0	transposon	LTR	ninja-Dsim-like
953	UGUUUCAUCGUUAGACGACUCGGGCU	4	2	AT-chX		AT-chX-B
954	UAGACUAAUCGGACAGGUGACUCGGA	4	0	transposon	LTR	GATE
955	UCAUCUUCUUCUCUACGUUGACCA	4	1	transposon	LTR	ninja-Dsim-like
956	CGUUGGACCAAUUUCCGAACAUACGCCUG	4	no match			
957	CUGCACACAGAGGAGAGCAGAUGAGAGCA	4	no match			
958	UCAAAUCCGAGAUACUCAUUUUUGU	4	no match			
959	UCUCAUCGUCGUAGAACUAGCCCGU	4	1	Su(Ste)		Su(Ste)
960	CAAACUGUGAGGGGGAGUCCUCAA	4	no match			
961	GUGCUCUGGACCCUGGACAUUGG	4	0	no annotation		
962	GUAGAUCUUCACCGUUGACUAGCCCC	4	0	Su(Ste)		Su(Ste)
963	CUCAUCUUCUUCUCGACGUUCGACCA	4	no match			
964	CAGAUACCCGUUACUCAGCUAGU	4	0	transposon	LINE	INE-1
965	UACUCAUCGUCGUAGAACUAGCCCGA	4	1	Su(Ste)		Su(Ste)
966	AAAGAACCCGAUUCCUCUUCCAUGAGA	4	0	transposon	LINE	Doc4
967	ACCAAAGUCUCUGAUGGUUUGACA	4	0	ncRNA		pseudogene
968	UGUUUCAUCGUUAGACCGCUCGGGC	4	1	AT-chX		AT-chX-B
969	UUCCUCGGGAUCAAGGGCAACGAGUC	4	1	coding gene		CG40635-RA|CG40635
970	UCUCAUCGUUGUAGAACAAGCCCG	4	1	Su(Ste)		Su(Ste)
971	UCUCAUCGUCGUAGAACUAGC	4	0	Su(Ste)		Su(Ste)
972	UAGAUCUUCACCGUUGACUAGCCCCAU	4	1	Su(Ste)		Su(Ste)
973	AAUCUCAACUGUGAGGGGAUCCUCAA	4	no match			
974	UGAUACCGAGAAUGACAAUGGUUGAU	4	0	transposon	LTR	gypsy3
975	CUCUAGAGAGAGGUGGCUCCAUGC	4	1	transposon	DNA	1360
976	UCCGAGGAAAAAAUUGAUCCAGCAGC	4	0	Su(Ste)		Su(Ste)
977	UGUUUCAUCAUUAGACGGCUCGGGC	4	1	AT-chX		AT-chX-B
978	UCUCAUCGUCGUAGAACAAGCCCGACUA	4	2	Su(Ste)		Su(Ste)
979	UCCCAACUGUGAGGGGAUCCUCAA	4	no match			
980	UAUGACCAGCUUGGACCGAACACUU	4	0	Su(Ste)		Su(Ste)
981	UGUUUCAUCGUUAUACGGCUCGGC	4	2	AT-chX		AT-chX-B
982	UCUUCACCGUUGACUUGCCCCGCAAA	4	0	Su(Ste)		Su(Ste)
983	UCUCAUCGUCGUAGAACAAGCCCAAA	4	1	Su(Ste)		Su(Ste)
984	UGGAUUUUUUUUCCGAUGAGCCAGUG	4	2	transposon	LINE	G2
985	UCAAAGAUCCCCCUGAAGUCCAAUGC	4	0	transposon	LINE	baggins
986	GCAUCCUCAACUGUGAGGGGAUCCUCAA	4	no match			
987	UCGACUAUUUACGAGACCACCCGA	4	0	transposon	DNA	H
988	CGAAAUCCAAGCUGAUGAUCCGGCA	4	0	transposon	LINE	BS
989	GCCCGAUGAGAAGAAUGGCCCGUAAA	4	0	transposon	LINE	RR49783
990	GUAGAUCUACACCGUUGACUAGCCC	4	1	Su(Ste)		Su(Ste)
991	GCAUACAGAGGAGAGCAGAUGAGAGCA	4	no match			
992	UGGAUUUUUUUCCGAUGAGCCAGUG	4	1	transposon	LINE	G2
993	UGGUUUCAUCGUUAGACGGCUCGGGCG	4	1	no annotation		
994	UCAUCUUCUUCUCGACGUUGCCA	4	2	transposon	LTR	ninja-Dsim-like

995	CUGUGAACGAGUGAACUGGCAACAUG	4	0	Su(Ste)		Su(Ste)
996	UGCUUGAUAUGACCAGCUUGGACCG	4	0	Su(Ste)		Su(Ste)
997	CCUCAUCGUCGUAGAACAAGCCCGAG	4	1	Su(Ste)		Su(Ste)
998	UCCAGUCUCCCGACUAUCAGAUACCCG	4	0	transposon	LINE	INE-1
999	UCUCAUCGUCGUAGUACAAGCCCGC	4	2	Su(Ste)		Su(Ste)
1000	AUCGUCAUCGUCGUAGAACAAGCCCGG	4	no match			
1001	AUUCCGAUCUUGGACAUUCCCGAUG	4	0	transposon	LTR	roo
1002	UCUCAUCAUCGUAUAACAAGCCCGA	4	1	Su(Ste)		Su(Ste)
1003	UAGAUCUUCACCGUUGACUAGCC	4	0	Su(Ste)		Su(Ste)
1004	AUAGAUCUUCACCGUUGACUUGCCCCA	4	1	Su(Ste)		Su(Ste)
1005	AUCGUCUCGGAUGAAAUGUUUCAUCGUUAGACGGCUCGGGC	4	no match			
1006	GCAAAGAGAUGGACGCCGUGAGACA	4	0	transposon	LTR	ninja-Dsim-like
1007	CAGCACACAGAGGAGGGCAGAUGAGACA	4	no match			
1008	AUUCAUCGUCGUAGAACAAGCCCGA	4	1	no annotation		
1009	UCUCAUCGUCGUAGAACAAGUCCCGA	4	no match			
1010	GCACACAGAGGAGAGAGAUGAGAGCA	4	no match			
1011	UAAGGAUGAUGAUACCGACUAAAGGCAA	4	0	no annotation		
1012	UCUCAUUGUCGUAGAACAAGCCCG	4	1	Su(Ste)		Su(Ste)
1013	CUCAACUGUGAGGGGAGUCCUCAA	4	no match			
1014	AGCCGAACGUAGUCUCUCGAGAGCGA	4	1	transposon	LTR	invader1
1015	GCCAAAGUACCGUUACGAGUGAGGC	4	0	transposon	DNA	invader2
1016	CAAGAUAUGACCAAAGCAAAGA	4	0	transposon	LINE	Cr1a
1017	UCUCAUCAUCGUAGAACAAGCCCCGA	4	no match			
1018	UGGUUUCAUCGUUAGACGGCUCGGGCCGUGUAACGUCGGGUCAAU	4	no match			
1019	GCCCCGUACUCUGUGGUGACCAACACA	4	0	transposon	LINE	baggins
1020	UAGGCUGAGCGGAGAACCAGAGUUG	4	0	transposon	Low_complexity	F-element
1021	UGUUUCAUCGUUAGACGGCUCGGGCCUGUAACUCGGGGUCAAU	4	no match			
1022	AAUCUCAUCGUCGUAGAACAAGCCCG	4	2	Su(Ste)		Su(Ste)
1023	UUCGAGAAGCGUUUGAUCGAGUGUGG	4	0	transposon	LINE	BS3
1024	UCACGCGGGAGACCGGGGUUCAAUU	4	0	ncRNA		tRNA
1025	CGUUUCAUCGUUAGACGGCUCGGCC	4	2	AT-chX		AT-chX-B
1026	UGUUUCAUCGUUAGACGGCUCGGCU	4	2	AT-chX		AT-chX-B
1027	CAAGAUAUGACCAAAGCAAAGCCAC	4	1	transposon	LINE	Cr1a
1028	UCUCAUCGUCGUAGAACAAGCCCGACUGUAACGUCGGGUCAAU	4	no match			
1029	UUCCGAGGAACCAAUUGGUCCAGCUG	4	1	Su(Ste)		Su(Ste)
1030	CAAGAUGUGACCAAAGCAAAGACUA	4	2	transposon	DNA	1360
1031	UGUUUCAUCGUUUGACGGCUCGGGC	4	1	AT-chX		AT-chX-B
1032	GUAUGACCCCUGGUACUGAAAUGCAGA	4	0	transposon	DNA	Bari1
1033	CUCUCAUCGUCGUAGAACUACGCCCGA	4	no match			
1034	UAGAGAGUGAUGCCGAGUAUAAAA	4	0	repeat		trf
1035	UCCAGGCCCAUCUGGUUGAACGUGU	4	0	Su(Ste)		Su(Ste)
1036	GAGAGGCGCCGUAGUUACCACCCCA	4	0	no annotation		
1037	AUGGUUUCAUCGUUAGACGGCUCGGGCCG	4	no match			
1038	UACAAUCUCCGAGCAAGACCCCGAUCU	4	0	transposon	LINE	Circe
1039	GCCCCGUACUCUGUGGUGACCAACAA	4	1	transposon	LINE	baggins
1040	CAGCACACAGAGGAGAGCAGAUGAGGCA	4	no match			
1041	GCGACACAGAGGAGAGCAGGAUGAGAGCA	4	no match			
1042	AGCCCCGUACUCUGCGGUGACCAA	4	0	transposon	LINE	baggins
1043	AUUAAUACCUCGAGUACGUCUGCUCU	4	1	transposon	LTR	Max
1044	GUCGCGAUCCCCAGUGAGUAUGUUU	4	0	transposon	Low_complexity	F-element
1045	GAUCCCCCUGAAGUCCAAUGCACUG	4	0	transposon	LINE	baggins
1046	ACCGUCAUUCCAGCGCAAGUUGCCCC	4	0	transposon	LTR	ninja-Dsim-like
1047	AGGUGGAAAUGAAGAAAGUACA	4	no match			
1048	CAGCACGACAGAAGGAGAGCAGAUGAGAGCA	4	no match			
1049	UUCUUCUCGACGUUGACCAGCCUC	4	0	transposon	LTR	ninja-Dsim-like
1050	GAAUUACUUACCGAGGGUUGUCCAGG	4	1	Su(Ste)		Su(Ste)
1051	UGGACUUUUUUUCCGAUGAGCCAGU	4	1	transposon	LINE	G2
1052	AUAGAUCUUCACCGUUGACUUGGCCC	4	1	Su(Ste)		Su(Ste)
1053	UAUAAUGCCCCGAAGGUUGCCCUCUG	4	0	transposon	LINE	R1-element
1054	UUCGCAAAAGUUGACCGAACUUGAU	4	0	ncRNA		rRNA
1055	UUCUCGUCGUCGUAGAACAAGCCCGA	4	1	Su(Ste)		Su(Ste)
1056	UUCUCAUCGUCGUAGAACAGGCCCG	4	1	Su(Ste)		Su(Ste)
1057	AUUAUACCCAGGUGAUUACCACUAU	4	0	Su(Ste)		Su(Ste)
1058	ACCUGAGAGGGUGACCGGCCACAGGGGCA	4	no match			
1059	CACACAGAGGAGACAGAUGAGAGCA	4	no match			
1060	AUUAUACCCAGGUGAUUACCACUAUU	4	0	Su(Ste)		Su(Ste)
1061	UCUCAUCUUCGUAGAACAAGCCCGAG	4	1	Su(Ste)		Su(Ste)
1062	UCUCAUCUUCGUAGAACAGCCCGA	4	no match			
1063	UCUCAUCGUCGUAGAACAAGCCUG	4	1	Su(Ste)		Su(Ste)
1064	UGUUCCAUCGUUAGACGGCUCGGG	4	1	AT-chX		AT-chX-B
1065	UUUCAUCGUUAGACGGCUCGGGC	4	0	AT-chX		AT-chX-B
1066	UCUCAUCGUCGUAGAACAAGUCCG	4	1	Su(Ste)		Su(Ste)
1067	UAACAACGUCAACCUGAGGCAUUGG	4	0	transposon	LTR	diver2
1068	UAAUGCCCCGAAGGUUGCCCUCUGU	4	0	transposon	LINE	R1-element
1069	UGUUUCAUCGUUAGACGGGCUCGGGC	4	no match			
1070	CGAAAUCUCAUCGUCGUAGAACUAGCCCGA	4	no match			
1071	UGUAUGACCCCUGGUACUAAAUGCAGA	4	no match			
1072	UAGACCUAGAGCAUCCAAAAAACUU	4	0	transposon	LTR	gypsy8
1073	UAAGAUCUUCACCGUUGACUAGCCCCA	4	2	Su(Ste)		Su(Ste)
1074	UAACGAGGUAGAUGACUCCUGGAA	4	0	transposon	LINE	Cr1a
1075	UCCGUGCUCCUGCGAACCAUCACCA	4	0	coding gene		CG4445-RA|pgant3
1076	AUCUCAUCGUCGUAGAACAAGCCCGU	4	2	Su(Ste)		Su(Ste)
1077	UCUUACGAGAACUCGAGCCUUA	4	no match			

1078	UCUCAUCGUCGUACCAAGCCCG	4	no match			
1079	AUCGUCUCGGGACUGAAACUCUCAUCCGUUCGUACGAACAACCGC	4	no match			
1080	UAAUGCCAGAAUUUUAUGGGGU	4	no match			
1081	UACGGUGCCCAAAUUAACAACUCAUG	4	0	ncRNA		rRNA
1082	UGUUUCAUCGUUAGACAGCUCGGG	4	1	AT-chX		AT-chX-B
1083	AAAAAAAAAAAAACUA	4	0	transposon	LTR	qbert
1084	UUAUAAGGAUGAUGAUACCGACUAA	4	0	no annotation		
1085	GAUUACCGAGGGCCGAGUACAUAUCGA	4	0	transposon	LTR	G4
1086	GCACACAGAGGAGAGCAGAUGAGAG	4	no match			
1087	GCUGCGAGAUACCCGUGAACAGCAGUC	4	0	coding gene		CG12734-RA|CG12734
1088	AAAUCUCAUCGUCGUAGAACAAGCCCG	4	no match			
1089	UGCUCUCGAAUGUAUGCCCGAUA	4	0	transposon	LINE	R1-element
1090	CAACCGGACGGGCCUAGAGUUAUUCCCC	4	0	coding gene		CG40635-RA|CG40635
1091	UUCGGCUUUCCGACGAGAACUCGAGC	4	0	no annotation		
1092	GGCUUUCUGACGAGAACUCGAGCC	4	0	no annotation		
1093	GCCAGACAAGCGUCCCGGACUAGAAGC	4	0	transposon	LINE	R1
1094	GUGACUGAUCUAGAGUACAAUGUU	4	0	transposon	LTR	DM88
1095	UAGAUCUUCACCGUUGACUUGCCCCAC	4	0	Su(Ste)		Su(Ste)
1096	UCUUCACCGUUGACUUGGCCCACA	4	0	Su(Ste)		Su(Ste)
1097	UGUCACGCGGGAGACCGGGGUUCAA	4	0	ncRNA		tRNA
1098	CAAGAUAUGACCAAAACAAAGACAC	4	0	transposon	DNA	1360
1099	UAACGACCAGACAUACGCCAUAAUC	4	0	transposon	LINE	R1
1100	UCUCGUCGUCGUAGAACAAGCCCG	4	1	Su(Ste)		Su(Ste)
1101	AUACCUCAACUGUGAGGGAUCCUCAA	4	no match			
1102	CAGCACACAGAGGAGAGCAGACGAGAGCA	3	no match			
1103	GCCAAUUACAUUAGACGAGACGGCCC	3	0	transposon	LTR	opus
1104	GGAUUCAACUGUGAGGGGAUCCUCAA	3	no match			
1105	CCUAUUACAGGAGAUAUUUGGACACGA	3	0	Su(Ste)		Su(Ste)
1106	AGAAAAAAACCUGAAGAUGAUCCUGAGUCGCA	3	no match			
1107	UGUACUGACCCAGUCGGAUCUAUUGU	3	0	transposon	LINE	baggins
1108	UAACGACCAUACAUACGCCAUAAUCU	3	1	transposon	LINE	R1
1109	UUUCUCAUCGUCGUAGAACAAACCCGA	3	1	Su(Ste)		Su(Ste)
1110	UUCAAGGUUACCCCAGGAUGCUG	3	0	repeat		trf
1111	UUCGCUAGUCCGUCCGAUUCUCGAGU	3	0	transposon	LINE	Cr1a
1112	CCAAUGAUUAGCAAGUUAAUUGAACC	3	0	transposon	Unknown	RR50960
1113	UCUCAUCAUCGUAGAACGAGCCCGA	3	1	Su(Ste)		Su(Ste)
1114	UAAAUGCCAGAUUUAUUGGGGUUC	3	no match			
1115	UAAAUUCCAGAAUUUUAUUGGGGU	3	1	transposon	LINE	baggins
1116	UCUCAUCGUCGUAGAACUAGCCCGUA	3	2	Su(Ste)		Su(Ste)
1117	UACUCGAUCGUCGUAGAACAAGCCCGACG	3	no match			
1118	AGAAAAAAACCUGAAGAUGAUCCUGAGUCG	3	no match			
1119	UUACCGGUAUAGCACACUUUUGUA	3	0	coding gene		CG41580-RA|CG41580
1120	UCGAUCUUCACCGUUGACUUGGCCCA	3	1	Su(Ste)		Su(Ste)
1121	UUCCGAGGAACCAAUUGAUUCAGCU	3	0	Su(Ste)		Su(Ste)
1122	GUCUCAUCAUCGUAGAACAAGCCCGAU	3	1	Su(Ste)		Su(Ste)
1123	UCUCAUCGUCGUAUAACAAGCCCAA	3	2	Su(Ste)		Su(Ste)
1124	GUAGAGAGUGAUGCCGAGUAUAAAA	3	0	repeat		trf
1125	UGUUUCUGCCGAACGUAGUCUCGAG	3	2	transposon	LTR	invader1
1126	UACUCAUCAUCGUAGAACAAGCCCGA	3	1	Su(Ste)		Su(Ste)
1127	UUCUCAUCGUCGUAGAACAAGCCCGAG	3	0	Su(Ste)		Su(Ste)
1128	CUCAACUGUGAGGGGGAUCCUCAA	3	no match			
1129	GUCUCGGGAGAAACUCAA	3	1	no annotation		
1130	UCCGGAGAGGGAGCCUGAGAAACG	3	0	ncRNA		rRNA
1131	UCUCAUCAUCGUAGACCAAGCCCGA	3	1	Su(Ste)		Su(Ste)
1132	GUGUUUCAUCGUUAGACGGCUCGGGU	3	2	AT-chX		AT-chX-B
1133	UAUAAGGAUGAUGAUACCGACUAAA	3	0	no annotation		
1134	ACGGUGCCCAAAUUAACAACUCAUG	3	0	ncRNA		rRNA
1135	ACAUGCAAGUCGAACGGCUCUUCGGAG	3	no match			
1136	AGCACCGUUUCCCGUCCGAUCAA	3	no match			
1137	UGUCAUCGUUAGACGGCUCGGGC	3	1	AT-chX		AT-chX-B
1138	UUAUCGACUCGUGUACCUGGUUAAUU	3	0	transposon	LTR	micropia
1139	GGAUCCUAACUGUGAGGGGAUCCUCAA	3	no match			
1140	UGUUUCAUCGUUAGACGCCUCGGGC	3	1	AT-chX		AT-chX-B
1141	UCAUCUUCAUCUCGACGUUGACCAG	3	1	transposon	LTR	ninja-Dsim-like
1142	UGUACUGUUUGUAGGUGUACUGACCCAG	3	0	transposon	LINE	baggins
1143	UUAUCGACUCGUGUACCUGGUUAA	3	0	transposon	LTR	micropia
1144	AUCGUCGUCGGGAUAGAAAUGUUUCAUCGUUAGACGGCUCGGGC	3	no match			
1145	GACUAGCUGGCCGAGGAUCCGAUUUU	3	0	Su(Ste)		Su(Ste)
1146	GGCAACUGUGAGGGGAUCUCAA	3	no match			
1147	UGGACCGAACACUUUGAACCAAGUAU	3	0	Su(Ste)		Su(Ste)
1148	ACUUCAACUGUGAGGGGAUCCUCAA	3	no match			
1149	UCCGAUCUCCCUAGUCGGACUCCGAGC	3	1	transposon	LINE	R1-element
1150	CAUGGAUUGGCUUGUUUGUUCA	3	no match			
1151	CGAAAUCUCAUCGUCGUAGAACCAAGCCCGA	3	no match			
1152	UUCGCCCAAGUUGCUCUCCCUUUG	3	0	transposon	LTR	micropia
1153	UCAUAUGAUCUGAGAAUAUGUUUC	3	0	transposon	LINE	Cr1a
1154	UCUCAUCGUCGUAAAACAAGCCCGA	3	1	Su(Ste)		Su(Ste)
1155	UUCUCAUCGUCGUAGAACAAGCCAGA	3	1	Su(Ste)		Su(Ste)
1156	AUCUCAUCGUCGUAGAACAAGCCCGACUGUAACUCGGUCAAU	3	no match			
1157	AAGAAGCACCGUUUCCCGUCCGAUCAA	3	no match			
1158	UCGUCCAUCCUGACUAGCUGAUCCCCU	3	1	transposon	LTR	flea
1159	AUAAAUGCCAGAAUUUUUAUUGGGGUUC	3	2	transposon	LINE	baggins
1160	UUCUUGACAACCCGAUCGGACCU	3	0	transposon	LTR	invader4

1161	AAAACCUAGAAGAUGAUCCUGAGUCGCA	3	no match			
1162	AUACUCAUCGUCGUAGAACAAGCCCGA	3	2	Su(Ste)		Su(Ste)
1163	UUCAAGGUUACCCCCAGGAUGCUGUGC	3	no match			
1164	UGUUUCAUCGUUAGACGGCUCGGGCCUC	3	no match			
1165	UGGAUCUUCACCGUUGACUUGCCC	3	1	Su(Ste)		Su(Ste)
1166	UGUUUCGAUCGUUAGACGGCUCGGGC	3	no match			
1167	GACACUGCAUUUGCGUCGAGGCCCAG	3	0	transposon	LINE	R1-element
1168	UUAUUGUGCCACUGAGUCGACUAUU	3	0	transposon	LINE	Cr1a
1169	UAGAUUCGGCUAGUUACUCUUCUAA	3	0	transposon	LINE	Circe
1170	CAGCACACAGAGGAGAGCAGAUGAGACGCA	3	no match			
1171	UAAUGCCCCGUAGGUUGCCCUCUG	3	0	transposon	LINE	R1-element
1172	CAAGAUAUGACCAAGGUUCAAGAUUU	3	no match			
1173	CAGCACACAGACGGAGAGCAGAUGAGACGCUA	3	no match			
1174	CUUCAACUGUGAGGGGAUCCUCAA	3	no match			
1175	UGAUUCAUCGUUAGACGGCUCGGGC	3	1	AT-chX		AT-chX-B
1176	UCUCAUCGUCGCAGAACAAGCCCGC	3	2	Su(Ste)		Su(Ste)
1177	GCCGUCGAGAGAUAUCGGAAGAGUUU	3	0	ncRNA		rRNA
1178	CGAAAUGUUUCAUCGUUAGACGGCUCGGG	3	no match			
1179	AAAAUCUCAUCUUCGUAGAACAAGCCCGA	3	no match			
1180	UACAAUGAUCUGAUCGGGUUUUUC	3	0	transposon	LINE	Cr1a
1181	UGCCGAACGUAGUCUCUCGAGAGCGAG	3	0	transposon	LTR	invader1
1182	AAGGUAAUAUCCGAGUCGUUCCCUAGU	3	0	transposon	LTR	opus
1183	CAAGAUAUGACCAAAGCAAAGACUAA	3	2	transposon	DNA	1360
1184	UUUUCUCAUCGUCGUAGAACAAGCCCCGA	3	2	Su(Ste)		Su(Ste)
1185	UGAGGAGGAUCCUCAA	3	1	coding gene		CG3848-RC|trr
1186	AUCUCAUCGUCGUAGAACAAGCCGA	3	no match			
1187	UUCAUCGUCGUAGAACAAGCCCG	3	0	no annotation		
1188	AGGAUCCUCAACUGUGAGGGGAUCCUCAA	3	no match			
1189	GGUCUCAUCGUCGUAGAACAAGCCCGA	3	2	Su(Ste)		Su(Ste)
1190	AAAACCUAGAAGAUGAUCCUGAGUC	3	no match			
1191	AUAGAUCUUCACCGUUGACUUGCC	3	1	Su(Ste)		Su(Ste)
1192	UCUCAUCGCGUAGAACAAGCCCGA	3	no match			
1193	UAGAUCUUCACCUUUGACUAGCCCCA	3	1	Su(Ste)		Su(Ste)
1194	CAGCACACAGAGGAGAGCAGAGAGAGCA	3	no match			
1195	CCGAACGUACCACGGUGCUCCAGUGA	3	1	transposon	Low_complexity	F-element
1196	UCGUCAUCGUACGUAGAACAAGCCCGGA	3	no match			
1197	GAAUGUUGGCCCCAGAAUACGCAGG	3	0	no annotation		
1198	ACCAAAGUCUCUGAUGGUUUGACAA	3	0	ncRNA		pseudogene
1199	UGCUGACGGGGCCCGAAAACACGAGA	3	0	transposon	LTR	opus
1200	ACCGCCGACGGAGACACGAGCCCUG	3	0	coding gene		CG4792-RA|Dcr-1
1201	UAGAUCUCCACCGUUGACUUGGCCC	3	1	Su(Ste)		Su(Ste)
1202	UCGUCAUCGUCGUAGAACAAGCCCGGACUG	3	no match			
1203	ACCUGAUGAAGAGGAGAUGCCCAGUGA	3	2	transposon	LTR	ninja-Dsim-like
1204	UUUCAUCGUCGUAGAACAAGCCCG	3	0	no annotation		
1205	CGAAAUCCUCAACCGUGAGGGAUCCCUCAA	3	no match			
1206	UUGAUAUGGCCAGCUACCACCUACU	3	0	Su(Ste)		Su(Ste)
1207	ACUGCAUUUGCGUCGAGGCCCAGG	3	0	transposon	LINE	R1-element
1208	CAGCGCACAGAGGAGAGCAGAUGAGAGCA	3	no match			
1209	GGUGCAUAUCAUUCACGCCAAGCUCCGA	3	0	Su(Ste)		Su(Ste)
1210	UCUCAUCGUCGUAGAACAAGCCGGA	3	1	Su(Ste)		Su(Ste)
1211	UUCUCAUCGUCGUAGAACGAGCCCGA	3	1	Su(Ste)		Su(Ste)
1212	CAGAUACCCGUUACUCAGCUAGUUG	3	0	transposon	LINE	INE-1
1213	UCUCAUAGUCGUAGAACAAGCCCGAGG	3	1	Su(Ste)		Su(Ste)
1214	UGCGAGGAGGCCCGUAUUACUGUCA	3	0	transposon	LTR	ninja-Dsim-like
1215	CAGACACAGAGGAGAGCAGAUGAGAA	3	no match			
1216	UACUGACCCAGUCGGAUCUAUUGU	3	0	transposon	LINE	baggins
1217	UGAGGACUUGAGCGAUUUAAGUGGC	3	1	Su(Ste)		Su(Ste)
1218	UCUCAUCAUCGUGGAACAAGCCCGA	3	1	Su(Ste)		Su(Ste)
1219	UGGACUUUUUUCCGUAUGAGCCAGU	3	no match			
1220	UUCGGCUUUCCGGCGAGAACUCGAGC	3	0	no annotation		
1221	CCUCAACUGUGAGGGAUCCUCAU	3	no match			
1222	UCCGAGGAACCAGUUGAUCCAGCUG	3	1	Su(Ste)		Su(Ste)
1223	UCUCAUUCGUCGUAGAACAAGCCCGA	3	no match			
1224	UCUCAUCGUCGUAGAACAAGCUCGAG	3	1	Su(Ste)		Su(Ste)
1225	CGAAACCAGGCCCAUCUGGUUGAACGUGU	3	no match			
1226	UAGAUCUUCACCGUUGACUUGCCCCCA	3	0	Su(Ste)		Su(Ste)
1227	UGUUUGUAGGUGUACUGACCCAGUCGG	3	0	transposon	LINE	baggins
1228	AUCCUCAACUGUGGGGGAUCCUCAA	3	no match			
1229	UUGAUGUCGGCUCAUCUCAUCCUGGGG	3	no match			
1230	UCAACCGUCAAGAUCCACUGCCCAC	3	0	repeat		trf
1231	UCUCAUCGUCGUAGAACAAGCCAGU	3	2	Su(Ste)		Su(Ste)
1232	CUAACUGUGAGGGGAUCCUCAA	3	no match			
1233	UGAUGACCGAAAUUUGGAAAACG	3	0	repeat		SAR_DM
1234	UCUCAUCUUCGUAGAACAAGCCCGAA	3	1	Su(Ste)		Su(Ste)
1235	UCGUUACGUGCAUUGGCUCCGCUUGC	3	0	transposon	LTR	diver2
1236	GCCGAAUGUAGUCUCUCGAGAGCGA	3	1	transposon	LTR	invader1
1237	UAGAUCCUCACCGUUGACUAGCCCCA	3	1	Su(Ste)		Su(Ste)
1238	AACCGAUAUGGUUGGACCACAAUC	3	0	ncRNA		7SLRNA-RA
1239	AUCGCCUGAGGAUGAGCCCGCGUU	3	no match			
1240	ACCGAUUUAGAUGCUAAUCCCGAGGA	3	0	transposon	LTR	opus
1241	CAAGAUAUGACCAAAACAAAGCAA	3	0	Su(Ste)		Su(Ste)
1242	CAGCACACAGAGGAGAGCAGAUGAGAGCAA	3	no match			
1243	UAACCGAAGAGCAUCCGUUGUUCAAA	3	1	coding gene		CG14017-RA|CG14017

1244	UGUUUCAUCGUUAAACGGCUCGGGC	3	1	AT-chX		AT-chX-B
1245	CAGCACACAGAGGAGAGUAGAUGAGAGCA	3	no match			
1246	UACUGUUUGUAGGUGUACUGACCCAG	3	0	transposon	LINE	baggins
1247	UGAUCUUCACCGUUGACUAGCCCC	3	1	Su(Ste)		Su(Ste)
1248	CCCCGUACUCUGUGGUGACCGACA	3	1	transposon	LINE	baggins
1249	UAGACCUUCACCGUUGACUAGCCC	3	1	Su(Ste)		Su(Ste)
1250	UCCUCCUGUGAGGGAUCCUCAA	3	no match			
1251	UCCUCGGGAUCAAGGGCAACGAGU	3	0	coding gene		CG40635-RA|CG40635
1252	UGGACUUUUUUUCCGAUGAGCCAGUGA	3	2	transposon	LINE	G2
1253	UUUUCAUCGCUGGAGGACGCCCGGG	3	0	transposon	LINE	R1-element
1254	CCCGUACUCUGUGGUGACCGACACA	3	1	transposon	LINE	baggins
1255	UAGCGGCCGAGGAUCCGAUUUU	3	no match			
1256	ACGGAGCGAUGUUCGGGCCCAGCU	3	0	repeat		trf
1257	CAGCACACAGAGGGAGAGCAGAUGAGAGCA	3	no match			
1258	UGUGACCGAGCGUGGGACCUCGAUAA	3	1	transposon	LINE	Cr1a
1259	AUAAAUGCCAGAAUUUUAUUGGGGUUC	3	0	transposon	LINE	baggins
1260	UAAAUGCCAGAAUUUUAUUGGGGUUCAG	3	0	transposon	LINE	baggins
1261	UAAAAGAAUCCUGACAAGAUCCCGG	3	2	transposon	LTR	invader1
1262	UUCCGGAGAGGGAGCCUGAGAAACG	3	0	ncRNA		rRNA
1263	UCUCAUCAUCGUAGAAGAAGCCCGA	3	1	Su(Ste)		Su(Ste)
1264	UUCACCGUUGACUAGCCCCACACAU	3	0	Su(Ste)		Su(Ste)
1265	UCUCAUCGUCGUAGAACAAGCACGA	3	1	Su(Ste)		Su(Ste)
1266	UAGGUCUUCACCGUUGACUAGCCCCA	3	1	Su(Ste)		Su(Ste)
1267	CAUAUCAUUCACGCCACGCUCCGACC	3	0	Su(Ste)		Su(Ste)
1268	GGCCACUCCCGCCAACUGGACCCCUGG	3	no match			
1269	GCACACAGAGGAGAGCAGAUGAAGCA	3	no match			
1270	CCUCAACUGUGAGGGGAUCCUCAAC	3	no match			
1271	CUCUCAUCGUCGUAGAAGAACGCCCGA	3	no match			
1272	AGUCUCAUCGUCGUAGAACAAGCCCGA	3	2	Su(Ste)		Su(Ste)
1273	UUCUCAUCGUCGUAGAACAAGCCCGG	3	1	Su(Ste)		Su(Ste)
1274	UUCACGUCUGGCGAUUUUGACUCA	3	0	transposon	LTR	mdg3
1275	UGAAUACCUCGGGUACGUCUGCUCUC	3	0	transposon	LTR	GATE
1276	AAACCUGAAGAUGAUCCUGAGUCG	3	0	transposon	LINE	Helena
1277	CGUAACUUCGGGAUAAGGAUUGGCUC	3	0	ncRNA		rRNA
1278	UAGAUCUUUACCGUUGACUUGGCC	3	0	no annotation		
1279	UCUCAUCGUCUAGAACAAGCCCG	3	no match			
1280	ACCUCAUCGUCGUAGAACAAGCCCGA	3	2	Su(Ste)		Su(Ste)
1281	UCCUGUAGUACGAGGUCAUGCCUAG	3	0	transposon	LTR	opus
1282	AUCCUCAACUGUGAGGGAAUCCUCAA	3	no match			
1283	UCCUCAAUGUGAGGGAUCCUCAA	3	no match			
1284	UCUCAUCCUCGUAGAACAAGCCCG	3	1	Su(Ste)		Su(Ste)
1285	AGAUCUUCACCGUUGACUAGCCCC	3	0	Su(Ste)		Su(Ste)
1286	UCUCAUCGUCGUACAACAAGCCCG	3	1	Su(Ste)		Su(Ste)
1287	AUCCUCAACUGUGAGGGGGAUCCUCAA	3	no match			
1288	UCGUCAGUCGUCGUAGAACAAGCCCGGA	3	no match			
1289	UCUCAUCGUUGUAGAACAAGCCCGAGG	3	1	Su(Ste)		Su(Ste)
1290	UGUUUCACCGUUGACUAGCCCCA	3	no match			
1291	UCUGCCCGAUGACAAAAAUGGCCCGU	3	0	transposon	LTR	mdg1
1292	GAAUCUCAUCGUCGUAGAACAAGCCCG	3	no match			
1293	UACAGAGAAGAUUAGCAUGGCCCCUG	3	0	ncRNA		snRNA
1294	UGAGGGGAUCCUCAA	3	1	no annotation		
1295	GCCACACGCAUCCUAGAAUCGACCGACC	3	0	transposon	LTR	flea
1296	UUCAAGUUUACCCCAGGAUGCUGU	3	1	repeat		trf
1297	UGCUUCAUCGUUAGACGGCUCGGG	3	1	AT-chX		AT-chX-B
1298	UUCUCAUCGUCGUAGAACAAGCCCC	3	1	Su(Ste)		Su(Ste)
1299	UUUCGCAUCGUCGUAGAACAAGCCCGA	3	0	Su(Ste)		Su(Ste)
1300	UGUUUCAUCGUUAGCGGCUCGGGC	3	no match			
1301	UCGUUUCAUCGUUAGACGGCUCGGGC	3	1	no annotation		
1302	GAACUUGAGACCGCCCCUGGACAACC	3	0	Su(Ste)		Su(Ste)
1303	ACCGGAGAAGACGUUGAAUGGAUCG	3	0	no annotation		
1304	CACACACAGAGGAGACAGAUGAGAGCA	3	no match			
1305	CAGGCCCAUCUGGUUGAACGUGU	3	0	Su(Ste)		Su(Ste)
1306	UCUCAUCGUCGUAGAACAAGCCCCC	3	2	Su(Ste)		Su(Ste)
1307	UCUCAUCGUCGCAGAAAGCCCGA	3	no match			
1308	UUGAAUUUGGCCAGUAUUUCAUC	3	0	transposon	LTR	DM88
1309	UUCAAGGUUACCCCAGGAUGCUGUGCC	3	0	repeat		trf
1310	CAGGCCCAUCUGGUUGAACGUGUC	3	0	Su(Ste)		Su(Ste)
1311	AGAAAUGUUUCAUCGUUAGACGGCUCGGGC	3	no match			
1312	CAAAGCCUUUCAGACGUUUGACUCUC	3	0	transposon	DNA	transib4
1313	UCUCAUCUUCAUAGAACAAGCCCG	3	2	Su(Ste)		Su(Ste)
1314	UACAAUCUCCGAGCAAGACCCGAUC	3	no match			
1315	UGCUUCAUCGUUAGACGGCUCGCGC	3	2	AT-chX		AT-chX-B
1316	UCCGUCGGGAUGAUUUGUACCGUA	3	0	transposon	Low_complexity	F-element
1317	UGGACCUUUUCCCGAUGAGCCAGUGG	3	0	transposon	LINE	G
1318	CAGCAUACAGAGGAGAGCAGAUGAGAGCA	3	no match			
1319	GUAUAAGGUUGGCCCAAGAACGCU	3	0	transposon	Low_complexity	F-element
1320	AUCUCAUCGUCGUAGAACAAGCCCAA	3	2	Su(Ste)		Su(Ste)
1321	AUCGUCUUCGGGAUGAACAGCACACAGAGGAGAGCAGAUGAGAGC	3	no match			
1322	UCUCAUCGUCGUAGAACAAGCCCGACUGUAAUCGGGUCAAU	3	no match			
1323	AUGGUUGCAACGGGUAACGGGGAAU	3	0	ncRNA		rRNA
1324	UCCUGUAGUACGAGGUCAUGCCUAGA	3	0	transposon	LTR	opus
1325	UCUAAUUCGGCCCGAACCCUUCUGG	3	0	transposon	LINE	baggins
1326	GAGAAUCGCAUCUUAGAGUAAAUGGCC	3	0	transposon	LTR	gypsy8

1327	AUGGCUCUCUCAGUCGCUUCCCGGGA	3	0	ncRNA		rRNA
1328	UGUUUCGAUCGUUAGACGGCUCGGGCCUG	3	no match			
1329	AUAGACCCGCUAGGAGAUGUUGAGAU	3	1	transposon	LTR	roo
1330	UGGGAACACCGCGUGUUGUUGGCCC	3	1	ncRNA		5SrRNA
1331	CAGCCACAGAGGAGAGCAGAGAGAGCA	3	no match			
1332	CUGUAACUCGGGAUAAGGAUUGGCUC	3	no match			
1333	GGAUCCUCAACUGUGAGGGGAUCCUCGA	3	no match			
1334	CAGCACACAGAGGAGAGCAGUGAGAGCA	3	no match			
1335	UAAAUCUCAUCGUCGUAGAACAAGCCCG	3	no match			
1336	UUAAUGCCUGAAGCGAUCCCACUGC	3	1	AT-chX		AT-chX-B
1337	GGAUCCUCAACUGUGAGGGGGAUCCUCAA	3	no match			
1338	AUUCGGCUUUCUGACGAGAACUCGAGC	3	1	no annotation		
1339	GCCGAACGUAGUCUCUCAAGAGCGAG	3	1	transposon	LTR	invader1
1340	GUUCGUAUCUCGCCGGGGCCUCCA	3	2	ncRNA		tRNA
1341	UUAAAUGCCGGAAGCGAUCCCACUGC	3	2	AT-chX		AT-chX-B
1342	UGGACCUUUCCCGAUGAGCCAGU	3	no match			
1343	AAAAACCUGAAGAUGAUCCUGAGUCG	3	1	transposon	LINE	Helena
1344	GCACACAGAGGAGGGCAGAUGAGAGCA	3	no match			
1345	UUCUGUUGAUGUGGGGGCCCCAAGU	3	0	transposon	LINE	G2
1346	UGUUUCACCUUAGACGGCUCGGGC	3	no match			
1347	ACUGCAUUCGCGUCGAGGCCCAGG	3	0	transposon	LINE	R1-element
1348	UGUUUCAUCGUUAGACGACUCUGGC	3	2	AT-chX		AT-chX-B
1349	CUCUAGAGAGAGGUGGCUCCAGGCUC	3	0	transposon	DNA	1360
1350	UAAUCUUGAGAAGGAAGACCCCGGA	3	0	transposon	LTR	gypsy
1351	UAGAUCUUCACCGUUGACUUGGCCCU	3	1	Su(Ste)		Su(Ste)
1352	CGAAAAUCCUCAACCGUGAGGGAUCCCUCAA	3	no match			
1353	CGUUUCAUCGUCAGACGGCUCGGGC	3	2	AT-chX		AT-chX-B
1354	CCUGCUGACGGGGCCCGAAAACACGAGA	3	0	transposon	LTR	opus
1355	UCGAUCUUCACCGUUGACUAGCCCC	3	1	Su(Ste)		Su(Ste)
1356	GCCCGAUGAGAAGAAUGGCCCGUAAC	3	1	transposon	LINE	RR49783
1357	AUCUCAUCGUCGUAGAACAAGCCCGGA	3	2	Su(Ste)		Su(Ste)
1358	CUCAACUGUGGGGGGAUCCUCAA	3	no match			
1359	UACAAUCUCCGAGCAAGACCCCCGAUC	3	no match			
1360	UGUUUCAUCGUUAGACGGCUCCGGGC	3	no match			
1361	UCAACUGUGAGGGGAUCC	3	2	no annotation		
1362	UAGAUCUUCACCGUUGACUUCGGCCCA	3	no match			
1363	UCUGAGGAACCAAGUGAUCCAGCUG	3	2	Su(Ste)		Su(Ste)
1364	CCCUGAUGAAGAAGAGAUGCCCAGUG	3	0	transposon	LTR	ninja-Dsim-like
1365	UCUUCACCGUUGACUAGCCCCACAC	3	0	Su(Ste)		Su(Ste)
1366	UCAACUGUGUGGGGAUCCUCAA	3	no match			
1367	UAUCUUCACCGUUGACUUGGCCCA	3	1	Su(Ste)		Su(Ste)
1368	AAAACCUGAAGAUGAUCCUGAUUCGCA	3	1	transposon	LINE	Helena
1369	AAGACGUGCAACUCUACGGCCCGG	3	0	transposon	LTR	ninja-Dsim-like
1370	GAUCCUCAACUGUGAGGAGAUCCUCAA	3	no match			
1371	CCCGUACUCUGUGGUGACCAACACAC	3	0	transposon	LINE	baggins
1372	CAAGAUAUGACCAAAGUAAAGACAC	3	0	transposon	DNA	1360
1373	UAACGACCAGACAUACGGCAUUAUCUG	3	1	transposon	LINE	R1-element
1374	CAGAUACCCGUUACUCAGCUAGUG	3	0	transposon	LINE	INE-1
1375	CAGCACACAGAGGAGAGCAGAUGAGAG	3	no match			
1376	UGUACUGUUUCUAGGUGUACUGACCCA	3	1	transposon	LINE	baggins
1377	UUUGUUGAUCAGUUUACACAUUAUUG	3	0	transposon	LINE	Cr1a
1378	UAAGCCCCGUACUCUGUGGUGACCAU	3	1	transposon	LINE	baggins
1379	UCCUUGCAGGUUGCUGAUCGAUGACCA	3	0	transposon	LINE	Rt1b
1380	UAAAUGCCAGAAUUUUAUUGGGGUUCA	3	0	transposon	LINE	baggins
1381	CAGCACACAGAGGAGAGCUAGAUGAGAGCA	3	no match			
1382	UCAAGGCCUUCUGGUGCAACCGAA	3	0	Su(Ste)		Su(Ste)
1383	AUCUCAUCGUCGUAGAACAAGCCCGACUGUAACUCGGGGUCAAU	3	no match			
1384	AUCCUAACUUGAGGGGUCCUCAA	3	no match			
1385	UGCAUAUCAUUCACGCCACGCUGCGACC	3	0	no annotation		
1386	UCCUCAUCGUCGUAGAACAAGCCCCGA	3	no match			
1387	AUCCUCAACUGUGAGGGGAUCCUCCAA	3	no match			
1388	CUUCUCAUCGUCGUAGAACAAGCCCGA	3	1	Su(Ste)		Su(Ste)
1389	UAAGCCCCGUACUCUGUGGUGACCAA	3	0	transposon	LINE	baggins
1390	UAGAUCUUCACCGUUGACUUGGC	3	0	Su(Ste)		Su(Ste)
1391	UACACAAACAUGUUCCUAGACAGGCC	3	0	transposon	LTR	opus
1392	CAACACACAGAGGAGAGCAGAUGAGAGCA	3	no match			
1393	UAAUGCCAGUAAUUUUAGUGGGGU	3	no match			
1394	UUAUACCCAGGUGAUUACCCUAU	3	no match			
1395	AAACCGAUAUGGUUGGACCACAAUCUU	3	1	ncRNA		7SLRNA-RA
1396	AUCGUCUCGGGAGUGAAAUACUCAUCGUCGUAGGAACAAGCCCGA	3	no match			
1397	UCAAGCCCGGGUAUUCUGAAGCCAAGA	3	0	repeat		trf
1398	AUAAUACCUCGAGUACGUCUGCUC	3	0	transposon	LTR	Rt1c
1399	UUAUGACUCAUAGACUCGUCUGGGAC	3	0	transposon	LTR	flea
1400	UCUCAUCGUCGUAGGACUAGCCCGA	3	1	Su(Ste)		Su(Ste)
1401	UAGAUCUUCACCGUUGACUUGGUCCC	3	2	Su(Ste)		Su(Ste)
1402	UGUUUCAUCGUUAGACGCCUCGGG	3	1	AT-chX		AT-chX-B
1403	ACCUCAACUGUGAGGGGAUCCUCAA	3	no match			
1404	GCUGACGGGGCCCGAAAACACGAGA	3	0	transposon	LTR	opus
1405	UCUAUCGAAGAAUCGCGGAUGUGACC	3	0	no annotation		
1406	UCUCAUCGACGUAGAACAAGCCCG	3	1	Su(Ste)		Su(Ste)
1407	CUGAGACACGCAACAGGGGA	3	no match			
1408	CCCCGCGGAAAGACGGAAAUACCCCAUG	3	no match			
1409	AAAGAACCCGAUUCCUCUUCCAUGAG	3	0	transposon	LINE	Doc4

1410	UGACUGUUCCCGUUGAUUCCCGUUU	3	0	no annotation		
1411	UCUCAUCGUCGUAGAAGAAGCCCGAGG	3	1	Su(Ste)		Su(Ste)
1412	GUAUGACCCCUGGUACUGAAAUGC	3	0	transposon	DNA	Bari1
1413	UCUCAUCGUCGGUAGAACAAGCCCGA	3	no match			
1414	UCUCAUCGUCGUAGAACAAGCCCGAGC	3	1	Su(Ste)		Su(Ste)
1415	AUCUCAUCGUCGUAGAACAAGCCCGAG	3	1	Su(Ste)		Su(Ste)
1416	UCACACAGAGGAGAGCAGAUGAGAGCA	3	no match			
1417	UCUCAUCGUCGUAGAACAAGCCCGUG	3	1	Su(Ste)		Su(Ste)
1418	GAAGAUGUCCGGACAGCUGGGCCCGA	3	0	Su(Ste)		Su(Ste)
1419	UGUUUUCAUCGUUAGACGGCUCGGGGC	3	no match			
1420	UCUCGUCAUCGUAGAACAAGCCCGA	3	1	Su(Ste)		Su(Ste)
1421	AUCGUCUCGGGGAUGAAAUGUUUCAUCGUUAGACGGCUCGGG	3	no match			
1422	UCCAACUGAAAGAGUACGAAUGAUC	3	0	transposon	LINE	R1-element
1423	UUCUCAUCGUCGUAGAACAAUCCCGA	2	1	Su(Ste)		Su(Ste)
1424	UCUCCAACUGAAGGAGUACGAAUGAUC	2	0	transposon	LINE	R1-element
1425	GCCCGAAAACACGAGAGCCGAGCGC	2	0	transposon	LTR	opus
1426	UAGGAUCUUCACCGUUGACUUGCCCCCG	2	no match			
1427	AGGCUGUGAGGGGAUCCUCAA	2	no match			
1428	UAACAAGAUGCAUACGAUUUUUUCG	2	0	transposon	DNA	1360
1429	AUCUCAUCGUCGUAGAACGAGCCCGA	2	2	Su(Ste)		Su(Ste)
1430	AUCGCAUCGUCGUAGAACAAGCCCG	2	1	Su(Ste)		Su(Ste)
1431	CGAAAUCUCAUCGUCGUAGAACUAAGCCCGA	2	no match			
1432	UAGCUGGCCGAGUAUUUGAUGUU	2	0	Su(Ste)		Su(Ste)
1433	UAGUUGGGCCGCUGACGAAGUUGA	2	0	transposon	LINE	Doc
1434	CAAACUGUGAGGGGAGUCCUCAA	2	no match			
1435	CGAAAUCCUCAUCGUCCGUUAGAACCAAGCCCCGA	2	no match			
1436	GCACACAGAGGAGAGCAGAUGAGA	2	no match			
1437	AUAGAUCUUCACCGUUGACUUGGCC	2	1	Su(Ste)		Su(Ste)
1438	AAUUGAUCCAGCUGCUGUUGUUG	2	0	Su(Ste)		Su(Ste)
1439	UGUUUCAUCGUUAGACGGCUCGGGCCUGUAACUCGGG	2	no match			
1440	CGUUGACCAAUUUCCGAACAUAGCC	2	2	transposon	LTR	DM88
1441	GGUGUCUCCCGGAGAGCAAUGACU	2	0	ncRNA		pseudogene
1442	CUCAACUGUGGGGGGGAUCCUCAA	2	no match			
1443	GAUCUUCACCGUUGACUAGCCCCACA	2	0	Su(Ste)		Su(Ste)
1444	GCCCGUACUCUGUGGUGACCAA	2	1	transposon	LINE	baggins
1445	AAAAAACCUGAAGAUGAUCCUGAGUCGC	2	2	transposon	LINE	Helena
1446	UGUGAGGAUCCUCAA	2	1	coding gene		CG5146-RA|CG5146
1447	UAGUAACUAUGACGAUUUGAUUC	2	0	transposon	LTR	flea
1448	UUUCUCAUCGUCGUAGAACUAGCCCGA	2	0	Su(Ste)		Su(Ste)
1449	GAGGAAAUCCCGGAUCGACAUGGC	2	0	transposon	LINE	X
1450	AUCUCAUCGUCGUAGAAUAAGCCCGA	2	2	Su(Ste)		Su(Ste)
1451	AUCUCAUCGUCGUAGAACAAGCCCCGACUGUAACUCGGUCAAU	2	no match			
1452	GACUAGCUGGCCGAGGAUCCGAUUUUC	2	0	Su(Ste)		Su(Ste)
1453	UUUGUAGGUGUACUGACCCAGUCGGA	2	0	transposon	LINE	baggins
1454	CAUCUUCUUCUCGACGUUGACCAGCC	2	0	transposon	LTR	ninja-Dsim-like
1455	CUGAUACCGAGAAUGACAAUGGCUGA	2	0	transposon	LTR	gypsy3
1456	AUCCUUAACUGUGAGGGAUCCUCAA	2	no match			
1457	GUUCAACUGGACGGGCCUAGAGUUA	2	1	coding gene		CG40635-RA|CG40635
1458	UCCGAACGUACCACGGUGCCCCCAG	2	no match			
1459	UCUUCAUCGUCGUAGAACAAGCCCGA	2	1	no annotation		
1460	UGCUCUCGAAUGUAUGCCCGAUC	2	1	transposon	LINE	R1-element
1461	AUCAGAUACCCGUUACUCAGCUAG	2	0	transposon	LINE	INE-1
1462	UCUCAUCGUCGUCGAACAAGCCCGA	2	1	Su(Ste)		Su(Ste)
1463	AUCCUCAACUGUGAGGGUCCUCAA	2	no match			
1464	CAAGAUAUGACCAAAGCAAAGCAACG	2	0	Su(Ste)		Su(Ste)
1465	GAUAUCCUGAGCAAGUGAUCUGACC	2	0	transposon	DNA	S
1466	GAUCCUCAACUGUGAGGGAUUCUCAA	2	no match			
1467	UUAGUUGACGUAGAGACUGAUACAU	2	0	transposon	LINE	Circe
1468	UCCGGGGGAAGUAUGGUUGCAAAG	2	0	ncRNA		rRNA
1469	UGUAUGUCCCAGUUGCAUUGCAUA	2	0	transposon	DNA	RR51476
1470	UCUCAUCUUUGUAGAACAAGCCCGA	2	2	Su(Ste)		Su(Ste)
1471	CUGUGAACGAGUGAACUGGCAACAA	2	0	Su(Ste)		Su(Ste)
1472	CAGCACACACGAGGAGAGCAGAUGACGACA	2	no match			
1473	UCCUCAACUGUGAGGGAUCCCCAA	2	no match			
1474	AUCUCAUCGUCGUAGAACAAGCCCCGACUGUAACUCGGGGUCAAU	2	no match			
1475	AGAAAAAAACCUGAAGAUGAUCCUGAGUCGC	2	no match			
1476	GCGAAAAUUGGCCCAAACUGAU	2	0	repeat		SAR_DM
1477	UGUUUCAUCGUUAGGGCUCGGGC	2	no match			
1478	AGAUCAACUGUGAGGGGAUCCUCAA	2	no match			
1479	AAAGGAAUUGACGGAAGGGCACCACC	2	0	ncRNA		rRNA
1480	UAGAUCUUCACCGUUGACUUGGCCAA	2	1	Su(Ste)		Su(Ste)
1481	UCUCAUCGUCGUAGAACUAGCCCAA	2	1	Su(Ste)		Su(Ste)
1482	CUUGAGACCGCCCCUGGACGACCCACG	2	0	Su(Ste)		Su(Ste)
1483	UGUUUCAUAGUUAGACGGCUCGGCC	2	2	AT-chX		AT-chX-B
1484	GUGACGAAGCCUAGA	2	1	no annotation		
1485	UAGAUCUUCACCGUUGACUAGCCUCA	2	1	Su(Ste)		Su(Ste)
1486	AUCUCAUCGUCGUAGAACAAAGCCCCGA	2	no match			
1487	GAUUCACCGUUGACUUGGCCCACACA	2	2	Su(Ste)		Su(Ste)
1488	CCAGGCCCAUCUGGUUGAACGUGUC	2	0	Su(Ste)		Su(Ste)
1489	CAGCACACAGAGAGAGCAGAUGAGAGCA	2	no match			
1490	CUGACCCAGUCGGAUCUAUUGUAC	2	0	transposon	LINE	baggins
1491	UCUCAUCGUCGUAGAACAAGCCCGGAG	2	1	Su(Ste)		Su(Ste)
1492	UGGACCGAACACUUUGAACCAAGU	2	0	Su(Ste)		Su(Ste)

1493	CGACACAGAGGAGAGCAGAUGAGAGCA	2	no match			
1494	UCGCCCAGAGCGGUUUCCCUGUGUUUA	2	1	no annotation		
1495	UAAUGUCCCUUAAAUCAGCACUCGUA	2	0	no annotation		
1496	UUAAAUGCCGGAAGCGAUCCCACUGCU	2	2	coding gene		CG41580-RA|CG41580
1497	UUCGUCAUCGUCGUAGAACAAGCCCGA	2	2	Su(Ste)		Su(Ste)
1498	GACUGCAGAUCGAGAGGUCCCCGG	2	0	ncRNA		tRNA
1499	UAAUGGUAUCCGAGUCGUACUCUAG	2	0	transposon	LTR	springer
1500	GGACAGACGUUUUUCUUCCUCCAGUG	2	0	transposon	LINE	baggins
1501	CUUCGGGAUAAGGAUUGGCUCUGA	2	0	ncRNA		rRNA
1502	CGAAACUCAACUGUGAGGGGAUCCCUCAA	2	no match			
1503	UCUGACGAGAACUCGAGCCGCGUAA	2	1	no annotation		
1504	UUCUCAUCGUCGUAGAACAACGCCCGA	2	no match			
1505	UAAACAAGAUGCAUACGAUGUGGCU	2	2	transposon	DNA	1360
1506	GUCCGGGAGAAAACAAUGAUGGCCA	2	0	coding gene		CG12717-RA|CG12717
1507	AACGAACCAGCCCGAGGAAAUCAAUCUC	2	no match			
1508	GACUAGCUGGCCGAGUAUUUGAUGU	2	0	Su(Ste)		Su(Ste)
1509	UCCUCAACUGUGAGCGAUCCUAA	2	no match			
1510	AAAUGCCAGAACUUUAUUGGGGU	2	0	transposon	LINE	baggins
1511	GUGGAAGAAAGUACAAUCCUCAA	2	no match			
1512	AUUGCGCCCCUUGGUAUUCCAG	2	no match			
1513	AUUCUCAUCGUCGUAGAACAAGCCCCGA	2	no match			
1514	CAGCACACAGAGGAGGGCAGAUGAGAGCA	2	no match			
1515	CCCUGACCUAGAUGACUGCAGAACC	2	0	transposon	LINE	TART-A
1516	AAUUUCGUAAACCAAAGGCCGUGAGUU	2	no match			
1517	CUCGAAGAUGACGGAGAAGUGCUU	2	0	coding gene		CG11611-RA|Tim13
1518	UACGAUCUUCACCGUUGACUAGCCCCCA	2	no match			
1519	UAUCGUCUUCAUUGCGACGGCGAC	2	0	AT-chX		AT-chX-A
1520	UCAAGGCCUUCUGGUGCAACCGAGA	2	0	Su(Ste)		Su(Ste)
1521	UCUCAUCGUCCUAGAACAAGCCCG	2	1	Su(Ste)		Su(Ste)
1522	GGAUCCUCAACUGUGAGUGGAUCCUCAA	2	no match			
1523	ACUCCCCAUCCUGCACUCUCC	2	no match			
1524	CACCACAGAGGAGAGCAGAUGAGAGCA	2	no match			
1525	UCUCAUCGUCGUAGAAUAGCCCGA	2	no match			
1526	UAGAUUUUCACCGUUGACUAGCCCCA	2	1	Su(Ste)		Su(Ste)
1527	UCCGGGAGAAAACAAUGAUGGCCAAG	2	0	coding gene		CG12717-RA|CG12717
1528	UACAAUGAUCUGAUCGGGUUUUCU	2	2	transposon	LINE	Cr1a
1529	UUUAUACCUGAGCCAGUUUGUUUUUG	2	2	transposon	Low_complexity	F-element
1530	UCUCAUCGUCGUAGAACAAGCCCAG	2	2	Su(Ste)		Su(Ste)
1531	UGCUGAACGUAGUCUCUCGAGAGCGA	2	1	transposon	LTR	invader1
1532	UCUCAUCGUCGUAAGAACAAGCCCGA	2	no match			
1533	UGGAAUAGUUUUGACCGUUGACCA	2	0	transposon	LTR	opus
1534	CACACAGAGGAGAGGAGAUGAGAGCA	2	no match			
1535	UCCAGUGACUUUAGUAUGCCCCGUAC	2	0	transposon	LINE	baggins
1536	UAAAUUAUAAGGAUGAUGAUACCGAU	2	1	no annotation		
1537	UGUCUUCAUCACUACGGGCCCAG	2	0	transposon	Low_complexity	F-element
1538	GGAUCCUUAACUGUGAGGGGAUCCUCAA	2	no match			
1539	AUUCAUCGUCGUAGAACAAGCCCG	2	1	no annotation		
1540	AUCUCAUCGUCGUAGAACAAUCCCGA	2	2	Su(Ste)		Su(Ste)
1541	CUCAUCGUCGUAGAAGAAGCCCGAG	2	1	Su(Ste)		Su(Ste)
1542	AUAGAUCUUCACCGUUGACUAGCCC	2	1	Su(Ste)		Su(Ste)
1543	CUGAGACACGCAACAGGGGAUAGGCAAGGCA	2	no match			
1544	CCAGGCCCAUCUGGUUGAACGUGU	2	0	Su(Ste)		Su(Ste)
1545	AUUUCUCAUCGUCGUAGAACAAGCCCGA	2	1	Su(Ste)		Su(Ste)
1546	UCUCAUCAUUCGUAGAACAAGCCCGA	2	no match			
1547	UCUCAUCGUCGUAGAACAACCCGA	2	no match			
1548	GAUCUUCACCGUUGACUAGCCCC	2	0	Su(Ste)		Su(Ste)
1549	CAGACACAGAGGAGAGCAGUUGAGAGCA	2	no match			
1550	UACUCGAAGACACCACGGUGAAAGUG	2	0	transposon	LTR	ninja-Dsim-like
1551	AUCCUCAACUGUGAGGGAUCCUCA	2	no match			
1552	UAGAUCUUCACCGGUUGACUAGCCCCA	2	no match			
1553	AGGUGUUUCAUCGUUAGACGGCUCGGGC	2	no match			
1554	UAAAUGCCAGAUUUAUGGGGUUC	2	no match			
1555	GAUUCCGCAGUGCGUUGAUAUGGCCAG	2	0	Su(Ste)		Su(Ste)
1556	UAGGAUCUUCACCGUUGACUUGGCCCGA	2	no match			
1557	UAGAUCUUCACCGUUGACUACGCCCC	2	no match			
1558	UAGAUCUUCACCGUUGACUAGCACCA	2	1	Su(Ste)		Su(Ste)
1559	UAAAUGCCAGAAUUUUAUUGGGGUUA	2	1	transposon	LINE	baggins
1560	AAUCUCAUCGUCGUAGAAGAAGCCCGA	2	no match			
1561	GCACACAGAGGAGAGCAGAUGAGAA	2	no match			
1562	UCUCCAUCGUCGUAGAACAAGCCCGA	2	2	no annotation		
1563	AUCCUCAACUGUGAGGUGAUCCUCAA	2	no match			
1564	GCUGGCCGAGGAUCCGAUUUUCUUUG	2	0	Su(Ste)		Su(Ste)
1565	GCAAUAAGACCCCAUUAGCAGCCC	2	1	no annotation		
1566	UCUCAUCGUCGUAGAACUAGCCCGACUGUAACUCGGGGUCAAU	2	no match			
1567	CAGCACACAGAGGAGGAGCAGAUGAGAGCA	2	no match			
1568	AUCUCAUCGUUGUAGAACAAGCCCGA	2	2	Su(Ste)		Su(Ste)
1569	UCGUCUCGGUGAAAGGAUCCUCAA	2	no match			
1570	GACAGAAUAGAGAGGGAUCCUCAA	2	no match			
1571	AUCGUCUCGGGAGUGAAAUACUCAUCGUCGUAGAACAAGCCCGA	2	no match			
1572	CUGUAACUCGGGGUCCGAAGG	2	no match			
1573	UAGCUGGCCGAGUAUUUGAUAUU	2	0	Su(Ste)		Su(Ste)
1574	CUGUUUCAUCGUUAGACGGCUCCGGGCG	2	no match			
1575	CAAGAUAUGACCAAAGUAAAGACUA	2	2	transposon	DNA	1360

1576	UCGAAUCAACAAAUUGCGCCGAUAU	2	0	coding gene		CG41545-RA|CG41545
1577	UCUCAUCGUCGUAGAACUAGCCCGAA	2	1	Su(Ste)		Su(Ste)
1578	UCCAAAGCGGGACCUCUUAGGCAAG	2	0	transposon	LINE	G3
1579	GGAUUAGAUACCCUGGUAGUCCACGC	2	no match			
1580	AAUGUUUCAUCGUUAGACGGCUCGGGC	2	2	AT-chX		AT-chX-B
1581	CUUGGACGGACAGGUUACACGAUC	2	0	transposon	LTR	297
1582	AAUUCCGGAGAGGGAGCCUGAGAAACGG	2	1	ncRNA		rRNA
1583	UGUACUGACCCAGUCGGAUCUAUAAA	2	no match			
1584	GGCAUUCCGAAUUAAUGUUGGCCAA	2	0	transposon	LINE	Rt1b
1585	UCGUCAUCGUCGUAGAACAAGCCCGGCUG	2	no match			
1586	UACAGUAUUGGUCCAAGAACGCUGCC	2	0	transposon	LINE	G
1587	UUACAAUCUCCGAGCACGACCCCGA	2	0	transposon	LINE	Circe
1588	UCUCAUCGUCGAGAAGUAGCCCGA	2	no match			
1589	UCUCAUCGUCGUAGAACAAGCCCCGUA	2	no match			
1590	UACACAAACGUCCCUGACCCAACCUGA	2	0	coding gene		CG15669-RC|MESK2
1591	UAAAGAAUCCUGACAAGAUCCCGCG	2	1	transposon	LTR	invader1
1592	UGUACUGGAUGCCACUCCUGGUGU	2	2	transposon	LINE	G2
1593	UGUUUCAUCGUUAGACGGCUCGAA	2	2	AT-chX		AT-chX-B
1594	UCUCAUCGUCGUAGAACAAGCCCGAGCUGUAACGUCGGGUCAAU	2	no match			
1595	ACUGCAGAUCGAGAGGUCCCCGGU	2	0	ncRNA		tRNA
1596	CCUCAACGUGAGGGGAUCCUCAA	2	no match			
1597	UUUCUGACGAGAACUCGAGCCGCGUA	2	0	no annotation		
1598	UCUCAUCAUCGUAGAACGAGCCCCGA	2	no match			
1599	UCCCCGCCGAGGUACGAUUCCUGCACC	2	1	transposon	LINE	R1-element
1600	UCUCAUCGUCGUAGAACUACGCCCGA	2	no match			
1601	UUUCUGACGAGAACUCGAGCCGCG	2	0	no annotation		
1602	UCUCAUCGUCGUAGAACAAGCCCCGU	2	2	Su(Ste)		Su(Ste)
1603	UCCGCAUGUGAAUGUUGGCCCCAG	2	0	no annotation		
1604	CGACAGAAUAGAGAGGGAUCCUCAA	2	no match			
1605	CGAGCACACAGAGGAGAGCAGAUGAGAGCA	2	no match			
1606	UGGAUUUUUUUUCCGUAUGAGCCAGUG	2	no match			
1607	AACUCUAGGCUGAGCGGAGAACCAGAG	2	no match			
1608	AGGGGCGAAGGCGGCUGUCUGGU	2	no match			
1609	UGCAGACGUGCCUACGGACCGACG	2	0	transposon	LINE	G
1610	UCUCAUCGUCGUAGAACAAGCACGAA	2	1	Su(Ste)		Su(Ste)
1611	GAUAUGACCAGCAUGGACCUAACACU	2	0	Su(Ste)		Su(Ste)
1612	UCUCAUCGUCGUAGAAGAAGCCCGU	2	2	Su(Ste)		Su(Ste)
1613	UGAUACCAAGAAUGACAAUGGUUGA	2	1	transposon	LTR	gypsy3
1614	UCUAAUCGUCUUAGAACAAGCCCGA	2	2	Su(Ste)		Su(Ste)
1615	AUGUGAGGGGAUCCUCAA	2	2	no annotation		
1616	AUCUCAUCGUCGUAGAACAAUCCCCGA	2	no match			
1617	AUCUCAUCGUCGUAGAACAAGCCUGA	2	2	Su(Ste)		Su(Ste)
1618	UCUCAUCGUCAUAGAACAAGCCCG	2	1	Su(Ste)		Su(Ste)
1619	AUCUCAUCGUCGUAGAACAAGCCCGCUGUA	2	no match			
1620	GCUCCGAACUCUCUUGGCCCAAGAC	2	0	coding gene		CG11306-RA|CG11306
1621	AACUAGUGAGGGGAUCCUCAA	2	no match			
1622	UGUUUUCAUCGUUAGACGGCUCGGGU	2	no match			
1623	UGGGAACACCGCGUGUUGUUGGCCU	2	0	ncRNA		5SrRNA
1624	UCUCAUCGUCGUAGAAGUAAGCCCGA	2	no match			
1625	UGUUACAAUGAUCUGAUCGGGUCAAU	2	no match			
1626	CCUCUACUGACUUCAGUAAGCCCCG	2	0	transposon	LINE	baggins
1627	UAAACAAGAUGCAUACGAUGUGGCUC	2	2	transposon	DNA	1360
1628	GGAUCAACUGUGAGGGAUCCUCAA	2	no match			
1629	UGUUACAAUGAUCUGAUCGGGUU	2	0	transposon	LINE	Cr1a
1630	CAAACUGUGAGGGGAUCCUCAA	2	no match			
1631	UCGCAUCGUCGUAGAACAAGCCCGCUGUAACUCGGGGUCAAU	2	no match			
1632	UCCUCAACUGUGAGGGGGAUCCUCAA	2	no match			
1633	UUACAAUCUCCGAGCAAGACCCCGA	2	0	transposon	LINE	Circe
1634	UCCGUAGGGAUGAUUUGUACCGUAGA	2	0	transposon	LINE	F
1635	UAGAUCUUCACCGUUGACUAGCCCCAG	2	1	Su(Ste)		Su(Ste)
1636	UCCGAUCUCCUGAGUCGGACUCCGAGU	2	1	transposon	LINE	R1
1637	GCAUCCUAGCUGGCGGGACCGACA	2	0	coding gene		CG1638-RB|CG1638
1638	AAUCCUCAACUGGAG	2	1	coding gene		CG14512-RA|CG14512
1639	UCUCAUCGUCGUAGAACAACGCCC	2	no match			
1640	AGAUUCUUGAGAAUUUGGACGCCGUU	2	0	transposon	LINE	Cr1a
1641	UCUCAUCGUCGUAGAACAAGCCCGGGG	2	1	Su(Ste)		Su(Ste)
1642	UGUAUCAUCGUUAGACGGCUCGGGC	2	1	AT-chX		AT-chX-B
1643	UAGCAUCGUCGUAGAACAACGCCCGA	2	no match			
1644	UUACAAUGAUCUGAUCGGGUUUUUCU	2	0	transposon	LINE	Cr1a
1645	AGGAGCCUCAACUGUGAGGGGAUCCUCAA	2	no match			
1646	UUCGUCAUCGUCGUAGAACAAGCCCG	2	2	Su(Ste)		Su(Ste)
1647	CAAUUGAUCCAGCUGCUGUUGUUG	2	0	Su(Ste)		Su(Ste)
1648	UAGAUCUUCACCGUUACUUGGCCC	2	no match			
1649	UGUUUCAUCGUUAGACGCUCGGG	2	no match			
1650	AAUCUCAACUGUGAGGGAUCCUCAA	2	no match			
1651	UGGACUUUUUUCCGUAUGAGCCAGUG	2	no match			
1652	UAGGACCUGGGUUUGUCAACGGACAACUG	2	no match			
1653	CAGCACACAGAGGAGAGCAGAUGAGAGCG	2	no match			
1654	UAACGACCAGACAUACGCCAUAAUCUC	2	1	transposon	LINE	R1
1655	UAAAAUGCCAGAACUUUAUUGGGGU	2	2	transposon	LINE	baggins
1656	UGAUCUUCACCGUUGACUAGCC	2	1	Su(Ste)		Su(Ste)
1657	CCCGUACUCUGUGGUGACCAAC	2	0	transposon	LINE	baggins
1658	CACACACAGAGGAGAGCAGAUAGAGCA	2	no match			

1659	UCCUAGUAUUCUUCCGGAUGAUCCACC	2	0	transposon	DNA	S
1660	CCGAACGUACCACGGUGCCCAGUGA	2	no match			
1661	CUGUGAGGGGGAUCCUCAA	2	2	no annotation		
1662	GACUAGCUAGCCGAGGAUCCGAUU	2	1	Su(Ste)		Su(Ste)
1663	UUCACUCGUUCACAGAAAUAUGACU	2	0	Su(Ste)		Su(Ste)
1664	UCUCAUCGUCGUAGAACAAGCCCUC	2	2	Su(Ste)		Su(Ste)
1665	GCGUUGACCAAUUUCCGAACAUACGA	2	1	transposon	LTR	Dm88
1666	GCCCGAUGAGAAGAAUGGCCCGUAA	2	0	transposon	LINE	RR49783
1667	AGAAAAAAAAAAAAAAGGAUCCUCAA	2	no match			
1668	UAACAAGAUGCAUACGAUGUGGCUU	2	1	transposon	DNA	1360
1669	UCUCAUCGUCGUGGAAGAAGCCCGA	2	2	Su(Ste)		Su(Ste)
1670	UGUUUCAUCGUUAGACGGCUCGGGCCUGG	2	no match			
1671	UCUCAUCAUCGUAGAACAACGCCCGA	2	no match			
1672	UGCUCUCGAAUGUAUGCCCGAUAUAC	2	0	transposon	LINE	R1-element
1673	UAAACAAGAUGCAUACGAUGUGGCUCC	2	2	transposon	DNA	1360
1674	ACCUCAACUGUGAGGGAUCCUCAA	2	no match			
1675	UCCAGGCCCAUCUGGUUGAACGUGUC	2	0	Su(Ste)		Su(Ste)
1676	UCGAGAAGCGUUUGAUCGAGUGUGG	2	0	transposon	LINE	BS3
1677	UCUCAUCGUCUAGAACUAGCCCGA	2	no match			
1678	CUCAUCGUCGUAGAACAAGCCCGAG	2	0	Su(Ste)		Su(Ste)
1679	AUCUCGUCGUCGUAGAACAAGCCCCGA	2	no match			
1680	AUCGUCUCGGGGAUGAAAUGUUUUCAUCGUUAGACGGCUCGGGC	2	no match			
1681	CUCAACUGAGGGGAUCCUCAA	2	no match			
1682	GUAACUUCGGGAUAAGGAUUGGCUCUG	2	0	ncRNA		rRNA
1683	AUCGUCUCGGGACUGAAAUCUCAUCGUCGUAGAACAAGCCCGA	2	no match			
1684	UGUACACACCGCCCGUCGCUACUACCG	2	0	ncRNA		rRNA
1685	UUUUCAUCGUCGUAGAACAAGCCCGA	2	0	no annotation		
1686	GCAGCUCUUCCGUGGCGACACCGUGA	2	0	coding gene		CG2331-RA|TER94
1687	UCACGCGGGAGACCGGGGUUCAAUUU	2	1	ncRNA		tRNA
1688	UUGUUUCAUCGUUAGACGGCUCGGGC	2	0	no annotation		
1689	GUGUUUCAUCGUUAGACGGCUCGGGGC	2	no match			
1690	UAGUUUUCAUCGUUAGACGGCUCGGGC	2	no match			
1691	AUCCAACUGUGAGGGGAUCCUCAA	2	no match			
1692	ACUGCAUUUGCGUCGAGGCCCAGGA	2	0	transposon	LINE	R1-element
1693	CAAGGAUAUGACCAAAGCAAAGACGA	2	no match			
1694	GAUGUCCGGACAGCUGGGCCCGAA	2	0	Su(Ste)		Su(Ste)
1695	CUCAACUGUGAGGGAAUCCUCAA	2	no match			
1696	UCCAAGACCCCAAGGACAUUGUCGAU	2	0	coding gene		CG4536-RA|iav
1697	CGUUGAGCAAUUUCCGAACAUACGCC	2	1	transposon	LTR	DM88
1698	UACAGAGUUUUGCUCGGCUCGAAGU	2	0	transposon	LTR	rover
1699	UAAGAUUGGACCCAGUACACUGCCC	2	1	transposon	LINE	Doc
1700	UCUCAUCGUCAUAGAACAAGCCCCC	2	no match			
1701	UCUCAUCGUCGUACGAACAAGCCCGA	2	no match			
1702	UAGGUUGUACCCCUACAGCAUGAA	2	0	transposon	LTR	opus
1703	UUCAUCGUCGUAGAACAAGCCCGAC	2	1	no annotation		
1704	ACUUCAGAGUCGACGUACCCGUU	2	1	transposon	LTR	G4
1705	UGUUUCAUCGUAGACGGCUCGGGC	2	no match			
1706	GGCCUCAACUGUGAGGGAUCCUCAA	2	no match			
1707	UACAAUACCCAGUUACUCGUGCCAU	2	0	no annotation		
1708	AUCUCGUCGUCGUAGAACAAAGCCCCGA	2	no match			
1709	CAGACAGAGGAGAGCAGAUGAGAGA	2	no match			
1710	UCUCAUCUUCGUAGAACAACGCCCGA	2	no match			
1711	CAACCAGAGAUCUAGAGGGCCCAAG	2	0	no annotation		
1712	UCGGACACCGAACCGGGAAGGCGAA	2	0	transposon	LTR	Tirant
1713	AAACAGCACACAGAGGAGAGCAGAUGAGAGCA	2	no match			
1714	ACAAGAUAUGACCAAAGCAAAUACAC	2	1	transposon	DNA	1360
1715	GGAUACCUCAACUGUGAGGGGAUCCUCAA	2	no match			
1716	UCCGGAGUGUUGAUCUGAGCCUCAAA	2	0	transposon	LTR	GATE
1717	CCUCAACUGUGAGGUCAA	2	1	no annotation		
1718	UCUCAUCGUCGUACAACAAGCCCGA	2	1	Su(Ste)		Su(Ste)
1719	UGUCUCAUCGUUAUACGGCUCGGGC	2	2	AT-chX		AT-chX-B
1720	UCGUAUCGUCGUAGAACAAGCCCGCG	2	2	Su(Ste)		Su(Ste)
1721	UCUCACUCGACUGCGACACCAUCCC	2	1	transposon	LINE	G2
1722	UCAACUGUGAGGAGGAUCCUCAA	2	no match			
1723	CAGCACACAGAGGUGAGCAGAUGAGAGCA	2	no match			
1724	UGUUUCAUCGUUACACGGCUCGGG	2	1	AT-chX		AT-chX-B
1725	UCAUCUUCUUCUCGACGUUGACCG	2	1	transposon	LTR	ninja-Dsim-like
1726	UUCUCAUCGACGUAGAACAAGCCCGA	2	1	Su(Ste)		Su(Ste)
1727	CUCAACUGUGAGGGGGAGUCCUCAA	2	no match			
1728	UUCUCAUCGUCGUAGAACAAGCCCA	2	1	Su(Ste)		Su(Ste)
1729	UAAAAGAAUCCUGACAAGAUCCCGGG	2	2	transposon	LTR	invader1
1730	AGCCCCCGUACUCUGUGGUGACCAACA	2	1	transposon	LINE	baggins
1731	GAGAGGCGCCGUAGUUACCACCCCACC	2	0	no annotation		
1732	UGGACUCUUUUCCGAUGAGCCAGU	2	1	transposon	LINE	G2
1733	GCUGGCCGAGGAUCCGAUUUUCUUU	2	0	Su(Ste)		Su(Ste)
1734	UCUCAUCGUCGUAGAACAAGCCCCAGAAAGUAAACA	2	no match			
1735	AUCCUCAACUGUGAGGGAUCCCCAA	2	no match			
1736	UUUUUUCAUCGUUAGACGGCUCGGGC	2	1	no annotation		
1737	CUGUGAGGAUCCUCAA	2	1	no annotation		
1738	UCCGAUCUCCUGAGUCGAACUCCGACC	2	1	transposon	LINE	R1-element
1739	UUAAUGCUGGAAGCGAUCCCACUGCC	2	1	AT-chX		AT-chX-B
1740	UUCGAUCUCCUGAGUCGGACUCCGAGC	2	1	transposon	LINE	R1
1741	AUCGUCUCGGGAGUGAAAAUACUCUAUCGUCGUAGAACAAGCCCG	2	no match			

1742	UGCUAGGAGGCCCGUAUUACUGUC	2	1	transposon	LTR	ninja-Dsim-like
1743	UGUUCAUCGUUAGACGGCGUCGGGC	2	no match			
1744	UAAAUGCCAGAAUUUAUUGGGGUUC	2	no match			
1745	ACUUGACAUCGCCCCUGGACGACCCACG	2	0	repeat		trf
1746	UGCGUGACGGGACCCAGAUAGCCUC	2	0	transposon	LINE	Rt1a
1747	CCAUCAGAGACUUUGGUGGCCAACAAA	2	1	coding gene		pseudogene
1748	AUUCUUCUCGACGUUGACCAGCCUCC	2	1	transposon	LTR	ninja-Dsim-like
1749	UAAGGAUUGGACCCAGUACACUGCCGU	2	no match			
1750	CCGGAGAAGACUUUGAAUGGAUCG	2	1	no annotation		
1751	UUCUCAUCGUCGUGAACAAGCCCAA	2	no match			
1752	UGUUUCAUCGUUAGAACGGCUCGGGC	2	no match			
1753	UCUCAUCGUCGUAGAACAACCCGAA	2	no match			
1754	UUUCUCACCGUCGUAGAACAAGCCCGA	2	1	Su(Ste)		Su(Ste)
1755	GGAUCCUCAACUGUGAGGG	2	2	coding gene		CG34138-RA|CG34138
1756	CGAAAUCCUCAUCGUCCGUUAGAACCAAGCCCCGACU	2	no match			
1757	UCUCAUCGUCGUAGUAGAAGCCCG	2	2	Su(Ste)		Su(Ste)
1758	UCUCAUCGUCGUAGAACAACGUCCGA	2	no match			
1759	UCUUCACCGUUGACUAGCCCCACA	2	0	Su(Ste)		Su(Ste)
1760	CAGAUACCCGUUACUCAGCUAGUAUAA	2	0	transposon	LINE	INE-1
1761	UUUUUCAUCGUUAGACGGCUCGGG	2	1	AT-chX		AT-chX-B
1762	UGUUUCAUCUUAGACGGCUCGGG	2	no match			
1763	UGAUGACCGAAAUUCGGAAAAACG	2	0	repeat		ARS406_DM
1764	UCCAGAGCAAGUGAUCUGACCGAGAU	2	0	transposon	LTR	Max
1765	UUCACCGGGUUUUGGCACUUUGCCGC	2	0	transposon	LINE	R1-element
1766	UAGAUCUUCACCGUGGACUAGCCCC	2	1	Su(Ste)		Su(Ste)
1767	CUCAACUGUGAGGGUAUCCUCAA	2	no match			
1768	CAGCACACAGAGGAGAGCCAGAUGAGAGCA	2	no match			
1769	UCCGAUCUCCUGAGUCGGACUCCGAU	2	1	transposon	LINE	R1
1770	AAAACCUGAAGAUGAUCCUGGGUCGC	2	1	transposon	LINE	Helena
1771	GGACUUCCGAGAAAGCCCGCAGAA	2	0	transposon	LTR	flea
1772	UCACGCCAGUCCCAUCAGGUCGCCCA	2	no match			
1773	GUCUCAUCGUCGUAGAACAAGCCCGUCUGUAACCUCGGGUCAAU	2	no match			
1774	CAAGGAUAUGACCAAAGCAAAGACGAC	2	no match			
1775	UUCCCGUGUGUCAUUAAGGUCUUGAACC	2	0	transposon	LINE	baggins
1776	CAUCACACAGAGGAGAGCAGAUGGGAGCA	2	no match			
1777	UGGACGGAGCGAUGUUCGGGCCCA	2	0	repeat		trf
1778	UACGAUCUUCACCGUUGACUAGCCCCA	2	no match			
1779	AACCUAGUUACUUACGGGCUCGACA	2	0	Su(Ste)		Su(Ste)
1780	UCCAGUCUCCCGACUAUCAGAUACCCGU	2	0	transposon	LINE	INE-1
1781	AUUUUCAUCGUCGUAGAACAAGCCCGA	2	1	no annotation		
1782	UUAUUCCCAAUGCUUUGCGACGGCGCA	2	no match			
1783	AUCGUCUCGGAUGAAAUCUCAUCGUCGUAGAACAAGCCCGGA	2	no match			
1784	UCGACUACAUUCAGGACCCGCUCAACC	2	1	coding gene		CG40635-RA|CG40635
1785	GACUAGCUGGCCGAGUAUUUGAAA	2	1	Su(Ste)		Su(Ste)
1786	AGAAAAAAAAAAAAAAGGAUCCUCAA	2	no match			
1787	AUGUUAAGGACGAGAAUCUACCCC	2	0	transposon	LTR	diver2
1788	UAUACCCAGGUGAUUACCACUAUUA	2	0	Su(Ste)		Su(Ste)
1789	UCAACGAAGACCCAGUAUACCGUA	2	no match			
1790	AUCGUCUCGGGGAUGAAAUCUCAUCGUCGUAGAACAAGCCCG	2	no match			
1791	UACAUCUUCACCGUUGACUAGCCCCA	2	1	Su(Ste)		Su(Ste)
1792	UCCUGAUGACUCUUCGACAUGUAC	2	0	no annotation		
1793	UUAAUGCCGGAAGCGAUCCCA	2	0	AT-chX		AT-chX-B
1794	GCACACAGAGGAGAGCAGAUGAGAGUA	2	no match			
1795	AGUUCCUGACUUUAUGCCCCAGUUU	2	0	transposon	LINE	G2
1796	AGAUCUUCACCGUUGACUUGCCCAAA	2	2	Su(Ste)		Su(Ste)
1797	UCUCAUCGACGUAGAACAAGCCCGAA	2	1	Su(Ste)		Su(Ste)
1798	UCUCAUCGUCGCUGAACAAGCCCGA	2	2	Su(Ste)		Su(Ste)
1799	AUACAAUCUCCGAGCAAGACCCCCGAUC	2	no match			
1800	CUCAUCGUCGUAGAAGAAGCCCGA	2	1	Su(Ste)		Su(Ste)
1801	UAAAUGCCAGAAUUUUACUGGGGUUC	2	1	transposon	LINE	baggins
1802	AUCCUCAACUGUGAGGGAUUCAA	2	no match			
1803	AUCGUCUCGGGACUGAAACUCUCAUCCGUUCGUACGAACAACGCC	2	no match			
1804	AUCUCAUCGUCGUAGAAACAAGCCCCGA	2	no match			
1805	GUCUCAACUGUGAGGGGAUCCUCAA	2	no match			
1806	UUACCAGGUCCGAACAUAAGUGUGUA	2	0	ncRNA		rRNA
1807	UCUCAUCGUCGUAGAACAAGGCCCGACUGUAACGUCGGGUCAAU	2	no match			
1808	UCCUUCCCAAGGCGUCUGGAUCCAGU	2	0	transposon	LTR	flea
1809	UCUCAUCGUCGUAGGACAAGCCCUA	2	2	Su(Ste)		Su(Ste)
1810	CCGAUCUCCUGAGUCGGACUCCGAGCU	2	0	transposon	LINE	R1
1811	GGAUCCCAACUGUGAGAUCCUCAA	2	no match			
1812	UUCUCAUCGUCGUAGAACAAGUCCGA	2	1	Su(Ste)		Su(Ste)
1813	UAAUGCCCCGUAGGUUGCCCUCUGU	2	0	transposon	LINE	R1-element
1814	UGUUUCAUCGUUAGACGGCUACGGGC	2	no match			
1815	UUAUAAGGAUGAUGAUACCGAUUAA	2	0	no annotation		
1816	UGAGCAACUAAUGACUAGCGCUAA	2	0	repeat		SAR_DM
1817	UCUUCAUCGUCGUAGAACAAGCCCG	2	1	no annotation		
1818	GAAUCUCAUCGUCGUAGAACAAGCCCGA	2	no match			
1819	UCAAGGUUACCCCAGGAUGCUGU	2	0	repeat		trf
1820	CUGGACCGAACACUUUGAACCAAGU	2	1	Su(Ste)		Su(Ste)
1821	GCACACAGAGGAGAGCAGAUGAGGCA	2	no match			
1822	UCCGUCGGGAUGAUUUGUACCGUAG	2	0	transposon	Low_complexity	F-element
1823	GCACACAGAGGAGAGCAGAUGAGACGCA	2	no match			
1824	UCUCAUCGUCGUAGAACAAGCCCGCUGUAACUCGGGGUCAAU	2	no match			

1825	UAAUUACCGGUAUAGCACACUUUUG	2	0	coding gene		CG41580-RA|CG41580
1826	CAGCACACAGAGGAGAGCAGAUGAAA	2	no match			
1827	AUCAAGGUUACCCCAGGAUGCUGU	2	1	repeat		trf
1828	UGUUUCAUCGUUAGACGGCUCGGGCCUGUAACUCGGUCAAU	2	no match			
1829	UGUUUUCGUUAGACGGCUCGGGC	2	no match			
1830	AAACAUGUUCCUAGACAGGCCCCGA	2	0	transposon	LTR	opus
1831	AACUUACCGUCGUAGAACAAGCCCGA	2	no match			
1832	UUUACGUGGACGAGUUGCUCGAUUU	2	0	transposon	Low_complexity	F-element
1833	UUUCCUGAAAUCCGAGUUAAUAUG	2	0	no annotation		
1834	UGUUUCAUCGUUAGAUGGCUCGGG	2	1	AT-chX		AT-chX-B
1835	UCUCCGGGAGACACCAAUGCUGGCCAA	2	0	coding gene		pseudogene
1836	UGACGAGAACUCGAGCCGCGUAGC	2	0	no annotation		
1837	GUUCAUUAUCAGACCACGCCCAAAGU	2	no match			
1838	AUCGUCAUCGUCGUAGAACAAGCCCGGACUG	2	no match			
1839	UCUGUUUUCAUCGUUAGACGGCUCGG	2	no match			
1840	ACAGCUCGAUGCGAGGCCUAGUACA	2	0	transposon	LINE	X
1841	UCUCACUCGUCGUAGAACAAGCCCGA	2	no match			
1842	UAAAUGCCAGAACUUUAUUGGCGUUC	2	1	transposon	LINE	baggins
1843	AUCGUCUCGGGAGUGAAAUAAACAAGAUGCAUACGAUGUCGGC	2	no match			
1844	UCUCUAUCGUCGUAGAACAAGCCCGA	2	2	no annotation		
1845	UAGAUCUUCACCGUUGACCUGGCCC	2	1	Su(Ste)		Su(Ste)
1846	UCGAUCUUCACCGUUGACUAGCCCCA	2	1	Su(Ste)		Su(Ste)
1847	UGUUUCAUCGUUAGACGGCUCGGCCC	2	2	AT-chX		AT-chX-B
1848	UCUCAUCGUCGUAGAACAAGCCCU	2	1	Su(Ste)		Su(Ste)
1849	ACUUACGGGAUAAGGAUUGGCUCUG	2	2	ncRNA		rRNA
1850	UCUGUGUGAACGAAAGCGAGGCGGC	2	0	coding gene		CG13369-RA|CG13369
1851	UUCGAUUUUCAACCGGACGCGUGU	2	1	no annotation		
1852	UCUUAUCGUCGUAGAACAAGCCCGAG	2	1	Su(Ste)		Su(Ste)
1853	UCUUGCCGAGGCAUAGAUAAAUUAC	2	0	no annotation		
1854	AUAAAAUGCCAGAAUUUUAUUGGGGUUC	2	no match			
1855	UCCGAACUACCACGGUGCCCCAGU	2	no match			
1856	GGAUCCAACUGUGAGGGGAUCCUCAA	2	no match			
1857	UAACAAGAUGCAUACGAUGUGGCG	2	1	transposon	DNA	1360
1858	CAGCACACAGAGGAGAGCAAAUGAGAGCA	2	no match			
1859	AUCCUCAACUGUGAGGGGAUCCUCA	2	no match			
1860	UCGGCUUUCCGGCGAGAACUCGAG	2	0	no annotation		
1861	GCUGAACGUAGUCUCUCGAGAGCGAG	2	1	transposon	LTR	invader1
1862	CCCGUACUCUGUGGUGACCAACA	2	0	transposon	LINE	baggins
1863	GCCGAACGUAGUCUCUCGAGAGCGAA	2	1	transposon	LTR	invader1
1864	UAUAUCUUCACCGUUGACUUGGCCC	2	1	Su(Ste)		Su(Ste)
1865	UCUCAUCGUUUUAGAACAAGCCCGA	2	2	Su(Ste)		Su(Ste)
1866	AGACUAGCUGGCCGAGUAUUUGAUAU	2	1	Su(Ste)		Su(Ste)
1867	CAGCACACAGAGGAGAGCAGAUGAGAGCGA	2	no match			
1868	GUUCCUCGGGAUCAAAGGCAACGAGUU	2	1	Su(Ste)		Su(Ste)
1869	CUAGCUGGCCGAGUAUUUGAUAUUC	2	0	Su(Ste)		Su(Ste)
1870	ACCGGAGAAGACGUUGAAUGGAUCGG	2	0	no annotation		
1871	UUUCUGACGAGAACUCGAGCCGCGU	2	0	no annotation		
1872	GGGUGCCCAACCAGACGACGUUGUC	2	0	coding gene		CG30494-RA|CG30494
1873	CAAGAUAUGACCUAAGCAUACAC	2	no match			
1874	GCCGAACGUAGCCUCGAGAGCGAG	2	no match			
1875	UUCUCAUCGUCGUAGAACAACCCCGA	2	1	Su(Ste)		Su(Ste)
1876	UCCUCAACUGUGAGGGAUCCUCCA	2	no match			
1877	UACAUCUCUGUAUGCACACCCCGUU	2	1	transposon	LINE	Circe
1878	UCUCCAUCGUCGUAGAACAAGCCCGAC	2	no match			
1879	UACAAUCUCCGAGCACGACCCCGAUCU	2	0	transposon	LINE	Circe
1880	UGUCACGCGGGAGACCGGGGUUCAAU	2	0	ncRNA		tRNA
1881	GGAUCAUCAACUGUGAGGGGAUCCUCAA	2	no match			
1882	CCGAACGUAGUCUCUCGAGAGCGAGA	2	0	transposon	LTR	invader1
1883	UUGGCGCAUACGAGUAAACGAAGGAA	2	0	transposon	LINE	Doc3
1884	CCUCAACUGUGAGGGAUCCUAAA	2	no match			
1885	UUAUUGUGCCACUGAGUCGACUAA	2	1	transposon	LINE	Cr1a
1886	UCUCAUCCUCGUAGAACAAGCCCCGA	2	no match			
1887	CCUGAUGAAGAAGAGAUGCCCAGUGA	2	0	transposon	LTR	ninja-Dsim-like
1888	CACACAGAGGAGAGCAGAUGAGAGC	2	no match			
1889	UAAUGGUAUCCGAGUCCUACUCUAG	2	1	transposon	LTR	blood
1890	UCUCAUCGUCGUAGAACCCCCCCG	2	no match			
1891	CGGGCCCAAACACAACCCAGCCCA	2	1	transposon	LTR	ninja-Dsim-like
1892	UUCUCAUCGUCGUAGAACAGCCCGA	2	no match			
1893	GUUCGGCUUUCCGGCGAGAACUCGAGC	2	0	no annotation		
1894	UUCACCGGGUUUUGGCACUUUGCC	2	0	transposon	LINE	R1-element
1895	UUAAAAAAUGACCUGGUGUUAAAACC	2	0	transposon	LTR	roo
1896	CCUCAACUGUGAGGGAUCCUCUAA	2	no match			
1897	UGAUAUGACCAGCAUGGACCUAACACU	2	0	Su(Ste)		Su(Ste)
1898	UGUUUCAUCGUUAGACGGCUCGGUCC	2	2	AT-chX		AT-chX-B
1899	UAGAAGAACAUUGAAGCGCUUGACU	2	0	Su(Ste)		Su(Ste)
1900	AUCUCAUCGUCGUAGAACAAGCCCUA	2	2	Su(Ste)		Su(Ste)
1901	UGUGACUGAUCUAGAGUACAAUGUUA	2	0	transposon	LTR	DM88
1902	UCGAGUUGCAGAAUUGAGGACCGCG	2	0	transposon	LTR	Quasimodo
1903	UAGACCUUUACCGUUGACUUGGCCCA	2	1	no annotation		
1904	AAAAAAAAAAAAGGAUCCUCAA	2	2	no annotation		
1905	CCCAGAUGCCUCAGGUUGCACCCAU	2	0	transposon	LINE	R1
1906	CCCGUACUCUGUGGUGACCAAAC	2	2	transposon	LINE	baggins
1907	UCCUCAACUGUGAGGGGAUCUCAA	2	no match			

1908	AGAAAAAAAAAAAAGGAUCCUCAA	2	no match			
1909	CAAGCCCUGGUAUUCUGAAGCCAAGA	2	1	repeat		trf
1910	UAAAGAAUCCUGACAAGAUCCCGGGCC	2	0	transposon	LTR	invader1
1911	UCUCCACCGUUGACUUGUCCCACACA	2	1	Su(Ste)		Su(Ste)
1912	GACGAGUCAGAUAUAACGGGCCUAAUC	2	1	transposon	LTR	diver2
1913	UGUUUCAUCGUUAGUCGGCUCGGG	2	1	AT-chX		AT-chX-B
1914	UAGAUCUUCACCGUUGACUAACCCCA	2	1	Su(Ste)		Su(Ste)
1915	UAUUCUGUGGAAGGGGGACCGGGAC	2	no match			
1916	UUGUUUCGGAGCGUUGGCCAGUUC	2	2	transposon	Low_complexity	F-element
1917	UCUAUCGUCGUAGAACAAGCCCGA	2	1	no annotation		
1918	AGAUCCUCAACUGUGAGGGGAUCCUCAA	2	no match			
1919	UCCGAACGUACCACGGUGCCCAG	2	2	transposon	Low_complexity	F-element
1920	UAGCGCUACUCCAGCCGGUGACCAUC	2	0	transposon	LTR	gypsy8
1921	UCUCAUCGCGUAGAACAAGCCCC	2	no match			
1922	UCUCAUCAUCGUAGAACAACCCCGA	2	1	Su(Ste)		Su(Ste)
1923	UCUCAUCGUCGUAGAACAAGCCCGACUGU	2	no match			
1924	AAAUGCCGGAAGCGAUCCCACUGCUU	2	2	coding gene		CG41580-RA|CG41580
1925	UCCGAACGUACCACGGUCGCCCCAGU	2	no match			
1926	UCGUCAUCGUCGUAGAACUAGCCCGA	2	2	Su(Ste)		Su(Ste)
1927	AUUCUUGAGAAUUUGGACGCCGUU	2	0	transposon	LINE	Cr1a
1928	UAAGACUUUGGCUCGUCCCUAUUU	2	0	transposon	LTR	opus
1929	AAAAAAAAAAAAACUA	2	0	transposon	LTR	qbert
1930	AUCUCACCGUCGUAGAACAAGCCCGA	2	2	Su(Ste)		Su(Ste)
1931	CUUCGGGAUAAGGAUUGGCUCUAA	2	1	ncRNA		rRNA
1932	AAUAACAAGAUGCAUACGAUGUGGCUCCA	2	0	transposon	DNA	1360
1933	AAUUCUCAUCGUCGUAGAACAAGCCCG	2	2	Su(Ste)		Su(Ste)
1934	UCUCAUCGUCGUAGAACAAACCCG	2	1	Su(Ste)		Su(Ste)
1935	UAAAUGCCAGAAUUUAUUGGGUUC	2	no match			
1936	UAUAAGAUUGGACCCAGUACACUGC	2	0	transposon	LINE	Doc
1937	UGUUUCAUCGUUAGACGGCUCGGGUA	2	2	AT-chX		AT-chX-B
1938	UAGAUCUUCACCUUUGACUAGCCC	2	1	Su(Ste)		Su(Ste)
1939	UUAAUAAAACAACGACCCGCUACCGGA	2	0	transposon	LTR	opus
1940	UAAAAUGCCAGAAUUUUAUUGGGGU	2	2	transposon	LINE	baggins
1941	UAGCACACAGAGGAGAGCAGAUGAGAGCA	2	no match			
1942	UUCAACAAGGAUAAUAUGCCCGCUC	2	0	coding gene		CG40601-RA|CG40601
1943	GGAUCCUCAACUGUGACAAA	2	2	no annotation		
1944	GAUGCCCCUGAUGAAGAAGAGAUGCCCA	2	0	transposon	LTR	ninja-Dsim-like
1945	UAUCUGUAUUUUAGGCAGGGUUGG	2	0	transposon	LINE	Rt1b
1946	GUCAUGCGAGGCGAACCGGGUUCA	2	no match			
1947	UUCCCAUCGUCGUAGAACAAGCCCG	2	1	Su(Ste)		Su(Ste)
1948	UAGAUCUUCACCGUUGACUUGGCCCAA	2	1	Su(Ste)		Su(Ste)
1949	UGUCCACCAAAGGCAUGCCCGAUA	2	1	transposon	LTR	opus
1950	UCUCAUCGUCGUAGAACUAGCCCUAG	2	1	Su(Ste)		Su(Ste)
1951	AUCAUCUUCUUCUCGACGUUGACCAGC	2	1	transposon	LTR	ninja-Dsim-like
1952	UCCGAGGAACCAAUUGAUCCAGCUC	2	1	Su(Ste)		Su(Ste)
1953	CUCAUGUGAGGGGAUCCUCAA	2	no match			
1954	CGAAACUCAACUGUGAGGGGAUCCUCAA	2	no match			
1955	AAAUCCCAACUGUGAGGGGAUCCUCAA	2	no match			
1956	GUACAUCCAGGAAGAUUGCAUUGC	2	0	transposon	LINE	BS3
1957	CAACUGUUAGGGGAUCCUAA	2	no match			
1958	UCUCAUCGUCGUAGAACAAGCCCGCUGUAACCUCGGGUCAAU	2	no match			
1959	CGCAGGAUGUCCGCCGCCUUUGCC	2	no match			
1960	UCUCUAUCGUCGUAGAACAAGCCCGUA	2	no match			
1961	AUCUCAUCGUCGUAGAAACAAGCCCG	2	no match			
1962	CGAAAUAGAUCUUCACCGUUGACUUGGCC	2	no match			
1963	UCUUGACGAGUGCCGAGAUGUUGGU	2	0	transposon	LINE	G6
1964	UCUCAUCGUCGUAGAGCAAGCCCCGA	2	no match			
1965	UAGAUCUUCACCGUUGACUGGCCCCA	2	1	Su(Ste)		Su(Ste)
1966	UAUGGGCCCAGUAAACAUUUAGUAU	2	0	ncRNA		alphagamma-element
1967	UCGACAACCAGAGAUCUAGAGGGCC	2	0	no annotation		
1968	GGAUCCUCAACUUGAGAUCCUCAA	2	no match			
1969	UAGUUUCUAUCGUUAGACGGCUCGGGC	2	no match			
1970	UAAAAUCCUGACAAGAUCCCGG	2	2	transposon	LTR	invader1
1971	UCUCAUCGUCGUAGAACAAGCCCGACUGUAACUCCGGGUCAAU	2	no match			
1972	UAGAUCUUCACCGUUGACUAGCCCCCUC	2	2	Su(Ste)		Su(Ste)
1973	UAAGAUUGGACCCAGUACACUGC	2	0	transposon	LINE	Doc
1974	UGGUUUCAUCGUUAGACGGCUCGGG	2	1	no annotation		
1975	AUCGUCUCGGGAGUGAAAUACUCUAUCGUCGUAGAACAAGCCCGG	2	no match			
1976	UUCUCAUCCUCGUAGAACAAGCCCGA	2	1	Su(Ste)		Su(Ste)
1977	UCCGAGGAACCAAUUGAUCUAGCUG	2	1	Su(Ste)		Su(Ste)
1978	AGUUUGACUGGGGCGGUACAUCUCU	2	0	ncRNA		rRNA
1979	AUCGUCUCGGGGAUGAAAAUCUCAUCGUCGUAGAACAAGCCCGAG	2	no match			
1980	UUCUCACUCGUCGUAGAACAAGCCCGA	2	no match			
1981	UCAAGCCCGGGUAUUCUGAAGCCA	2	0	repeat		trf
1982	AUCGUCAUCAUCGUAGAACUAGGCCCGA	2	no match			
1983	UGAUUCCGAUCUUGGACAUUCCCGAUG	2	0	transposon	LTR	roo
1984	UUCGGCUUUCCGGCGAGAACUCGAG	2	0	no annotation		
1985	AAACUUCGGGAUAAGGAUUGGCUC	2	1	ncRNA		rRNA
1986	UGUCUCAUCGUCGUAGAACUAGCCCGA	2	1	Su(Ste)		Su(Ste)
1987	CGUUGGACCAAUUUCCGAACAUACGC	2	no match			
1988	AUGUUUCAUCGUUAUACGGCUCGGGC	2	2	AT-chX		AT-chX-B
1989	UGUUUCAUCGUUAGACGGCUCCGCC	2	2	AT-chX		AT-chX-B
1990	AUCUCAUCGUCGUAGAAACAAAGCCCCGA	2	no match			

1991	CAAAGGUGGAAAUGAAGAAAGUACA	2	no match			
1992	ACCGAUAUGGUUGGACCACAAUCUUU	2	0	ncRNA		7SLRNA-RA
1993	UGUUUCAUCGUUAGACGGCUCGAGC	2	1	AT-chX		AT-chX-B
1994	AUCGUCGUCGGGAUGAAAUCUCAUCGUCGUAGAACAAGCCCG	2	no match			
1995	UGUUUCAUCGUUAGACGGCGUCGGGC	2	no match			
1996	CAGUAUUGGUCCAAGAACGCUGCC	2	0	transposon	LINE	G
1997	AAAGUGGAAAUGAAGAAAGUACA	2	2	no annotation		
1998	AUCUCAUCGUCGUAGAACAAGCCCGAAA	2	2	Su(Ste)		Su(Ste)
1999	UCCGAUCUCCUGAGUCGGACUUCGAG	2	0	transposon	LINE	R1-element
2000	UGUUUCAUCGUUAGACGGCUAGGG	2	1	AT-chX		AT-chX-B
2001	AAAUUUCUCGACGACCCGACUUCC	2	0	transposon	LINE	G2
2002	CUCUCUACCGGACUUCGUUUGACCCC	2	0	no annotation		
2003	ACGCCGGAGGUCGCGGGUUCGAGUCCC	2	no match			
2004	UGAUACCGAGAAUGACAAUGGUUG	2	0	transposon	LTR	gypsy3
2005	UCUCAUCAUCGUAGAACAAGCCUGA	2	1	Su(Ste)		Su(Ste)
2006	CAACUGUGAGGGGAUUCCUCAA	2	no match			
2007	UCCAUGCAGCAGAAGCGACAGAGU	2	0	transposon	LINE	baggins
2008	UGUCAUCGUCGUAGAACAAGCCCG	2	1	Su(Ste)		Su(Ste)
2009	UCGUCAGUCGUACGGUAGAACAAGCCCGGACUG	2	no match			
2010	CCUUAACUGUGAGGGCAUCCUCAA	2	no match			
2011	CAGCACACGGAGGAGAGCAGAUGAGAGCA	2	no match			
2012	AUCCUCAACUGUGAGUGAAAA	2	no match			
2013	UCUCAUCAUCGUAGAACAAGCCCAA	2	1	Su(Ste)		Su(Ste)
2014	AGGUGGAAAUGAAGAAAGUACG	2	no match			
2015	GUAGGCUCGAUGGUGACGACCGACC	2	1	transposon	LINE	G2
2016	UGGACCAUAAGCCUGCCGUCUGCAGA	2	0	no annotation		
2017	CAGCAUGAUCAACAUCAAUGGCCCCA	2	1	coding gene		CG9056-RA|CG9056
2018	UCUCAUCGUCGUAGAACAACGCCCGUA	2	no match			
2019	UCACAUCCGGUAUGAAGAUUGGAGGC	2	0	transposon	LINE	G6
2020	CAACUGUGAGGGGAUCCUCAACUG	2	no match			
2021	GACCUUUACCGUUGACUUGGCCC	2	1	no annotation		
2022	UAAACCGUCAUUCCAGCGCAAGUUGCCCC	2	0	transposon	LTR	ninja-Dsim-like
2023	AUAACAAGAUGCAUACGAUGUGGC	2	0	transposon	DNA	1360
2024	UUCUCAUUGUCGUAGAACAAGCCCGA	2	1	Su(Ste)		Su(Ste)
2025	CGAAAUUCUCAUCGUCGUAGAACAAGCCCGA	2	no match			
2026	UCGGCCCGAAAUAGGUUCCUGAAG	2	0	transposon	LTR	flea
2027	AGAAAAAAAACCUGAAGAUGAUCCUGAGUCGC	2	no match			
2028	AAAGCACCGUUUUCCCGUCCGAUCAA	2	no match			
2029	UUCGUGACGAGAACUCGAGCCGCGAAAAA	2	no match			
2030	UCUCAAGACCCACGGGAUGACUUC	2	0	transposon	LINE	BS
2031	CGGCACACAGAGGAGAGCAGAUGAGAGCA	2	no match			
2032	UCUCAUCCUCGUAGAACAAGCCCGAA	2	1	Su(Ste)		Su(Ste)
2033	AACUGCAUAUCGAGAGGUCCUCGGUU	2	no match			
2034	AUCUCAUCGUCGUAAGAACAAGCCCGA	2	no match			
2035	GAUCUCCUGAGUCGGACUCCGAG	2	0	transposon	LINE	R1
2036	UUCAACUGAAGGAGUACGAAUGAUCG	2	0	transposon	LINE	R1-element
2037	UCCUGACGAAGAGCGCAAGAAGA	2	0	transposon	LTR	opus
2038	UUAACACCUCGAGUACGUCUGCUC	2	0	transposon	LTR	Max
2039	CAAAUCCCUUUCGACAACCAGAGAUC	2	0	no annotation		
2040	UCUCAUCGUCGUAGAACAAGCCCGUACUGUAACCUCGGGUCAAU	2	no match			
2041	UCUGCCCGAUGACAAAAAUGGCCCG	2	0	transposon	LTR	mdg1
2042	UUACCGUACGAUCAGGACAACCUUC	2	2	transposon	LTR	mdg1
2043	UAAUGCCCCGAAGGUUGCCCUCUG	2	0	transposon	LINE	R1-element
2044	UCUCAUCGUAGUAGAACAAGCCCGA	2	1	Su(Ste)		Su(Ste)
2045	UACAAUCUCCGUGCACGACCCCGA	2	0	transposon	LINE	Circe
2046	AUGUUUUCAUCGUUAGACGGCUCGGGGC	2	no match			
2047	UUAAUGCCGGAAGCGAACCCACUG	2	1	AT-chX		AT-chX-B
2048	UAGGCUGAGCGGAGAAUCAGAGUUG	2	1	transposon	Low_complexity	F-element
2049	UCCGAUGUUCUGAAACUGCGUGACCC	2	0	transposon	LTR	invader3
2050	UCUAGGCUGAGCGGAGAACCAGAGUUG	2	0	transposon	Low_complexity	F-element
2051	UGCAUGCCCUGAAGACCAUUACUC	2	0	coding gene		pseudogene
2052	AGAAAAAACCUGAAGAUGAUCCUGAGUCGCA	2	no match			
2053	UAAACAAUUGAAAGAUAAGACCCA	2	0	transposon	LINE	Cr1a
2054	GGAUCCUCAACUGUGAGUCAA	2	no match			
2055	UCUUCUCGACGUUGACCAGCCUCCU	2	0	transposon	LTR	ninja-Dsim-like
2056	UCUCAUCUUCGUAGAACAAGCCCCGA	2	no match			
2057	CUCGAACUGUGAGGGGAUCCUCAACG	2	no match			
2058	UCAUCGUCGUAGAACAAGCCCAA	2	1	Su(Ste)		Su(Ste)
2059	CGAAAUAGAUCUUCACCGUUGACCUUGGC	2	no match			
2060	GCAGUUUGGCAACUACGAGAUCGACACC	2	0	coding gene		CG3025-RA|mof
2061	UUCUCAUCGUCAUAGAACAAGCCCGA	2	1	Su(Ste)		Su(Ste)
2062	AACUCAACUGUGAGGGGAUCCUCAA	2	no match			
2063	CAAACUGUGAGGGGGAUCCUCAA	2	no match			
2064	ACUCAGCCACGAAUUAGCGAU	2	no match			
2065	UAAAUUCCAGAAUUUUUAUUGGGGU	2	no match			
2066	UUCUCAUCGUCGUAGAACAAGCCCGAA	2	0	Su(Ste)		Su(Ste)
2067	UCAACUGUGGUGGAUCCUCAA	2	no match			
2068	CCUCGAUAGUAUAGUGGUUAGUAUCACC	2	1	ncRNA		tRNA
2069	UUCCUGGAGAAUUUGCCCGUAUGUC	2	0	coding gene		CG9925-RA|CG9925
2070	GACCGAACACUUUGAACCAAGUAU	2	0	Su(Ste)		Su(Ste)
2071	CACACAGAGGAGAGCAGAUGAGAGGCA	2	no match			
2072	GCCCCUGAUGAAGAAGAGAUGCCCA	2	0	transposon	LTR	ninja-Dsim-like
2073	UCAACUGUGGAGGGGAUCCUCAA	2	no match			

2074	CACACUCAUCAACCAAAGGACGUAGGAC	2	0	no annotation		
2075	UAGAUCUUCCCGUUGACUAGCCCCA	2	no match			
2076	UCUAGGCUGAGCGGAGAACCAGAG	2	0	transposon	Low_complexity	F-element
2077	AACUGAGGGGAUCCUCAA	2	no match			
2078	CGUUGACCAAUUUCCGAACAUAA	2	1	transposon	LTR	DM88
2079	AUGUGACUGAUCUAGAGUACAAUGU	2	1	transposon	LTR	DM88
2080	CAGAUCUUCACCGUUGACUAGCCCCA	2	1	Su(Ste)		Su(Ste)
2081	UUUACCGGAAGAAUCGCGGAUGUGAC	2	0	no annotation		
2082	GCUGGCCGAGGAUCCGAUUUUCUU	2	0	Su(Ste)		Su(Ste)
2083	CGAAAUCAACUGUGAGGGGAUCCCUCAA	2	no match			
2084	UUUUAAGGAUGAUGAUACCGAUUAAA	2	1	no annotation		
2085	UUCAUAUGAAGAAAUGACCCCUGAAA	2	0	transposon	LTR	McClintock
2086	GCUUUCUGACGAGAAUUCGAGCCGCG	2	1	no annotation		
2087	CUCUCAUCCGUCGUAGAACAACGCCCGUA	2	no match			
2088	UCGUCAUCGUCGGUAGAACAAGCCCGGA	2	no match			
2089	GUCGUCGGGAUGAAA	2	1	transposon	LTR	ninja-Dsim-like
2090	GGUCCCCUGUGUGCCGUUGACGGA	2	0	transposon	LINE	Rt1b
2091	UCUCAUCUCCGUAGAACAAGCCCGA	2	2	Su(Ste)		Su(Ste)
2092	UGAAAAUGAUGAUACCGACUAAA	2	2	no annotation		
2093	AUCGUCGUCGGGAUAGAAAAUAGAUCUUCAACCGUUGAACUUGCC	2	no match			
2094	UACAAUCUCCGAGCAACGACCCCGA	2	no match			
2095	CCAAUUGAUCCAGCUGCUGUUGU	2	0	Su(Ste)		Su(Ste)
2096	GGAUCCUCAACUGUGAGGGUAUCCUCAA	2	no match			
2097	UAAUGCCCCUAAGGUUGCCCGA	2	2	transposon	LINE	R1-element
2098	AUCGUCUCGGGAGUGAAAUACUCUAUCGUCGUAGAACAGCCCGA	2	no match			
2099	UCCGAGGAACCAAUUGAUCCAGCUGCU	2	0	Su(Ste)		Su(Ste)
2100	UUAUACCCAGGUAUUACCACUAU	2	no match			
2101	UGAUUAAAGCCUUCGCGUCUUCCACG	2	0	AT-chX		AT-chX-A
2102	GCCCGUACUCUGUGGUGACCAACA	2	1	transposon	LINE	baggins
2103	UCUCAUCUUCGUAGAACAAGCCCGU	2	2	Su(Ste)		Su(Ste)
2104	UCUUCUUCUUCUCGACGUUGACCA	2	1	transposon	LTR	ninja-Dsim-like
2105	GGAUCCUCAACUGUGAGGCGAUCCUCAA	2	no match			
2106	GCCCCAGUGGCCUAAUGGAUAAGGCAUC	2	0	ncRNA		tRNA
2107	UCAACUGUGAGGGGAGUCCUCAA	2	no match			
2108	CCCCGUACUCUGUGGUGACCGACACA	2	1	transposon	LINE	baggins
2109	AGUCUCAUCGUCGUAGAACAAGGCCCGA	2	no match			
2110	CUGAGACACGCAACAGGGA	2	2	no annotation		
2111	UUAGAAAUGUAAGUCACGCCCUGAGG	2	1	transposon	LTR	opus
2112	UCCUGUUGAUGUGAGGGCCCAAGU	2	2	transposon	LINE	G2
2113	AUCUCAUCGUCGUAGAACAACCCCGA	2	2	Su(Ste)		Su(Ste)
2114	UGUGAUCGGCAUCGAAAUGAAAAC	2	0	transposon	LINE	R1-element
2115	AUCUCAUUGUCGUAGAACAAGCCCGA	2	2	Su(Ste)		Su(Ste)
2116	UUCCCCUGCGUAGGCGCUGAGCUCU	2	0	transposon	LINE	R1-element
2117	AGAGGCGCCGUAGUUACCACCCCAC	2	0	no annotation		
2118	UCUCAUCGUCGUAAACAAGCCCGA	2	no match			
2119	AAAUCCUCAACUGUGAGGGGUCCUCAA	2	no match			
2120	UAAGGAUGAUGAUACCGAUUAAAGGCAA	2	0	no annotation		
2121	AUUCGUCAUCGUCGUAGAACAAGCCCGG	2	no match			
2122	CGUUGACCAAUUUCCGAACAUACA	2	1	transposon	LTR	DM88
2123	UCCGAACGUACCACGGUGCCCCCAGU	2	no match			
2124	UCGGCCCGAACCCUUCUGGUGUGU	2	0	transposon	LINE	baggins
2125	CAAGAUAUGACCAAAACAAAGCAACG	2	0	Su(Ste)		Su(Ste)
2126	UCCGAGGAACCAAUUUAUCCAGCUGC	2	1	Su(Ste)		Su(Ste)
2127	UAAGCGUUCGUUGACGUUGAUGG	2	0	transposon	LTR	HMS-Beagle
2128	UGGACUUUUUCCGAUGAGCCAGUG	2	no match			
2129	UAAAUCUUCACCGUUGACUAGCCCC	2	1	Su(Ste)		Su(Ste)
2130	GCAUAUCAUUCACGCCACGCUCCGA	2	0	Su(Ste)		Su(Ste)
2131	CAGCACACAGAGGAGAGCAGGUGAGAGCA	2	no match			
2132	CACACGCAUCCUAGAAUCGACCGACCA	2	0	transposon	LTR	flea
2133	CUCAACUGUGGAGGGGAUCCUCAA	2	no match			
2134	CAGCACACAGAGGAGAGCAGAUGAGAGCACUGUA	2	no match			
2135	UACCUUAUUAAUGGAUUGGCUGAGA	2	0	transposon	LTR	Stalker
2136	GCCUCCAACCCAAAACUCCCGAACUC	2	0	transposon	Low_complexity	F-element
2137	CAGCACACAGAGAAGAGCAGAUGAGAGCA	2	no match			
2138	CAAGAUAUGACCAAAGCAAAGACACA	2	1	transposon	LINE	Cr1a
2139	GUGGACCUCGAGACCCUCGACCGUUC	2	0	transposon	LTR	G4
2140	UUGAUUCGGAAUACCUUCAAUGAGGAG	2	0	transposon	LTR	diver2
2141	UCUCAUCGUACGUAGAACAAGCCCG	2	no match			
2142	CUUCUCAUCGUCGUAGAACAACGCCCGA	2	no match			
2143	UUCGGCCCGAACCCUUCUGGUGUG	2	0	transposon	LINE	baggins
2144	ACCUAACGAGGUAGAUGACUCCUGGAA	1	1	transposon	LINE	Cr1a
2145	AAAAAUCCUCAACUGUGAGCGGAUCCUCAA	1	no match			
2146	UCCGAGGAACCAAUUGAUCCAGCUGU	1	1	Su(Ste)		Su(Ste)
2147	UGUUUCAUCGUUAGUCGGCUCGGGC	1	1	AT-chX		AT-chX-B
2148	UAUGGGCUAAUCCGACUGAGAUGAAG	1	0	no annotation		
2149	GAACGCCUCAUUGGUGAUCCGGCCA	1	0	coding gene		CG31359-RA|Hsp70Bb
2150	UAGUUUCAUCGUUAGACGGCUCGGGC	1	1	no annotation		
2151	CGUAGACCAAUUUCCGAACAUACGCC	1	1	transposon	LTR	DM88
2152	CAUCUUCUUCUCGACGUUGACCAG	1	0	transposon	LTR	ninja-Dsim-like
2153	AUAAUGUCCCUUAAAUCAGCACUCG	1	0	no annotation		
2154	AUGAAUACCUCGGGUACGUCUGCUCUC	1	1	transposon	LTR	GATE
2155	CUAUAAGGAUGAUGAUACCGAUUA	1	1	no annotation		
2156	CAAGAUAUGACCAUAGCAAAGACAC	1	1	transposon	LINE	Cr1a

2157	UAGAUUGAACUGAAUUGCAAUUUAU	1	0	no annotation		
2158	UAUAUGCUCUCGAAUGUAUCGCCCGA	1	no match			
2159	UUCUCGCGGAUGUAUAUUGGCGGAGGC	1	0	transposon	Low_complexity	F-element
2160	CUACUCGAAUGAGUGCUUGAUAUGACCA	1	0	Su(Ste)		Su(Ste)
2161	UUUUUCAUCGUUAGACGGCUCGGGCG	1	1	AT-chX		AT-chX-B
2162	UUCUCAUCGUCGUAGAACAAGCCGGA	1	1	Su(Ste)		Su(Ste)
2163	GCAAGCUAACUACAGACGGGGUGA	1	0	transposon	LINE	R1
2164	UGUUUCAUCGUUAUACUGCUCGGGC	1	2	AT-chX		AT-chX-B
2165	AACUGACCCAGUCGGAUCUAUUGUAC	1	1	transposon	LINE	baggins
2166	GCACACAGAGGAGAGCAGAUAGAGCA	1	no match			
2167	AUUGGACCUAGGCUUCUGUUGACUCU	1	0	transposon	LTR	flea
2168	AACUAGUGAGGGGGAUCCUCAA	1	no match			
2169	UCUGACAAUGGUUCCCAAUUUCGAGC	1	0	transposon	LINE	Rt1b
2170	AACUUCGGGAAAAGGAUUGGCUC	1	1	ncRNA		rRNA
2171	AGGAUGAUGAUACCGACUAAAGGCAA	1	0	no annotation		
2172	AGAUCCAACUGUGAGGGGAAUACCUCCAAC	1	no match			
2173	UCUCAUCGUCGUAGAGCUAGCCCGA	1	1	Su(Ste)		Su(Ste)
2174	UCUCAUCGUCGUAGAACUAGCCCGG	1	1	Su(Ste)		Su(Ste)
2175	UCUCCGAUUGGUAUUAGCGAGUUUA	1	0	transposon	LTR	opus
2176	CCCCGUACUCUGUGGUGACCAACAC	1	0	transposon	LINE	baggins
2177	GGCAACUGUGAGAGGGGAUCCUCAA	1	no match			
2178	AUUUUAAACAAUUGAAAGAUAAGACCC	1	1	transposon	LINE	Cr1a
2179	AUUCGUCAUCGUACGUAGAACAAGCCCGGA	1	no match			
2180	CCCCUGAAGUCCAAUGCACUGUAUAAA	1	0	transposon	LINE	baggins
2181	GCACACAGAGGAGAGCAGAUGAGAGCCA	1	no match			
2182	UCCGAGGAACCAAUUGAUCCAGCCCC	1	2	Su(Ste)		Su(Ste)
2183	AUCGUCGUCGGGAUAGAAACUCAUCGUCGUAGAACAAGCCCGA	1	no match			
2184	UCUCAUCGUCUUAGAACAAGCCCAA	1	2	Su(Ste)		Su(Ste)
2185	AGGACCCCUGCCGUGUGUGAAGCCCC	1	0	coding gene		CG33340-RA|CG33340
2186	UGGAUCUUCACCGUUGACUAGCCCCA	1	1	Su(Ste)		Su(Ste)
2187	UGCAUAUCAUUCACGCCACGCUCCGA	1	0	Su(Ste)		Su(Ste)
2188	AUAGAUCUUCACCGUUGACUAGCCCCC	1	2	Su(Ste)		Su(Ste)
2189	CAAAUGGCUGUGAAGCCGGCAAUGCCCCG	1	0	transposon	LINE	INE-1
2190	AGAUCUUCACCGUUGACUUGCCCCGC	1	0	Su(Ste)		Su(Ste)
2191	CGAAAUCUCAUCGUUCGUAGAACUAACGCCCGUACUGUAACCUCGC	1	no match			
2192	GUUCAAGGUUACCCGAGGAUGCUGU	1	1	repeat		trf
2193	UCUCAUCGUCGUAGAACCAAGCCG	1	no match			
2194	UAUAUCUUCACCGUUGACUAGCCCCA	1	1	Su(Ste)		Su(Ste)
2195	CCAGGCCCAUCUGGUUGAACGUGUCC	1	0	Su(Ste)		Su(Ste)
2196	UAACAAGAUGCAUACGAUGCGGC	1	1	transposon	DNA	1360
2197	CCUGUGAGGGAUCCUCAA	1	2	coding gene		CG13650-RB|CG13650
2198	UCUUCCUGGAAGAUGUGACAGUGGC	1	0	coding gene		CG13397-RA|CG13397
2199	UCCAACUGAGGAGUACGAAUGAUC	1	no match			
2200	CUUUGACCACUGUUAAUCCAUGCU	1	0	transposon	LINE	Cr1a
2201	UCGUUAACACUUCCCAGAAUGCACAC	1	2	transposon	LINE	Cr1a
2202	AACAAGAUAUGACCUAAGCAAAGACAC	1	2	transposon	DNA	1360
2203	UCUCAUCAUCGUAGAACAGCCCGA	1	no match			
2204	GGCCUCAACUGUGAGAGGAUCCUCAA	1	no match			
2205	GCUUGAUAUGACCAGCUUGGACCGAA	1	0	Su(Ste)		Su(Ste)
2206	CUCAACUGUGAGGGGAUCCUCAAA	1	no match			
2207	UCCGAGGAACCAACUGAUCCAGCUG	1	1	Su(Ste)		Su(Ste)
2208	CAGCACACAAAGGAGAGCAGAUGAGAGCA	1	no match			
2209	UAUAAUCGCCCCGAAGGGUUUGCCCUCUG	1	no match			
2210	AAAAUCUCAUCGUCGUAGAACAAGCCCG	1	no match			
2211	UAGAUCUUCACCGUUGACUUGGCCCUGUAACUCGGGGUCAAU	1	no match			
2212	ACGGAGCGAUGUUCGGGCCCAGCUU	1	0	repeat		trf
2213	CCCUGAUGAAGAAGAGAUGCCCAGU	1	0	transposon	LTR	ninja-Dsim-like
2214	UAGAUCUUCACCGUUUACUUGCCCCC	1	1	Su(Ste)		Su(Ste)
2215	AUCUUCACCGUUGACUUGGCCCACACA	1	0	Su(Ste)		Su(Ste)
2216	UACAAUCUCCGAGCACGACCCCGAUA	1	0	transposon	LINE	Circe
2217	ACUAGCUGGCCGAGUAUUUGAUACU	1	1	Su(Ste)		Su(Ste)
2218	UGAGCAUGACCCUGGACACCACCAAG	1	0	coding gene		CG9327-RA|Pros29
2219	UACAUCAUCGUCGUAGAACAAGCCCG	1	2	Su(Ste)		Su(Ste)
2220	CUGAGAAGACUAUUGUACCCGUCGAAUU	1	no match			
2221	AGUAGUGGCGAGCGAAAUGGGAUGAGC	1	no match			
2222	UCCUCAACUGUAAGGGGAUCCUCAA	1	no match			
2223	UAACUGGAAUAGUUUUGACCGUUGA	1	0	transposon	LTR	opus
2224	AUCCUCAACUGUGAGGGGAUCUCAA	1	no match			
2225	UCGUCAUCGUACGUAGAACAAGCCCGG	1	no match			
2226	UUGAUUGUCUUGAGAACCAGCAGC	1	1	ncRNA		pseudogene
2227	GGCAGCUGUGAGGGGAUCCUCAA	1	no match			
2228	UCAUCGUCGUAGAACAAGCCCGAGG	1	0	Su(Ste)		Su(Ste)
2229	GAUCUUCACCGUUGACUUGGCCCACAU	1	1	Su(Ste)		Su(Ste)
2230	AUCGUCUCGGGAAAUGAAAUCUCAUCGUCGUAGAAGAAGCCCGA	1	no match			
2231	UCCGAACGUACCACGGUGCCCCAGUCUGUAACUCGGUCAAU	1	no match			
2232	AUUUACCCGACGUUACAGGCCCGACGCCGUCUAACGAUGAAACAAU	1	no match			
2233	GACUAGUGGCCGAGGAUCCGAUUUUC	1	no match			
2234	AUCUCAUCGUCGUAGAAAACAACGCCCCGA	1	no match			
2235	UCUCCAUCGUCGUAGAACAAGCCCG	1	2	no annotation		
2236	AAUAUUGGCGCGGUCAUCGGAACUG	1	0	transposon	LTR	ninja-Dsim-like
2237	UAUGCCCCGUACUCUGUGGUGACCG	1	1	transposon	LINE	baggins
2238	UCAACUGUGAGGGGAGGAUCCUCAA	1	no match			
2239	UCAAUUGUGAGGGGAUCCUCAA	1	no match			

2240	UACUGAGUCGUUUGCCGAGAGCUGAA	1	1	transposon	LINE	Rt1b
2241	GGCAACUGUGAGGGAUCCUCAA	1	no match			
2242	GGAUCCGAACUGUGAGGGGAUCCUCAACG	1	no match			
2243	CUCUACUGACUUCAGUAAGCCCCGU	1	0	transposon	LINE	baggins
2244	UACAACUCUGUAUGCACACCCCGUUA	1	0	transposon	LINE	Circe
2245	AUCGUCUCGGGAGUGAAAAUACUCAUCGUCGUAGAACAAAGCCCG	1	no match			
2246	UCAACUGAAGGAGUACGAAUGAUCG	1	0	transposon	LINE	R1-element
2247	UCUUAUCGUCGUAGAACAAGCCCGAC	1	2	Su(Ste)		Su(Ste)
2248	AAAAAAAAAAAAGGAAAUCCUCAA	1	2	no annotation		
2249	GCACACAGAGGAGAGCAGGUGAGAGCA	1	no match			
2250	UUCUUUGAAGAUGGCCCACAUAA	1	0	repeat		trf
2251	UCUCAUCUUCGUAGAACAAGCCCGAU	1	1	Su(Ste)		Su(Ste)
2252	GCCCCGUACGUCUGUAGGUGACCAA	1	no match			
2253	UCUCAACUGUGAGGGGGAUCCUCAA	1	no match			
2254	AUUUUUUCAUCGUUAGACGGCUCGGGC	1	2	no annotation		
2255	UUCAAGGUUACCCACGGAUGCUGUGC	1	2	repeat		trf
2256	UUCUCAUCGUCGUAGAACAAGCCCGC	1	1	Su(Ste)		Su(Ste)
2257	UAACAAGAUGCAUACGAUGUG	1	0	transposon	DNA	1360
2258	UAGAUCUUCACCUUUGACUUGCCCC	1	1	Su(Ste)		Su(Ste)
2259	AGACUAGCUGGCCGAGGAUCCGAUU	1	1	Su(Ste)		Su(Ste)
2260	UGUUUCGUCGUUAGACGGCGUCGGG	1	no match			
2261	GCCCUGGGAGGAAUACUCCGAGAACGGACC	1	0	coding gene		CG12072-RA|wts
2262	CAGACACAGAGGGAGCAGAGAGAGCA	1	no match			
2263	UGUUUCAUCGUUAGACGGUCUCGGGC	1	no match			
2264	GUUCCUCGGGAUCAAGGGCAACGAGUU	1	0	coding gene		CG40635-RA|CG40635
2265	UUUCAUCGUCGUAGAACAAGCCCGAG	1	0	no annotation		
2266	AAGAGCGUGACCACGGACAUACCCU	1	1	coding gene		CG8230-RA|CG8230
2267	AUCGUCUCGGGACUGAAACUCUCAUCCGUCGUACGAACAACCGCC	1	no match			
2268	GGAUACCUCAACUGUGAGGGGGAUCCUCAA	1	no match			
2269	GCCCCGUACGUCUGUAGUCGACCAA	1	no match			
2270	UGGAGAGAACGUGAGACUACAAACGG	1	0	no annotation		
2271	GACGGAGCGAUGUUCGGGCCCAG	1	0	repeat		trf
2272	UCUUACAAUCUCCGAGCAAGACCCCGA	1	0	transposon	LINE	Circe
2273	UGCUGACGGGGCCCGAAAACACGA	1	0	transposon	LTR	opus
2274	AAAACCUGAAGAUGAUCCUGAGU	1	0	transposon	LINE	Helena
2275	UCAACUGUGAGGUGAUCCUCAA	1	no match			
2276	AAGUAAGCCCGAAGAUUGUCCAAU	1	0	transposon	LTR	opus
2277	UCCAAAGGACUCGAAAAACCUCUUGA	1	no match			
2278	CCUCAACUGUGAAGGGGAUCCUCAA	1	no match			
2279	UAAGUAAGCCCGAAGAUUGUCCAAUGAC	1	0	transposon	LTR	opus
2280	UCUACAUCGUCGUAGAACAAGCCAA	1	no match			
2281	UAGAUCUUACCGUUGACUAGCCCCA	1	no match			
2282	GUCUCAUCGUCGUAGAACAAGGCCCGA	1	no match			
2283	CCCAAGUACAAGCCCGGCCGAAAGACC	1	no match			
2284	UCUCAUCGUCGUGACAAGCCCGA	1	no match			
2285	CCCGAUGAGAAGAAUGGCCCGUAAAGC	1	0	transposon	LINE	RR49783
2286	UGAUGACCGAAAUUUGGAAAAACGA	1	0	repeat		ARS406_DM
2287	UCGCAUCGUCGUAGAACAAGCCCGAG	1	0	Su(Ste)		Su(Ste)
2288	AUGUUUCACCGUUAGACGGCUCGGGAA	1	no match			
2289	CUCAACUGUGAGGGACCACAA	1	no match			
2290	UGGCCGAGGAUCCGAUUUUCUU	1	0	Su(Ste)		Su(Ste)
2291	AAAAUGCCAGAAUUUUUAUUGGGGGUUC	1	no match			
2292	CUCGCGCCGCCUGCGC	1	1	coding gene		CG3941-RA|pita
2293	GUGUGAGGGGAUCCUCAA	1	2	coding gene		CG9684-RA|CG9684
2294	UGUGUCAUCGUUAGACGGCUCGGGC	1	1	AT-chX		AT-chX-B
2295	UCUCAUCGUCGUAGAACAAGCCCGUUGUAACUCGGGGUCAAUAUC	1	no match			
2296	UUCCGAGGAACCAGUUGAUCCAGC	1	1	Su(Ste)		Su(Ste)
2297	CCUCAUCGUCGCAGAACAAGCCCGA	1	2	Su(Ste)		Su(Ste)
2298	UCUCAACCGUAGAACAAGCCCGA	1	no match			
2299	UAAAUCUCAUCAUCGUAGAACAAGCCCGA	1	no match			
2300	UCGCAUCGUCGUAGAACAACGCCCG	1	no match			
2301	CUCAACUGUGACCUCAA	1	2	transposon	LTR	gypsy
2302	UUAAUAAGGAUGAUGAUACCGAUU	1	2	no annotation		
2303	UCUCAUCGUCGUAGACAAGCCCGU	1	no match			
2304	CAAGAUAUGAUCAAAGCAAAGACA	1	0	transposon	DNA	1360
2305	AAACCAAGGGGCGCGAGAUAACUCUCU	1	no match			
2306	UGGGCUUUCUGACGAGAACUCGAGC	1	0	no annotation		
2307	UGCAUAGCCGAAUCGAGACCCAGG	1	0	coding gene		CG14025-RA|Bsg25D
2308	CUCUCAUCCGUCGUAGAAUAACGCCCGA	1	no match			
2309	CAGCACACAGAGGAGAGCAGAUGAGGAG	1	no match			
2310	AAACCUAAGUUACUUACGGGCUCGACA	1	2	Su(Ste)		Su(Ste)
2311	AUCCGAUCUCCUGAGUCGGACUCCGAGCU	1	0	transposon	LINE	R1-element
2312	UAGAUUUUCACCGUUGACUAGCCCC	1	1	Su(Ste)		Su(Ste)
2313	UCUCAUCGUCGUAGAACAAGCCCCGACUGUAAACUCGGGGUCAAU	1	no match			
2314	UCCGAUCUCCUGAGUCGGACUCCGAGCUC	1	0	transposon	LINE	R1
2315	CAAAGUACUCGAGGCAGCUAAGCGAA	1	0	transposon	DNA	pogo
2316	GAUACAGAACCCGUGGGCAGCAGCGA	1	0	transposon	LTR	297
2317	UAAUGACCGAAAUUUGGAAAAAAUA	1	0	repeat		SAR_DM
2318	ACCGAUAACGCCCACCGAGGUCCAAG	1	1	transposon	LTR	ninja-Dsim-like
2319	AUCGUCUCGGGGAUGAAAACUCAACUGUGAGGGGAUCCUCAA	1	no match			
2320	CGGGGCCCGAAAACACGAGAUCCGA	1	1	transposon	LTR	opus
2321	UCUCAUCGUCGUAGAACAAGCCCGAGCUGUAACGUCGGGUCAAUA	1	no match			
2322	CACAAGGUGACCCCAAGCUGCUAU	1	1	transposon	LINE	G2

2323	UCCGAGGAACCCACUGAUCCAGCUA	1	1	Su(Ste)		Su(Ste)
2324	AAUCCUCAACUGUGAGGGAUCCUCAACUGUAACUCGGGGUCAAU	1	no match			
2325	UUGACCUCAAAUCAGGUAGGAGUACCCG	1	no match			
2326	UGUUUUCAUCUUUUAGACGGCUCGGGC	1	no match			
2327	GACACAGAGGGGAGCAGAUGAGAGCA	1	no match			
2328	GAUCUUCACCGUUAACUAGCCCCA	1	1	Su(Ste)		Su(Ste)
2329	ACUCUCAUCGUCGUAGAACAACGCCCG	1	no match			
2330	UCCUCAUCAUCGUAGAACAACGCCCG	1	no match			
2331	AUGGUUGCAACGGGUAACGGGAAU	1	2	ncRNA		rRNA
2332	GCUACGAGGAUGGAAGCGGCGAUGCC	1	0	coding gene		CG14718-RA|CG14718
2333	UCAAAGCGUUGACUACUCCACGAGGG	1	1	transposon	LTR	opus
2334	UAAGAUCUUCACCGUUGACUUGGCCCA	1	2	Su(Ste)		Su(Ste)
2335	UAGAUCUUCACCGUUGACUACCCCCAC	1	1	Su(Ste)		Su(Ste)
2336	AUCGUCUCGGGGAUGAAAAUCUCAUCGCCGUAGAACAAGCCCGA	1	no match			
2337	CAGAUCUUCACCGUUGACUUGGCCCA	1	1	Su(Ste)		Su(Ste)
2338	AUCUCAUCGUCGUAGAACAAGCCCCGACUGUAAACUCGGGGUCAA	1	no match			
2339	UCUCAUCGUCGUAGAACAAGCUCGAGCUGUAACUGGGUCAAU	1	no match			
2340	GCCCCGUACUCUGUGGUGACCAACU	1	1	transposon	LINE	baggins
2341	CAGGCACACAGAGGAGAGCGAGAUAGAGCA	1	no match			
2342	CUCUCAUCGUCGUAGAACAAGCCG	1	2	Su(Ste)		Su(Ste)
2343	CAGCACACAGAGGACGAGCAGAUGAGAGCG	1	no match			
2344	UCAAGGUGUAACUAUGACGAUUUGA	1	no match			
2345	UAAGGUGACCGGCAAAGCUAAAAU	1	0	transposon	LINE	baggins
2346	CUCAACUGUGAGGGGACCUCAA	1	no match			
2347	UAAGCCGAAGUAGUCAGUGGUAGUA	1	1	transposon	LTR	diver2
2348	UGUUUCAUCGUUAGACGGCUCGGGCCUU	1	no match			
2349	GCCCCGUACUCUGUAGUGACCAACU	1	2	transposon	LINE	baggins
2350	GGAUCUCAACUGUGAGGGGACCUCAA	1	no match			
2351	AUCGUCUCGGGAAUGAAACGUUGACCAAUUUCCUAACAUACGC	1	no match			
2352	AUAAGAAAACCACUCGAGCCGCCUUUA	1	no match			
2353	CCUCAUCGUCGUAGAAGAAGCCCGA	1	2	Su(Ste)		Su(Ste)
2354	UGUUUCAUCGCUAGACGGCUCGGG	1	1	AT-chX		AT-chX-B
2355	UGUUUCAUCUUUGACGGCUCGGGC	1	no match			
2356	AUCGUCUCGGGACUGAAACCUCUCAUCCGUUCGUACGAAACAACC	1	no match			
2357	GACGACCACGUUGUAAUGCCACCGAGC	1	0	transposon	LTR	ninja-Dsim-like
2358	UGAUCUUCACCGUUGACUAGCCCCA	1	1	Su(Ste)		Su(Ste)
2359	UCGAACUGUGAGGGGAUCCUCAACG	1	no match			
2360	CUCUCAUCGGUCGUAGAACUAAGCCCCGACUGUAACUCGGGGUCA	1	no match			
2361	CGAAAUGGACUUUUUUCCGAUGAGCCAGU	1	no match			
2362	UGAAUUGUCCGAGGAUUUGGCCAAG	1	0	transposon	LTR	invader1
2363	ACUGAGAAGAUAUUGUACCCGUGAA	1	0	coding gene		pseudogene
2364	CGGAGCGAUGUUCGGGCCCAGCUU	1	0	repeat		trf
2365	UUCAAGGUUACCCCAGGAUGCUUU	1	1	repeat		trf
2366	GAAAACCCAAACAAGCCCGGCAAGG	1	0	transposon	LTR	ninja-Dsim-like
2367	UGUUUCAACGUUAGACGGCUCGGGC	1	1	AT-chX		AT-chX-B
2368	UCUCAUCGUUCGUAGAACAAGCCCGACU	1	no match			
2369	UCUCAUCGUCGUAGGACAAGCCCGCUGUAACGUCGGGUCAAU	1	no match			
2370	GUUGACUUGACUCUACCACAUGACCUGU	1	0	transposon	LTR	micropia
2371	AUCCUCAAUUGUGAGGGGAUCCUCAA	1	no match			
2372	UAAAUGCCAGAAUUUUUAUUGGGGGUUC	1	no match			
2373	UCCGCUAGCUCAUUUCCCGGGAUGU	1	0	transposon	LINE	Fw2
2374	CGCACACAGAGGAGAGCAGAUGAGAG	1	no match			
2375	UUACCGAGGGCCGAGUACAUAUCGAG	1	0	transposon	LTR	G4
2376	AAGGUGGAAUGAAGAAAGUACA	1	no match			
2377	UCUCAUCAUCGUAGAACAAGCCCGAUCUGUAACUCGGGGUCAAU	1	no match			
2378	GGACUUCCGAAAAGCCCGCAGAAU	1	no match			
2379	UCUCAUCGUCGUAGAACAAGCGAG	1	2	Su(Ste)		Su(Ste)
2380	AUCCUCAACUGUGAGGGGAUCCUCGA	1	no match			
2381	CGCACACAGACGGAGAGCAGAUGAGACGCUA	1	no match			
2382	AUCGUCUCGGGGAUGAAAAAAACCUGAAGAUGAUCCUGAGUC	1	no match			
2383	AUCGUCUCGGGACUGAAACUUCUCAUCGUCGUAGAACAACGCCCG	1	no match			
2384	AUCCAGUCUCCCGACUAUCAGAUACCCG	1	1	transposon	LINE	INE-1
2385	UUGGACCAAACACUUUGAACCAAGUACUGU	1	no match			
2386	CGAAAUCCGAGGAACCAAUUGAUCCAGCUG	1	no match			
2387	CUCAUCAUCGUAGAACAAGCCCGAU	1	0	Su(Ste)		Su(Ste)
2388	GCCAGAUGGAUGGUUGGCAGGGAU	1	0	transposon	LINE	G2
2389	GAAACUGUCUCGGGAUGAGAAAGAA	1	no match			
2390	AAUUCCGAUAACGAACGAGACUCA	1	0	ncRNA		rRNA
2391	UCUCAUCGUAACGUGAACAAGCCCG	1	no match			
2392	AAGAAUGUUGUACCCCAGCCGGA	1	2	coding gene		CG10841-RA|CG10841
2393	UCCCAGGCCCGUAUAACGUUAAUGAUU	1	1	ncRNA		rRNA
2394	UAACUUCGGGAUAAGGAUUGGCU	1	0	ncRNA		rRNA
2395	GAACCAGCGAGGAGAGGCCCAGAAG	1	0	coding gene		CG16932-RA|Eps-15
2396	AUCCUCAACUGUAAGGGGAUCCUCAA	1	no match			
2397	UAACAAGAUGCAUACGAUGUGAC	1	1	transposon	DNA	1360
2398	AAAAAACCUGAAAGAUGGACUCCUGUAGUCGC	1	no match			
2399	GUCGUCGGGAUGAAAACG	1	2	transposon	LINE	R2
2400	GGGAUCCUCAACUGUGAGGGGAUCCUCAA	1	no match			
2401	AAACAAAGCAUUGUGAUGGCCCUAG	1	0	ncRNA		rRNA
2402	CUCGAAUGAGUGCUUGAUAUGACCAG	1	0	Su(Ste)		Su(Ste)
2403	CUAACUCCGCAUCGAUGACCGCCGC	1	no match			
2404	CCGCGGAAAGACGGAAAGACCCCAUG	1	no match			
2405	UAGAUCUUUACCGUUGACUUGCCCCG	1	1	Su(Ste)		Su(Ste)

2406	CGUUGGACCAAUUUCCGAAACAUACGCCUG	1	no match			
2407	UGAUGACCGAAAUUUGGAAAAACAGAC	1	0	repeat		ARS406_DM
2408	GGAUCCUCAACUGUGGGUCAA	1	no match			
2409	GACUAGCUGGCCGAGUAUUUGAUG	1	0	Su(Ste)		Su(Ste)
2410	GUCUCAUCGUCGUAGAACAAGCCCGU	1	2	Su(Ste)		Su(Ste)
2411	CUCAACUGUGAGGGGUCCUCAA	1	no match			
2412	GGCCGAGGAUCCGAUUUUCUUUG	1	0	Su(Ste)		Su(Ste)
2413	CCUCAUCGUCGUAGAACAAAGCCCGA	1	no match			
2414	UUAAGGCCGAGAGCUGAUGACGAGUG	1	no match			
2415	AGGGACUAGCUGGCCGAGUAUUUGAUAUUC	1	2	Su(Ste)		Su(Ste)
2416	UUCUCAUCGUCGUAGAACAAGCCUGA	1	1	Su(Ste)		Su(Ste)
2417	UAACAAGAUGCAUACGAUGUGGCA	1	1	transposon	DNA	1360
2418	UGUUAAGUGUCGGGAUAGGAGUGCC	1	0	transposon	LINE	Doc
2419	UCUCAUCGUCGUAGAACAAGCCCGAGA	1	1	Su(Ste)		Su(Ste)
2420	CUACCUGAAACUCUGUUGCACCCGA	1	0	transposon	LTR	accord
2421	CUAAGAUUGGACCCAGUACACUCGCCUU	1	no match			
2422	UUUCUCAUCGUCGUAGAAGAAGCCCG	1	1	Su(Ste)		Su(Ste)
2423	UCUCAUCGUCGUAGAACAAGCGA	1	2	Su(Ste)		Su(Ste)
2424	CCAACCUAGAAGUACUGCUGUCACAG	1	0	transposon	LTR	ninja-Dsim-like
2425	UCCUUAGUUCAUUCCUGAGUGCGAGCUCCA	1	no match			
2426	UGGUCUCAUCGUUAGACGGCUCGGGC	1	2	no annotation		
2427	UCAAACUGUGAGGGGAUCCUCAA	1	no match			
2428	GUGUUUUCAUCGUUAGACGGCUCGGGC	1	no match			
2429	UAGGAAAAACCUAGUUACUUACGGGC	1	0	Su(Ste)		Su(Ste)
2430	UACUCGAAUGAUUGCUUGAUAUGACCA	1	1	Su(Ste)		Su(Ste)
2431	AAUGCCGGAAGCGAUCCACUGCUU	1	no match			
2432	ACCAAGAUCAUCAACUCGUGUUAUUC	1	0	transposon	LINE	Cr1a
2433	UAUAUCUUCACCGUUGACUUGCCCC	1	1	Su(Ste)		Su(Ste)
2434	UCUCAUCGUCGUAGAACAAGCCCAAG	1	1	Su(Ste)		Su(Ste)
2435	UCUAGGCUGAGCGUAGAACCAGAGU	1	1	transposon	Low_complexity	F-element
2436	UAGAUCUUCAGCGUUGACUAGCCCCA	1	1	Su(Ste)		Su(Ste)
2437	UAAACAAAGAUGCAUACGAUGUGGC	1	no match			
2438	ACUGAAGGAGUACGAAUGAUCG	1	0	transposon	LINE	R1-element
2439	UCUUCUCGACGUUUACCAGCCUCCU	1	1	transposon	LTR	ninja-Dsim-like
2440	CCUCAUCGUCGUAGAACUAGCCCGA	1	1	Su(Ste)		Su(Ste)
2441	AAGGUGGAAAUGGAUCCUCAA	1	no match			
2442	UAACAAGAUGCCUACGAUGUGGC	1	1	transposon	DNA	1360
2443	GGUGCCCGAAGGUGUUGACCAAGUU	1	0	coding gene		CG33340-RA|CG33340
2444	CCGAACUUGAGACCGCCCCUGGACA	1	0	Su(Ste)		Su(Ste)
2445	UGUUUCAUCGUUAGACGGCUCGGCCGG	1	1	AT-chX		AT-chX-B
2446	ACAAGAUAUGACCAAAGCAAAGACGC	1	1	transposon	DNA	1360
2447	UCUCAUCGUCGUAGAAGAAACCCGA	1	2	Su(Ste)		Su(Ste)
2448	UAGAUCUCACCGUUGACUUGGCCC	1	no match			
2449	CGUUUACCAAUUUCCGAACAUACGCC	1	1	transposon	LTR	DM88
2450	GUGUUUCAUCGUUAGACGGCUCGGGGU	1	no match			
2451	AAGCCCGAUGUUGAUUAAGAAGAAUUA	1	no match			
2452	AGGAAAUGUUUCAUCGUUAGACGGCGUCGGGCCG	1	no match			
2453	GGCAACAUUUCGAGCCCCCAGAACAAC	1	no match			
2454	CAGCACACAGAGGAGAGCAGAUGAGAGCU	1	no match			
2455	UGGAGGGAUCCUCAA	1	1	coding gene		CG1507-RA|Pur-alpha
2456	UUGGAACGAAAUUGGCCUGAUUAGC	1	0	transposon	LTR	GATE
2457	AAUCUAGGCUGAGCGGAGAACCAGAG	1	1	transposon	Low_complexity	F-element
2458	AGUAUACCGAGGCGUUCGGGUGAA	1	no match			
2459	UCGAGAGCAUAUAAUGCCCCGAAGGU	1	0	transposon	LINE	R1-element
2460	UGUUUCAUCGUUACGACGGCUCGGGC	1	no match			
2461	UCGAUGGAGCUCACUUACCUAAUGUA	1	1	transposon	LINE	R1-element
2462	GAAGUUCAAGCCCGGGUAUUCUGA	1	0	repeat		trf
2463	GGAUCCUCAACUGUGAGGAGAUCCUCAA	1	no match			
2464	UGAUUCCGAGGAACCAAUUGAUCC	1	0	Su(Ste)		Su(Ste)
2465	CAGCACACAGAGGAGAGCGGAUGAGAGCA	1	no match			
2466	UCUGCCCGAUGAACAAAAAAUGGCCCGU	1	no match			
2467	AACUGUGGGGAUCCUCAA	1	2	no annotation		
2468	UCUCAUCGUCGUAGAACAAGCACGAG	1	1	Su(Ste)		Su(Ste)
2469	UGCUCUCGAAUGUAUGCCCGAUAAA	1	1	transposon	LINE	R1-element
2470	AUCUAGGCUGAGCGGAGAACCAGAG	1	0	transposon	Low_complexity	F-element
2471	UAACUCCGAGGGAGGGUGAAACCAUC	1	0	transposon	LINE	R1-element
2472	CGAAAUCUCAUCGUCGUAGACAAGCG	1	no match			
2473	UGUUUACAUCGUUAGACGGCUCGGGC	1	no match			
2474	UGUCUCAUCGUCGUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
2475	UUGGACCGACACUUUGAACCAAG	1	no match			
2476	UUAUAAUGAUGAUGAUACCGACUAAAG	1	1	no annotation		
2477	AUCUUAUCGUCGUAGAACAAGCCCGA	1	2	Su(Ste)		Su(Ste)
2478	UUAAACAAUUGAAAGAUACGACCCA	1	1	transposon	LINE	Cr1a
2479	UAAUCUUCACCGUUGACUAGCCC	1	2	Su(Ste)		Su(Ste)
2480	UCUCAUCGUCGUAGAACAAGCCCGACUGUAACUCGGGGUCAAUAU	1	no match			
2481	GGGAUCCUCACUGUGAGGGAUCCUCAA	1	no match			
2482	AUCGUCUCGGGAGUGAAAUUACUCAUCAUCGUAGAACAAGCCCGA	1	no match			
2483	UCUCAUCGGCGUAGAACAAGCCCCGA	1	no match			
2484	UGAUACCGAGAAUGACAAUGGCUCC	1	2	transposon	LTR	gypsy3
2485	UGUUUCAUCGUUAGACGGCUCGGGCCUGUAACUCGGGGUCAAUAU	1	no match			
2486	UGGACCUUUUCCCGAUGAGCCAGUG	1	0	transposon	LINE	G
2487	CAAGAUAUGACCAAGGUUCAAGAU	1	no match			
2488	GAUCUCAACUGUGAGGAUC	1	no match			

2489	UAAGGAAAUGACCCAGUUGGCCAGA	1	0	transposon	LTR	Stalker
2490	AUCUCAUCGUCGUAGAACAAGCCCCGACUC	1	no match			
2491	UGUACUACCCCUUCAUGACUCAA	1	0	transposon	LINE	Fw2
2492	UCGAACCCAGAGGCUGGAUCCUCAA	1	no match			
2493	GCGCACCGACGCCCGGACCAGACCUU	1	no match			
2494	UCUUAUCGUCGUAGAACAAGCCCG	1	1	Su(Ste)		Su(Ste)
2495	UUAAUGCCGGAAGCGAUCCCACUCGC	1	2	AT-chX		AT-chX-B
2496	UCGUCUCGGGAGUGAAAUACUCAUCGUCGUAGAACAAGCCCGC	1	no match			
2497	CAGCACACAGACGGAGAGCAGUAUGAGACGCA	1	no match			
2498	AUGAUACCGAGAAUGACAAUGGUUGAU	1	1	transposon	LTR	gypsy3
2499	CUGUUCGGAAUUCUGUAACCUCGGGUCAAU	1	no match			
2500	CCUCAUCGUCGUCGAACAAGCCCGA	1	2	Su(Ste)		Su(Ste)
2501	UUAAGUCCCGUUACUUGUGCCCAGUU	1	2	transposon	LINE	G6
2502	UCUCAUCGUCGUAGAACUAGCCGA	1	2	Su(Ste)		Su(Ste)
2503	GGAUCUCAACUGUGAGGGAUCCUCAA	1	no match			
2504	UACUGACCCAGUCGAAUCUAUUGUAC	1	1	transposon	LINE	baggins
2505	AUCGUCUCGGGGAUGAAAUCUCAUCGUCGUAGAACAAGCCCGAA	1	no match			
2506	CCUGCUCUCCCGCUCUAUUGGCUCC	1	0	transposon	Low_complexity	F-element
2507	ACAUCCUUCCGGGAGGUAGAGCA	1	no match			
2508	AUAACAAGAUGCAUGCGAUGUGGC	1	1	transposon	DNA	1360
2509	AUCCUCAACUGUGAGGCAUCCUCAA	1	no match			
2510	UUUACAGUGGCGCCGUCAGUUACUGA	1	0	transposon	LINE	R1-element
2511	UCUCAUCGUCGUAGAGCGAACGCCCGA	1	no match			
2512	UGUUUCAUCGUAGAACAAGCCCGAU	1	no match			
2513	GAGGCCGAGCUGUGACCAUCGAG	1	0	transposon	LINE	R1-element
2514	GCCAUCCUGACUAGCUGAUCCCCUGA	1	0	transposon	LTR	flea
2515	GGCAAUGGUGACCCGAAACUUUAUCU	1	0	coding gene		CG9064-RA|Ucp4C
2516	UAGAAGAACAUUCAAGCGCUUGACU	1	1	Su(Ste)		Su(Ste)
2517	UCUCAUCGUUGUAGAAACAAGCCCGA	1	no match			
2518	UCUCAUCGUGUAGAACAAGCCCG	1	no match			
2519	UCUCAUCGUCGUGAACAAGCCCGAG	1	no match			
2520	UGAAAUGUUUCAUCGUUAGACGGCUCGGGC	1	no match			
2521	UCAUUUCAAUCUAAGACUGGAGUGCC	1	1	no annotation		
2522	UCGUCUCGGAUAGAAAAA	1	2	no annotation		
2523	UGCAACUGAUGACCUGAUGACUCUU	1	no match			
2524	GUAGAUCUUCACCGUUGACUAGCCCCUACUGUAACCUCGGGUCAA	1	no match			
2525	AGUUGAGGACUGUAACUCGG	1	no match			
2526	AUCUCAUCGUCGUGAACAAGCCCCGA	1	no match			
2527	UCUCAUCGUCGUAGAACAAGCGCGA	1	1	Su(Ste)		Su(Ste)
2528	UUCUCAUCGUCGUAUAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
2529	AACAGGAGAAAUGAAGAAAGUACA	1	no match			
2530	UCCGCAUGUGAAUGUUGGCCCAGA	1	1	no annotation		
2531	UGUUUCAUAGUUAGACGGCUCGGG	1	1	AT-chX		AT-chX-B
2532	CAUUUGACAGAGUGUGGCUCGAUG	1	0	transposon	LINE	Doc
2533	CCGGAUGAAGAUUUGGACUACCUCCCC	1	no match			
2534	UAUGACCAGCAUGGACCUAACACUU	1	0	Su(Ste)		Su(Ste)
2535	UAAAAAAGCACCGUUUCCCGUCCGAUCAA	1	no match			
2536	AUCUCAUCGUCGUAGGACAAGCCCGA	1	2	Su(Ste)		Su(Ste)
2537	UGUAAAAUUACAAGACAAACCUGA	1	0	no annotation		
2538	AGAAAAAAAAAAAAAAUGAUCCUCAA	1	no match			
2539	AUCGUCUCGGGGAUGAAAUCUCAUCGUCGUAGAACAACGCCCGA	1	no match			
2540	AUCCUCAACUGUGGAGGGGAUCCUCAA	1	no match			
2541	CAAGAUAUGACCAAAGCAGACAC	1	1	transposon	DNA	1360
2542	AUUCAUCGUCGUAGAACAAGCCCAA	1	2	no annotation		
2543	UGUUUCAUCCGUUAGACGGCUCGGGC	1	no match			
2544	AUAAGAUCAACGAGAGAUGGCCAUAAA	1	2	transposon	LTR	opus
2545	CGUUGACCAAUUUCCGAACAUACAC	1	1	transposon	LTR	DM88
2546	UCUCAUCGUCGAAGAACAAGCCCG	1	1	Su(Ste)		Su(Ste)
2547	UUACUGACGAGAACUCGAGCCGCGUA	1	1	no annotation		
2548	CACACACAGAGGAGAGCAGAUGAAGCA	1	no match			
2549	UUUCUCAUCCUCGUAGAACAAGCCCG	1	1	Su(Ste)		Su(Ste)
2550	ACUGGUGAGGGGAUCCUCAACUGUAACGUCGGGUCGAAU	1	no match			
2551	AUGUUAAGGACGAGAAUCUACCCCCUGUAACGUCGGGUCAAU	1	no match			
2552	UGUUUCAUCGUUAGACGGCUCGGGCGU	1	1	AT-chX		AT-chX-B
2553	GUGUUUCAUCGUUAGACGGCACGGGC	1	2	AT-chX		AT-chX-B
2554	GGCCUCAACUGUGAGGGGAUCCUCAA	1	no match			
2555	UAGAUCCUCACCGUUGACUUGGCCC	1	1	Su(Ste)		Su(Ste)
2556	UCUCAUCGUCGUAGGGAAGCCCGA	1	no match			
2557	AAUUCUCAUCGUCGUAGAACAAGCCCGA	1	2	Su(Ste)		Su(Ste)
2558	GGAGAGGUGGCCGAGUGGUUAAUGG	1	no match			
2559	UCGGGAACACCCCGUGUUGUUGGCCU	1	2	ncRNA		5SrRNA
2560	AGUGUAUUUCCUAGACCGAGAGCUG	1	no match			
2561	CUCAACUGUGAGGGGAUCCCUCAA	1	no match			
2562	UGUGUCAUCGUUAGACGGCUCGGG	1	1	AT-chX		AT-chX-B
2563	GAAUAUCGUCUCGGGAGUGAAAGUACUCGGGAUGAUAGGAUCCUC	1	no match			
2564	UCUCAUCGUCGUAGAACAAAGCCCGAC	1	no match			
2565	UGUUUUCAUCGUUAGACGGCUCGGGGAUGAAAUCUCAUCUUCGUA	1	no match			
2566	AUCGUCUCGGGAGUGAAAUAAGAGAGUGAUGCCGAGUAUAAAG	1	no match			
2567	UCUCAUCGUCGGAGAACAAGCCCGAAG	1	1	Su(Ste)		Su(Ste)
2568	AUCGUCAGUCGUACGGUAGAACAAGCCCGGA	1	no match			
2569	AUGUUUCAUCGUUAGACGGCUCGGGGCC	1	no match			
2570	UACAAUCUCCGUGCACGACCCCGAUCU	1	0	transposon	LINE	Circe
2571	UAUGCAACCCUUAUAAGCUGCACCCGA	1	0	transposon	LTR	gypsy12

2572	AAUAACCCGAUUCCUCUUCCAUGAGA	1	1	transposon	LINE	Doc4
2573	UCGCAUCGUCCUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
2574	UUGAUACGAAUGAGACCCGCAGUAAA	1	0	repeat		trf
2575	AAAAAUCUCAUCGUCGUAGAACAAGCCCGA	1	no match			
2576	CUCAACUGUGAGGGGAUCCCAA	1	no match			
2577	CCGAUGAACAAAAAUGGCCCGUAAAG	1	no match			
2578	GCCCGUACUCUGUGGUGACCAACAC	1	1	transposon	LINE	baggins
2579	GCUGCCGAGAUAUGCUGCGAGUUGCCAACUC	1	no match			
2580	AUCGUCUCGGGAGUGAAAUACUCAUCAUCGUAGAACAAGCCCGA	1	no match			
2581	UCUCAUCGUCGUAUAACAAGCCCGAGG	1	1	Su(Ste)		Su(Ste)
2582	CUGUUUCAUCGUUAGGCGGCUCGGG	1	1	AT-chX		AT-chX-B
2583	UCGGCCCGAACCCUUCUGGUGUGUU	1	0	transposon	LINE	baggins
2584	GCAUAUCAUUCACGCCACGCUCCGAC	1	0	Su(Ste)		Su(Ste)
2585	GCACACAGAGGAGAGCAGAUGGAGCA	1	no match			
2586	AACUGUGGGUUCAAAUCGAACCCAGGGAA	1	no match			
2587	AUCGUCUCGGGAGUGAAAUACUCAUCGUCGUAGAACUAGCCCGA	1	no match			
2588	AUCGUCAUCGUCGUAGAAGAAGCCCGA	1	no match			
2589	AUCGUCUCGGGGAUGAAAAUGUUUCAUCGUUAUACGGCUCGGGC	1	no match			
2590	UCUCAUCGUCGCAGAACAACGCCCGA	1	no match			
2591	AGAAAUCUCAUCUCGUAGAACAAGCCCGA	1	no match			
2592	AUCUCAUCGUCGUAGAAGAAGCCCGU	1	no match			
2593	GAUCUUCACCGUUGACUUGGCCCAC	1	0	Su(Ste)		Su(Ste)
2594	ACAACUGUGAGUCCUCAA	1	2	coding gene		CG7963-RA|CG7963
2595	UUCGUGGAGUGAUUUGUCUGGUUA	1	0	ncRNA		rRNA
2596	UAAUAAAAACAACGACCCGCUACCGGAG	1	no match			
2597	AUGAUUUUUUUCUUUGAAGAUGGCCCA	1	1	no annotation		
2598	CAGCACACAGAGGAGAAGAUGAGAGCA	1	no match			
2599	ACAGCACACAGAGGAAAGCAGAUGAGAGCA	1	no match			
2600	AUCCUCAACUGUGAGGGAUCCUCCAA	1	no match			
2601	GGAUUCAACUGUGAGGGAUCCUCAA	1	no match			
2602	UCCGGAAGAAUGGUUGCGCAUGGUUA	1	2	transposon	LINE	TART-A
2603	AUGUUUCAUCAUUAGACGGCUCGGGC	1	2	AT-chX		AT-chX-B
2604	UCCUCAACUGUGAGGGAUCCCAA	1	no match			
2605	UCUCAUCGUCGUAGAACAAGCCCGACUGUAACUCGGGGUCAAUAU	1	no match			
2606	AAAGAACCGAUUCCUCUUCCAUGAG	1	no match			
2607	UCUUCAUCGUCAGACGGCUCGGG	1	no match			
2608	AAAACCUGAAGAUGAUCCUGACGUCGC	1	no match			
2609	GAUAUCUCACAGUUGAGG	1	1	no annotation		
2610	UCAGAUGGUGCCUUUAGUUUCCCGAAG	1	0	transposon	DNA	1360
2611	CCUAACGUCGAGUUAAGGUGCCCAA	1	no match			
2612	CUCCAUGCAGCAGAAGCGACAGAGUU	1	0	transposon	LINE	baggins
2613	UCUCAUCGCCGUAGAACAAGCCCGAA	1	1	Su(Ste)		Su(Ste)
2614	AUCUCAUCUUCGUAGAACAAGCCCCGA	1	no match			
2615	ACAAUGAUCUGAUCGGGUUUUUCUG	1	0	transposon	LINE	Cr1a
2616	UUUGCCUGAGGGUGAUAUGCGGUUG	1	1	transposon	DNA	1360
2617	CAGCACACAGAGGAGAGCAGAUGAGAUCA	1	no match			
2618	CAGCACACAGAAGGAGAGCAGAUGAGAGCA	1	no match			
2619	AAAACCCUAGUUACUUACGGGCUCGA	1	1	Su(Ste)		Su(Ste)
2620	UCGAAUUGCCCUAACGGUGUAACAAA	1	0	transposon	LTR	flea
2621	UCAACUGUGAGGGGACCCUCAA	1	no match			
2622	AUCUCAUCGUCGUAGAACAAGCCCGC	1	2	Su(Ste)		Su(Ste)
2623	UUAAGAUCAACGAGAGAUGGCCA	1	0	transposon	LTR	opus
2624	UGUUUCAUCGUUAGACGUCUCGUGC	1	2	AT-chX		AT-chX-B
2625	AUCUAAUCGUCUUAGAACAAGCCCGA	1	no match			
2626	UCUCCAUCGUCGUAGAAGUAGCCCGA	1	no match			
2627	UAUUGCUUGUGGAACGUCGUUGCC	1	0	no annotation		
2628	UCAACUGUGAGAGGAUCCUCAA	1	no match			
2629	AUCGUCCUCGGGAUGAAACCGUCAACUGCGAGGGGAUCCUCAA	1	no match			
2630	UAAAAACCAAUCGAUGACAACAAAAAAUA	1	no match			
2631	UAGAUCUUCACCGUUGACCUUGCCCC	1	no match			
2632	AAAACCUAGAAGAUGAUCCUGAGUCG	1	no match			
2633	UCCGGUAACGAGCGAGACCCCAACG	1	no match			
2634	CUUAUCACCCCGGAAUUGGUUUAU	1	no match			
2635	AGGCAAGGGAAGUCGGCAAAAUGGAU	1	no match			
2636	UCAACUGUGAGGGUAUCCUCAA	1	no match			
2637	UGCACGAGAUGCUUGGACCGCACAUGC	1	0	coding gene		CG8103-RA|Mi-2
2638	UGUUUCAUCGUUAGACGUCUCGGC	1	2	AT-chX		AT-chX-B
2639	GCUGAGCUGGAUUAGACCUCGCGC	1	0	transposon	Low_complexity	F-element
2640	CAGCACACAGAGGAAGCAGAUGGAGC	1	no match			
2641	UCUCAUCGUCGGUAGAACAAGCCCGGA	1	no match			
2642	UGUUUCAUCGUUAGACGGCUCGGGCCUGUAACUCGGGGUCAAUAU	1	no match			
2643	UUCAAGGUUAUCCCAGGAUGCUGU	1	1	repeat		trf
2644	UUCUUUGGUGAGGACUUGAGCGA	1	1	Su(Ste)		Su(Ste)
2645	UCUCAUCGUCGUAGGAACAAGCCCCGA	1	no match			
2646	GUGGAAAUGAAGAAAGUA	1	1	no annotation		
2647	UACUCAUCGUCGUAGAACGAGCCCGA	1	2	Su(Ste)		Su(Ste)
2648	CAGCACACAGAGGAGAGCCAGAUGAGGAGCA	1	no match			
2649	UCAUCUUCUUCUCGACGUUGACCAGCCU	1	0	transposon	LTR	ninja-Dsim-like
2650	UCCGAACGUACCACGGUGCCCAGU	1	no match			
2651	CCGCAUAGCUAGAGAUGACAUACCGGC	1	0	no annotation		
2652	UGUUUCAUCUUUAGACGGACUCGGGC	1	no match			
2653	CGUGGCAGAGGAACUCGUUGACCUUGA	1	0	Su(Ste)		Su(Ste)
2654	UCUCAUCGUUGUAGAGCAAGCCCGA	1	2	Su(Ste)		Su(Ste)

2655	AUCUCAUCUUCGUAGAACACCCCGA	1	no match			
2656	AUGAGCCGACGAGAUGCCCAGGGCGUG	1	1	transposon	LTR	ninja-Dsim-like
2657	GUAGAUCUUCACCGUUGACUUGCCCCCCUUGUAACUCCGGGUCAA	1	no match			
2658	GAUCUCAUCGUCGUAGAACAAGCCCG	1	2	Su(Ste)		Su(Ste)
2659	AUCGUCAUCGUCGUAGAACAAAGCCCGGA	1	no match			
2660	UCCUCAUCGUCGUAGAACCAAGCCCCGAGU	1	no match			
2661	UUCUCAUCGUAGUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
2662	UGGUUUCAUCGUUAGACGGCUCGGGUCG	1	no match			
2663	UCUGCCCGAUGACAAAAAUGGCCCGUA	1	0	transposon	LTR	mdg1
2664	UCUCAUCCGUAGAACAAGCCCGA	1	no match			
2665	CAGCUACACAGAGGAGAGCAGAUGAGAGCA	1	no match			
2666	AUGAUGACCGAAAUUUGGAAAACGGA	1	1	repeat		SAR_DM
2667	UGACCAUACAAGAGUACAAGGCACG	1	0	transposon	LTR	blood
2668	UGACACGCGGGAGACCGGGGUU	1	1	ncRNA		tRNA
2669	AAAAAAAAGGAUCCUCAA	1	1	no annotation		
2670	UCUCAUCUUCGCAGAACAAGCCCGA	1	2	Su(Ste)		Su(Ste)
2671	UCUCAUCGUCGUUGAACAAAGCCCCGA	1	no match			
2672	UGUGACUGAUCUAGAGUACAUGU	1	no match			
2673	UCUCAUCUUCGUAGAACAAACCCGA	1	2	Su(Ste)		Su(Ste)
2674	GGGUGAGGAGGAUCCUCAA	1	no match			
2675	ACAGGCCCAUCUGGUUGAACGUGUC	1	0	Su(Ste)		Su(Ste)
2676	GAGGACUUGGGCGAUUUAAGUGG	1	0	Su(Ste)		Su(Ste)
2677	UUCUUCUCGACGUUGACCAGCCUAA	1	2	transposon	LTR	ninja-Dsim-like
2678	ACAAGAUAUGACCAAAGCAAAUACCAC	1	no match			
2679	UCCAGGCCCGUAUAACGUUAAUGAU	1	1	ncRNA		rRNA
2680	UACUUACAGUGGUUGUCCAGGG	1	no match			
2681	UCUCAUCGUCGUGGAACAAGCCCGAGGCUGUAACUCGGG	1	no match			
2682	UGCCUCAAGAUAUGACCAAAGCAAA	1	0	transposon	DNA	1360
2683	AUCAAAGCGUUGACUACUUCACGAGGAA	1	2	transposon	LTR	opus
2684	AUAAACUGUGAGGGGAGUCCUCAA	1	no match			
2685	UGUUUCGUUAGACGGCUCGGGC	1	no match			
2686	UGCUCUCGAAUGUAUGCCGAUA	1	no match			
2687	UUUUGUAUGCGACGAGUACGCUAU	1	0	transposon	LINE	Rt1b
2688	GGAAAACCUGAAGAUGAUCCUGAGU	1	0	transposon	LINE	Helena
2689	CAACCAGAGAUCUAGAGGGCCCAA	1	0	no annotation		
2690	GAGAUCUUCACCGUUGACUAGCCCCA	1	1	Su(Ste)		Su(Ste)
2691	AACCUGUGAGGGGAUCCUCAA	1	no match			
2692	AUUCGUCAUCGUCGUAGAAGAAGCCCGG	1	no match			
2693	ACUACUUCCAUCCUAAAACUGAUAC	1	0	Su(Ste)		Su(Ste)
2694	UUCAGGAACCUAUUUCGGGCUGAA	1	0	transposon	DNA	invader2
2695	UGUCACGCGGGAGACCGGGGUUCGA	1	1	ncRNA		tRNA
2696	CUGUAACUCGGGCCUGUAACUCGGGC	1	no match			
2697	CUCCUCAACUGUGAGGGGAUCCUCAA	1	no match			
2698	UAGAUCUUCGCCGUUGACUAGCCCCA	1	1	Su(Ste)		Su(Ste)
2699	UACAUUCGAGAGCAUAUAACCCCCG	1	no match			
2700	UUGACGAACACCCUUUUGCUUGCGCC	1	0	transposon	DNA	hopper
2701	UCUAUCUGAGCGGAGAACCAGAGU	1	no match			
2702	UAGAUAUUGAACAGCUCAGAGCGU	1	1	transposon	LTR	Stalker2
2703	UCGUCAGUCGUACGUAGAACAAGCCCGGA	1	no match			
2704	UACUCAUCGUCGUAGAACAAGCCCGAG	1	1	Su(Ste)		Su(Ste)
2705	CAGCACACAGAGGAGAGCAGAUGAAGA	1	no match			
2706	GGACCUUUUCCCGAUGAGCCAGU	1	0	transposon	LINE	G
2707	UCUCAUCGUCGUAAAACAAAGCCCGA	1	no match			
2708	AAAGAUAAGACCCAUAUUAAAAAA	1	1	transposon	LINE	Cr1a
2709	GAAACUUCACCUCAGUCGGAUUAGCC	1	0	ncRNA		pseudogene
2710	UCUCAUACCGUCGUAAGAAACAACGCCGA	1	no match			
2711	GAAUUACUUACCGAGGGUUGUCCAG	1	1	Su(Ste)		Su(Ste)
2712	AUCUCAUCGUCGUAGAACAAGCCCGCUGUAACUCGGUCAAU	1	no match			
2713	GGAUGAAAUCUCAUCGUCGUAGAACAAGCCCGA	1	no match			
2714	UCUCAUCGUCGUAGAACAAGCCCGACUGUAACUCGGUCAAU	1	no match			
2715	UUAAUGCCGGAAGCGAUCCCCACUGCU	1	no match			
2716	AUGAUCUUCACCGUUGACUUGGCCC	1	2	Su(Ste)		Su(Ste)
2717	AAAGAAUCCUGACAAGAUCCCGGGC	1	0	transposon	LTR	invader1
2718	ACCGGACGGGCCUAGAGUUACUUCCCC	1	2	coding gene		CG40635-RA|CG40635
2719	AUCGUCAUCGUCGUAGAAGAAGCCCGGA	1	no match			
2720	CGUUGACCAAUUUCCGAACAUACCC	1	1	transposon	LTR	DM88
2721	UUCUCACUCGUCGUAGAACAGCCCGA	1	no match			
2722	UGAUACCGAGAAUGACAUGGUUGA	1	no match			
2723	UACAUCUUCACCGUUGACUAGCCCC	1	1	Su(Ste)		Su(Ste)
2724	AUCUCAUCGUCGUAGAACAAGCCC	1	1	Su(Ste)		Su(Ste)
2725	UCCGGUCUCCUGAGUCGGACUCCGAG	1	0	transposon	LINE	R1-element
2726	CAAUAUAUGACCAAAGCAAAGACAC	1	0	transposon	DNA	1360
2727	AUCGUCAGUCGUACGUAGAACAAGCCCGGA	1	no match			
2728	UCCAACUGAAGGAGUACGAACGAUCG	1	1	transposon	LINE	R1-element
2729	AAACGAAAUGAAAAAUAACGGAUGA	1	0	transposon	LTR	GATE
2730	GGAUAUCUACGAGCUGUUUGAUCCUGUC	1	1	coding gene		CG6829-RB|Ark
2731	UAAGGAUGAUGAUACCGACUAAAGGCA	1	0	no annotation		
2732	AGAAAUGUUUCAUCGUUAGACGGCUCGGGUU	1	no match			
2733	GACCGAAUGAUUUAACCCGGAGACU	1	0	transposon	LTR	Rt1c
2734	GGAUCCUCAACUUGGAUCCUCAA	1	no match			
2735	GGCACACAGAGGAGAGCGAGAUGAGAGCA	1	no match			
2736	GGAUCUGUGAGGGGAUCCUCAA	1	no match			
2737	CCCCAUGUAGUGAUUGUAUCCCCAA	1	1	coding gene		CG3506-RA|vas

2738	UUGGACCGAACACUUUGAACCAAGA	1	1	Su(Ste)		Su(Ste)
2739	CAGCACACAGAGGAGAGCAGAUGAGAGCACUG	1	no match			
2740	UCAAAUCGUCGUAGAACAAGCCCGA	1	2	Su(Ste)		Su(Ste)
2741	AUGUUUCAUCGUUAGACGCCUCGGGCC	1	no match			
2742	UGUACUGUUUGUAGGUGUACUGGCCAA	1	1	transposon	LINE	baggins
2743	UCGUCAUCUUCGUAGAACAAGCCCGGA	1	no match			
2744	CAUGGAUUGCGCCGCCCUGCCCGC	1	no match			
2745	UCUCAUCGUCGUAGAACAAGCCCGACUGUAACUCGGGGU	1	no match			
2746	UCUCAUCGUCGUAGAACAUCCCG	1	no match			
2747	GGACCGAACACUUUGAACCAAGUA	1	0	Su(Ste)		Su(Ste)
2748	UGUUUUCAUCGUUAGACGGCUCGGUC	1	no match			
2749	UAAAAAGCACCGUUUCCCGUCCGAUCAA	1	no match			
2750	CAGAUACCCGUUACUCAGCUAGUGG	1	0	transposon	LINE	INE-1
2751	UGUCACGCGGGAGACCGGGGUUA	1	1	ncRNA		tRNA
2752	GAACACCGCGUGUUGUUGGCCCU	1	2	ncRNA		5SrRNA
2753	ACUGUAACUCGGGGUCAAUAGUUACAGU	1	no match			
2754	CGAAAUGGACUUUUUUCCGAUGAGCCCAGU	1	no match			
2755	AUCCUCAACUGUGAGGGGAAUCCUCAA	1	no match			
2756	GCCCGAACAUGGCUGUGUUAACUUU	1	no match			
2757	UUCCGAAGAGGAUGGAUACCGGCA	1	no match			
2758	GAAGAAUUCUUCCUCAGCGUUCCCGG	1	0	transposon	LINE	INE-1
2759	ACUACUUCCAUCCUAAAAACUGAUAC	1	no match			
2760	GAUGGAACCCCAGGAAGCUCACGCUCGUGUAACGUCGGGUCAAU	1	no match			
2761	UGUUUUCAUCGUUAGACGCUCGGGC	1	no match			
2762	AUGUUUUCAUCGUUAGACGGCUCGGGCC	1	no match			
2763	ACAAUAUCGUCUCGGGGAUGAAAUCUCAUCGUCGUAGAACAAGCC	1	no match			
2764	UCUCAUCAUAGAACAAGCCCGA	1	no match			
2765	AAACUGAAGGAGUACGAAUGAUCG	1	1	transposon	LINE	R1-element
2766	CAAGAUAUGACUAAAGCAAAGACAC	1	0	transposon	DNA	1360
2767	GGCUCCUCAACUGUGAGGGGAUCCUCAA	1	no match			
2768	CUGUGAGGUCUCCGGACGUGAUCA	1	2	ncRNA		rRNA
2769	UAUCAUUCACGCCACGCUCCGAC	1	0	Su(Ste)		Su(Ste)
2770	UCUGGAUACUAGGUAUCGACGCCGA	1	0	no annotation		
2771	CACCGGGUUUUGGCACUUUGCCGCG	1	0	transposon	LINE	R1
2772	UUAAUGCCGGAAGCGAUCCCCACUGC	1	no match			
2773	CAAGAUAUGACCUAAGCAAAGACACU	1	1	transposon	LINE	Cr1a
2774	CCCAAGAAACCCAAGCCUAAGGU	1	2	transposon	LTR	opus
2775	UCCUGUGAGGAUCCUCAA	1	2	transposon	LINE	Cr1a
2776	CUUAAUGCCGGAAGCGAUCCCACCUG	1	no match			
2777	UGUUUCAUCGUUAGACGGAUCGGGC	1	1	AT-chX		AT-chX-B
2778	UCUCAUCGUCGUAGAACAAGCCAGAG	1	1	Su(Ste)		Su(Ste)
2779	GAUCCAACUGUGAGGGAUCCUCAA	1	no match			
2780	CAGCACACAGACGGAGAGCAGUAUGAGACGCUA	1	no match			
2781	UCAAUACGACCCGGUUUAUUAAAAUC	1	0	transposon	LTR	mdg1
2782	CUCGGCCUCGGGCACAGUUCGCUUG	1	0	transposon	LINE	R1-element
2783	CUCUCAUCGUCGUAGAACAAGCCCGUA	1	2	Su(Ste)		Su(Ste)
2784	AUAAGGAUGAUGAUACCGAUUAAA	1	0	no annotation		
2785	GCUCUCGAAUGUACGCCCGAUAUACG	1	1	transposon	LINE	R1-element
2786	UGCAUUGCCCAACAAUACAUUCUC	1	0	transposon	DNA	1360
2787	CAGCAACAGAGGAGAGCAGAUGAGAGCUA	1	no match			
2788	GGAUACGGAGACCGAGGUUCAAACCGAG	1	no match			
2789	UGUCUUGAGAACCACCAGAUUGAUAGCC	1	2	ncRNA		pseudogene
2790	CGUUUCAUCGUUAGACGGCUCGUGC	1	2	AT-chX		AT-chX-B
2791	UCCUAGAAAAAAACGAGGUUCACCCAAU	1	0	transposon	LINE	F
2792	UUAGUAAAUGUAAGUCACGCCCUGAGG	1	no match			
2793	AAAAAGGAUCCUCAA	1	1	transposon	LTR	HMS-Beagle
2794	UGUUUCAUUUAGACGGCUCGGGC	1	no match			
2795	UCUCAUCAUCGUAGAACAAGCCCGG	1	1	Su(Ste)		Su(Ste)
2796	UGUUUAUCGUUAGACGGCUCGGGC	1	no match			
2797	UCCUAAGAUGAUGUGACCCAGAUGCU	1	no match			
2798	UACAUCAAGCUUCGCCAUCGCCCGGU	1	0	transposon	LINE	R1-element
2799	UAGAUCUUCACCGUUGACUAGCCCCCACUGUAACUCGGGGUCAAU	1	no match			
2800	UCUCAUCGUCGUAGAACAAGCCCCGACUGUAACUCGGGGUCAAUA	1	no match			
2801	UCAUCUUCUUCUCGACGUUGACAG	1	2	transposon	LTR	ninja-Dsim-like
2802	UGUUUUCAUCGUUAGACGGCUCGAGGC	1	no match			
2803	UCUCAGUCGUCGUAGAACAAGCCCGACUG	1	no match			
2804	UCUCAUCGUCGUAGAGGAAGCCCGA	1	2	Su(Ste)		Su(Ste)
2805	UGUGAGGGGACUCCUCAA	1	2	transposon	LTR	3S18
2806	CUCUCAUCGUCGUAGAAUAACGCCCGA	1	no match			
2807	AUCGUCCUCGGGAUGAAAUCUCAUCGUCGUAGAACGAAGCCCGGA	1	no match			
2808	UUUUGACAGGAAAAAUGUUCUGAGUU	1	0	transposon	LINE	G2
2809	UCUCAUCGGCGUAGAACAAGCCCAA	1	2	Su(Ste)		Su(Ste)
2810	CAAGACAUGACCUAAGCAAAGACAC	1	2	transposon	LINE	Cr1a
2811	UUCUGACGAGAACUCGAGCCGUAG	1	no match			
2812	UGUUUCAUCGUUAGACGGCCUCGGGC	1	no match			
2813	UCUCAUCGUUAGACGGCUCGG	1	1	AT-chX		AT-chX-B
2814	UUUAAGGCAACUCCCUGGACGCACGA	1	0	transposon	LTR	ninja-Dsim-like
2815	GCACACAGAGGAGAGCAGAUGAAAGCA	1	no match			
2816	UCCGAACGUACCCGUGCCCCAGU	1	no match			
2817	UCCAACGAGAGAGCGGAGAGCCAGC	1	0	transposon	Unknown	RR50960
2818	GUGUGACCCCUGGUACUGAAAUGCA	1	1	transposon	DNA	Bari1
2819	AUCUCAUGUCGUAGAACAAGCCCGA	1	no match			
2820	AGAAAUCAACUGUGAGGGGAUCCUCAA	1	no match			

2821	UCUACUGACUUCAGUAAGCCCCGU	1	0	transposon	LINE	baggins
2822	AUCGUCAUCGUCGUAGAACAAGCCCG	1	no match			
2823	GACUAGCUGGCCGAGUAUUUGUUA	1	1	Su(Ste)		Su(Ste)
2824	UAGAAAACAACCAGGACUAUAUUCGGA	1	0	Su(Ste)		Su(Ste)
2825	GAGAUAGCAGGACUGGGGGCGG	1	no match			
2826	UCGCAUCGUCGUAGGACAAGCCCGACUGUAACUCGGGGUCAAU	1	no match			
2827	UCCGAGGAACCAAUUUAUCCAGCUG	1	1	Su(Ste)		Su(Ste)
2828	GCGACACAGAGGAGAGCGAGAGAGAGCA	1	no match			
2829	UAUAUUCGGAGAAUUACUUACCGAGGG	1	1	Su(Ste)		Su(Ste)
2830	UCGAAAGGAUGAUGGAGAAGAUU	1	1	transposon	LINE	G2
2831	AUCGUCUCGGGGAUGAAAUCUCAUCGUCGUAGAACAAGCCCC	1	no match			
2832	GCUCGUCCUCGGGUUUGUUGUUGAU	1	0	transposon	LINE	G6
2833	CAAGUAAGCCCGAAGAUUGUCCAAU	1	1	transposon	LTR	opus
2834	AGUCUUCAUCGUCGUAGAACAAGCCCCGUACUGUA	1	no match			
2835	UGUCACGCGGGAGACCGGGUUCAA	1	no match			
2836	UCUCAUCGUCGUAGAACAAGCCCAACUGUAACGUCGGGUCAAU	1	no match			
2837	UACUCGAUCGUCGUAGAACAAGCCCGACGUGUAACGUCGGGUCAA	1	no match			
2838	UCUCAUCGUCCUAGAACUAGCCCGA	1	1	Su(Ste)		Su(Ste)
2839	CGAAAUCUCAUAGUCGUAGAACAAGCCCGA	1	no match			
2840	CGCGAAAGACCCAAGUUAGAUCCUAAAA	1	no match			
2841	UCUCAUCGUCGUAGAACAACGCCCGACUA	1	no match			
2842	UCUUAUCGUCGUAGAACAAGCCCUA	1	2	Su(Ste)		Su(Ste)
2843	GCGUCCGAAAACAAGGCCCGUCGCU	1	no match			
2844	CGAAAUGUUUCAUCGUUACGACGCCUCGGGC	1	no match			
2845	CUUCGAGAGCGAACCCGGAGAA	1	no match			
2846	CAGCACACAGAGGAGAGCAGAUGAGGCACUGUAACUCGGGGUCAA	1	no match			
2847	UCUCAUCGUCGUAGCACAAGCCCG	1	1	Su(Ste)		Su(Ste)
2848	UCCCGAUCUCCUGAGUCGGACUCCCGAGC	1	no match			
2849	UGUUUCUAUCGUUAGACGGCUCGGAC	1	no match			
2850	AUCUCAUCGUCCUAGAACAAGCCCG	1	2	Su(Ste)		Su(Ste)
2851	UUCACCUUUGACUUGGCCCACACAU	1	1	Su(Ste)		Su(Ste)
2852	CUCGAUGUCGGCUCGUCGCAUCCUGGG	1	no match			
2853	UCCAAUUAAAAAAUGACCUGGUGUU	1	0	transposon	LTR	roo
2854	AAAAGUUGACCGAACUUGAUUAUUUAG	1	0	ncRNA		rRNA
2855	UACGACCAAGUAGGACUAUUGUA	1	no match			
2856	CCGAACUUGAGACCGCCCCUGGAC	1	0	Su(Ste)		Su(Ste)
2857	AUCCUCAACUGUGAGGGAUCCGUCAA	1	no match			
2858	UAGAUCUUCACCGCUGACUAGCCCCG	1	2	Su(Ste)		Su(Ste)
2859	AACUUCGGGAUAAGGAUUGGCUCUC	1	1	ncRNA		rRNA
2860	GCACACAGAGGAGAGCAGACGAGAGCA	1	no match			
2861	AUCUCAUCGUCGUAGAACCAGCCCGAG	1	2	Su(Ste)		Su(Ste)
2862	AACUGCACCCUGAGUAUGGCAUUAUA	1	0	transposon	LINE	Rt1b
2863	CUCAACUGUGAGGGAUCCUCAACG	1	no match			
2864	GAAAAUCUACUACGUACGUCGUAGAACAAGACCCGA	1	no match			
2865	AGCACACGGAGGAGAGCAGAUGAGAGCA	1	no match			
2866	UGACUGUUCCCGUUGAUUCCCGUU	1	0	no annotation		
2867	UCUCAUCGUCGUAGAACACGCCCGA	1	1	Su(Ste)		Su(Ste)
2868	AUCGUCUCGGGGAUGAAAAUAAAUGCCAGAACUUUAUUGGGGUG	1	no match			
2869	CAGCACACAGAGGAGAGGCAGAUGAGAGA	1	no match			
2870	CUCUACUGUGAGGGGAUCCUCAA	1	no match			
2871	CUCGAUCGUCGUAGAACUAGCCCGAGG	1	no match			
2872	CAGCCCACAGAGGAGAGCAGAUGAGAGCA	1	no match			
2873	UCUCAUCGUCGUAGCAGAAGCCCGA	1	2	Su(Ste)		Su(Ste)
2874	UAGAUCUUCACCGUUGGACUUGGCCCCA	1	no match			
2875	AUCGUCGUCGGGAUAGAAAUGUUUCAUCGUUAGACGGCUCCGGGC	1	no match			
2876	CUCCGAACGUACCACGGUGCCCCAGU	1	0	transposon	Low_complexity	F-element
2877	UUCUCAUCGUCGUAGAAACAAGCCCGA	1	no match			
2878	AAAACCUGAAGAUAAUCCUGAGUCGC	1	1	transposon	LINE	Helena
2879	GUCAUCUUCUUCUCGACGUUGACCUA	1	no match			
2880	UCCGAGGAACCAGUUGACUCCAGCUC	1	no match			
2881	GAUCCAACUGUGAGGGGGAUCCUCAA	1	no match			
2882	UCAACCUGAGAGUUUGAUCCUG	1	no match			
2883	GCCCCGUCCUCUGUGGUGACCAACA	1	1	transposon	LINE	baggins
2884	UAUCCUCAACUGUGAGGGAUCCUCAA	1	no match			
2885	UAAGCCCGAAGAUUGUCCAAUGACCA	1	0	transposon	LTR	opus
2886	AAUACUGACCCAGUCGGAUCUAUUGUA	1	2	transposon	LINE	baggins
2887	AUCGUCUCGGAAUCGAAAUCUCAUCGUCGUAGAACAAGCCCGA	1	no match			
2888	UGAUAUGACCAGCAUGGACCUAACAC	1	0	Su(Ste)		Su(Ste)
2889	CAACCAGAGAUCUAGAGGGCCCAAGA	1	0	no annotation		
2890	UCUCAUCGUCGUAGAACAACCCCGACUGUA	1	no match			
2891	UCUCAUCGUCGUAGAACAAGCCCGAGACGAU	1	no match			
2892	UUCCCAUCGUCGUAGAACAAGCCCAA	1	2	Su(Ste)		Su(Ste)
2893	UGUAGUUAUAUUCAUAGAUGGCCCUA	1	0	transposon	LTR	HMS-Beagle
2894	UCGCAUCGUCGUAGAACAAGCCCCGA	1	no match			
2895	AAAAGCACCGUUUCCCGUCCGAUCAA	1	no match			
2896	AAUCCCAACUGUGAGGGGAUCCUCAA	1	no match			
2897	UCUCAUCGUCGUAGAACAAGCCCGCUGUAACUCGGUCGAAU	1	no match			
2898	AGAAAUCUCAUCGUCGUAGAACAACGCCCGA	1	no match			
2899	AGAAGCAUUCCGAAAGUUCCGAGUACA	1	0	ncRNA		ncRNA
2900	GCCCCGUACUCUGUGGUGACCGACA	1	1	transposon	LINE	baggins
2901	UUUUCAUCUUUAGACGGCUCGGGC	1	2	AT-chX		AT-chX-B
2902	UUUCGCCGAGGUGUGAAAAGUUGA	1	0	transposon	LTR	gypsy12
2903	UUAAACAAUUGAAAGAUAAGACCCAU	1	0	transposon	LINE	Cr1a

2904	CAAAGGACUCGAAAAACUCUUGAA	1	2	transposon	LINE	G2
2905	AAAGGUGGAAAUGAAGAAAGGUGAA	1	no match			
2906	UGUACUACCCCUUCAUGACUCAAAGA	1	0	transposon	LINE	Fw2
2907	UCUCAUCGUCGUAGAACAAGCCCCGACUGUAACUCGGGGUCAAUA	1	no match			
2908	AUCUCUAUCGUCGUAGAAGAAGCCCGA	1	no match			
2909	AUCUCAUCGUCGGUAGAACCAAGCCCGA	1	no match			
2910	CUGGCCGAGUAUUUGAUAUUCUUUG	1	0	Su(Ste)		Su(Ste)
2911	UGUUUCAUCGUUAGACGCUCGGGCG	1	no match			
2912	CCCCGCGGAAAGACGGAAAGACCCCAUG	1	no match			
2913	UGUUUCAUCGUUAGACGGCUCGGGCUGUAACUCGGGGUCAAUAUC	1	no match			
2914	CCUCUACUGACUUCAGUAACGCCCCG	1	no match			
2915	UGAUUCAUCCAGGGAAUGUCAGCGC	1	0	transposon	LTR	ninja-Dsim-like
2916	UACAACUGUGAGGGGAUCCUCAA	1	no match			
2917	UCUCAUCGUCGUAGAACAAGCCCGACUG	1	no match			
2918	UGCCCGAUGAGAAGAAUGGCCCGUA	1	0	transposon	LINE	RR49783
2919	CUAGAUCUUCACCGUUGACUACGCCCCA	1	no match			
2920	AUCGUCUCGGGGAUGAAAUGAAUUUUGGCCAGUAUUUCAUCUC	1	no match			
2921	AUCUCAUCGUCGUAGAACUAGCCCAA	1	2	Su(Ste)		Su(Ste)
2922	CCCAAUGAGAUGGCCUUUGGCGAGA	1	0	transposon	LTR	Quasimodo
2923	UCUGCCCGUCCGAACAAUCUUUGGCC	1	0	transposon	LTR	297
2924	UGAAUACAUCCAGGACAAGUUCAAUC	1	0	coding gene		CG13591-RA|Ssl
2925	UCCUGACGAAGAGCGCAAGAAGAGG	1	0	transposon	LTR	opus
2926	AUCGUCUCGGGGAUGAAAUCUCAUCGUCGUAGAACAAGCCCGAG	1	no match			
2927	GAGGCCCCGUAGUUACCACCCCAC	1	1	no annotation		
2928	UCCGAUCUCCUGAGUCGGACUCCGAA	1	1	transposon	LINE	R1
2929	AUCCUCAACUGUGAGGGGAUCUCCAA	1	no match			
2930	UCUCAUCGUCGUAGAAGAAGCCCGUA	1	2	Su(Ste)		Su(Ste)
2931	AUCUGGAUACUAGGUAUCGACGCCGA	1	0	no annotation		
2932	AAAAUCUCAUCGUCGUAGAACUAGCCCGA	1	no match			
2933	UCUGUUUCAUCGUUAGACGACUCGGG	1	1	AT-chX		AT-chX-B
2934	UCUCCAUCGUCGUAGAACAAGUCCCGA	1	no match			
2935	AUCGUCUCGGGGAUGAAAACGUUUUCAUCGUUUAGACGGCUCGGG	1	no match			
2936	UCUCAUCGUCGUAGAACAAGCCCCGAG	1	2	Su(Ste)		Su(Ste)
2937	UUAUUACCCAGGUGAUUACCACUAUU	1	2	Su(Ste)		Su(Ste)
2938	ACCAGAGAUCUAGAGGGCCCAAGAAA	1	0	no annotation		
2939	AUCGUCGUCGGGAUGAAACUCAACUGUGGAGGGGAUCCUCAA	1	no match			
2940	UCAACUGGUGAGGGGAUCCUCAA	1	no match			
2941	AUCGGUCAGUCGUACGGUAGAACAAGCCCGGGACUG	1	no match			
2942	UACAAAACAAGUACAGGAUCCUCAA	1	no match			
2943	AUCGUCUCCGGGAUGAAAUUUCUGACGAGAACUCGAGCCGCGU	1	no match			
2944	GGAUCCCAACUGUGAGG	1	2	coding gene		CG11326-RA|Tsp
2945	AAUGCCCAAGAAAAUGAUGACCAGCAA	1	0	coding gene		CG42266-RA|CG42266
2946	AUCUCAUCAUCGUAGAACAACGCCCGA	1	no match			
2947	AUCACGCGGGAGACCGGGGUUCAAUU	1	1	ncRNA		tRNA
2948	UCGGCCUUCUGACGAGAACUCGAGCC	1	1	no annotation		
2949	GCCCGUACUCUGUGGUGACCAAU	1	2	transposon	LINE	baggins
2950	UCAUCAUCGUAGAACAAGCCCGA	1	0	Su(Ste)		Su(Ste)
2951	CCUCAACGGAUCCUCAA	1	2	transposon	LTR	roo
2952	UUCCUCGGGAUCAAGGGCAACGAGUUC	1	0	coding gene		CG40635-RA|CG40635
2953	AACUGUGAGGGGGAUCCUCAA	1	no match			
2954	GCCGAACGUAGUCUCUCACGAGCGAGA	1	2	transposon	LTR	invader1
2955	UGGACUUUCCCGAUGAGCCAGU	1	no match			
2956	GAAUAUCGUCUCGGGAGUGAAAUACUCAUCGUCGUAGAACAAGCC	1	no match			
2957	UCUCAUCGUCGUAGAACAAGCCCGACUGUAACUCGGGGUCAAAUC	1	no match			
2958	UAAAGAUUGGACCCAGUACACUGCC	1	2	transposon	LINE	Doc
2959	AUAAAUGCCAGAAUUUUAUUGGGG	1	0	transposon	LINE	baggins
2960	UCUCAACUCCGAACUCAACUGUACCGA	1	0	transposon	LTR	gypsy11
2961	UUAUACAUCUGCGAGCCAUACCACG	1	no match			
2962	UUGGACCAAACACUUUGAACCAAGUA	1	1	Su(Ste)		Su(Ste)
2963	GAGGACUUGAGCGAUUUAAGUGG	1	1	Su(Ste)		Su(Ste)
2964	UCAACUGUGAGGGGGACCCUCAA	1	no match			
2965	UUCCGAGGAACCCAUCGAUCCAGCU	1	1	no annotation		
2966	CCAUCCUAAAACUGAUAGACAGAU	1	0	Su(Ste)		Su(Ste)
2967	UGUUUCAUCUUUAGACGGCUCGGCC	1	2	AT-chX		AT-chX-B
2968	UCCGAAGAGGAUGGAUACCGGCA	1	no match			
2969	CCCGAAGCGGACCAGCCCCAAUUAUUCUGUAACUCGGGGUCAAU	1	no match			
2970	UCUCAUCAUCGUAGAACAAGCCC	1	0	Su(Ste)		Su(Ste)
2971	GACUAGCUGGUCGAGGAUCCGAUU	1	1	Su(Ste)		Su(Ste)
2972	UCGGGAUAAGGAUUGGCUCUGAAGAU	1	0	ncRNA		rRNA
2973	UAAAUGCCAGAACUUAUUGGGGUUC	1	no match			
2974	AACUAGUGAGGAGGAUCCUCAA	1	no match			
2975	AACAACUGUGAGGGGGAUCCUCAA	1	no match			
2976	UGUUUCAUCGUUAGAAGGCUCGGGC	1	1	AT-chX		AT-chX-B
2977	CUCUCAUCUUCGUAGAACAACGCCCGA	1	no match			
2978	UGUUUUUAGGUGUACUGACCCAGUC	1	1	transposon	LINE	baggins
2979	CGAAAGUAGAUCUUCACCGUUGACUUGGCC	1	2	Su(Ste)		Su(Ste)
2980	UAAAAGAAAUCCUGACAAGAUCCCGGGC	1	no match			
2981	ACAAGAUAUGACCUAAGCAAAGACAC	1	1	transposon	DNA	1360
2982	CUGUAACUCGGUCAAUCGGGUCAAU	1	no match			
2983	GAUGAACCCCAGGAAGCUCACGCUU	1	0	transposon	LTR	springer
2984	UAAAUGCCAGAAUUUUAUUGGGGUUC	1	no match			
2985	UCUCAUCGUCGUAGAAUUAGCCCGA	1	1	Su(Ste)		Su(Ste)
2986	AUCGUCUCGGGGAUGAAAAUGUUUCAUCUUAGACGGCUCGGGC	1	no match			

2987	GCUCCAAAAUGGCGGCCGCCCCGACGC	1	0	transposon	LTR	ninja-Dsim-like
2988	AUCUCAUCGUCGUAGAACUAGCCCGAGG	1	1	Su(Ste)		Su(Ste)
2989	CACCAGGAAGAUUGAUGAGGACA	1	0	transposon	LINE	Rt1b
2990	CACACACAGAGGAGAACAGAUGAGAGCA	1	no match			
2991	UCCGAAAGAGAAAAUAUUCAUC	1	1	transposon	LTR	HMS-Beagle
2992	UUCGGACCGAACACUUUGAACCAAGUCUGUAACCUCGGGUCAAU	1	no match			
2993	UUCUCAUCGUCGUAGAACAAGCCCGCG	1	1	Su(Ste)		Su(Ste)
2994	CGAAAUGUUUCAUCGUUACGACGGCCUCGGGC	1	no match			
2995	CUUCAUCAUCGUAGAACAAGCCCGA	1	2	Su(Ste)		Su(Ste)
2996	GCACACAGAGGAGAGCGAGAUGAGAGCA	1	no match			
2997	UCUCAUCGUCGUAGAACAAGCCCGACUGUAACUCGGGGUCAAUAU	1	no match			
2998	GGAUCCUCAACUGUGAGGUGAUCCUCAA	1	no match			
2999	ACUGUGAGGUGAUCCUCAA	1	no match			
3000	UGUUUUCAUCGUAAGACGGCUCGGG	1	no match			
3001	GACUACCUGGCCGAGUAUUUGAUA	1	1	Su(Ste)		Su(Ste)
3002	GAUUCCGGAGAGGGAGCCUGAGAAACG	1	0	ncRNA		rRNA
3003	AUCGUCUCCGGGAUGAAAUCUCAUCGUCGUAGAACAAGCCCGA	1	no match			
3004	GAAGAAGAAAGUACAGUGAGG	1	2	no annotation		
3005	GAUAUGCUGCGAGUUGCCAAUCUCC	1	0	transposon	LINE	X
3006	GAUGUCCGCGAGAUUGCCGCAAACU	1	0	transposon	LTR	opus
3007	GAGACUCAACUGUGAGGGGAUCCUCAA	1	no match			
3008	ACGACCCGCUACCGGAGGACGCCA	1	0	transposon	LTR	opus
3009	AGCACACAGAGGAGAGCAGAUGAGACA	1	no match			
3010	UAAGAAAAUGUCCGGACAGCUGG	1	0	Su(Ste)		Su(Ste)
3011	CGAAGGUAUUCGUAAUGACCGCUU	1	0	transposon	LTR	297
3012	UCUCGAUCGUCGUAGAACAAGCCCGCG	1	no match			
3013	UCUCAUCGUCGUAGAACAAGCCCGACUGUAACUGGGUCAAU	1	no match			
3014	UUCGAGAAGCGAACCCGGUGAACUGAA	1	no match			
3015	GGAUUCUCAACUGUGAGGGGAUCCUCAA	1	no match			
3016	CCAUCAGAGACUUUGGUGGCCAACAA	1	0	coding gene		pseudogene
3017	GGAUCCCAACUGUGAGUCCUCAA	1	no match			
3018	UAUAAUGCCCCGUAGGUUGCCCUCU	1	0	transposon	LINE	R1-element
3019	UCUCAUCGUCCGUAGAACAAGCCCGCUGUAACUCCGGGUCAAU	1	no match			
3020	ACUUACGGGAGUAAGGAUUGGCUCUG	1	no match			
3021	CUCAACUGUGAGGGGAUCCUUAA	1	no match			
3022	UUCUCAUCGUCGUAGAACAAGCC	1	0	Su(Ste)		Su(Ste)
3023	GAAUCAUGGAGAUGAUACCGAGCUUC	1	0	transposon	LINE	BS3
3024	UGUUUCAUCUUAGACGGCUCGGGUC	1	no match			
3025	UAGAUCUUCACCGUUGACUUGGCCA	1	1	Su(Ste)		Su(Ste)
3026	UAUAAGGUUGGCCCAAGAACGCUGC	1	0	transposon	Low_complexity	F-element
3027	CAGCACACAGAGGAGAGCAGAGAGA	1	no match			
3028	UGAUACCUAGAAUGACAAUGGUUGA	1	1	transposon	LTR	gypsy3
3029	CGAAAUAGAUCUUCACCGUUGACCUAGCCCCA	1	no match			
3030	GAAUAUCGUCUCGGGAGUGAAAUACUCCAUCGUCGUAGAACAAGC	1	no match			
3031	UGUUUCAUCGUUAGACGGCUCGGGGCCUGUAACUCGGGGUCAAUA	1	no match			
3032	UGCACCGGAACGACCAACCAGCCCA	1	0	transposon	LINE	Doc3
3033	CGCCGUGGUCAACCUGGACGUCCCAGU	1	1	coding gene		CG17957-RA|Sry-alpha
3034	UAGUAUCUUCUACCGUCGACUUGGCC	1	no match			
3035	UACUGUUUGUAGGUGUACUGACCA	1	1	transposon	LINE	baggins
3036	UGCAACCCCCAACGACCACCCGCUGUU	1	0	coding gene		CG2098-RA|ferrochelatase
3037	UCUCGAUCGUCGUAGAACAACCCCGA	1	no match			
3038	CGCACACAGAGGAGAGCAGAUGAGGAGCA	1	no match			
3039	CAAAGAAGCCCCGCCGCCGCCGCC	1	no match			
3040	AUCUAGGCUGAGCGGAGAACCAGAGU	1	0	transposon	Low_complexity	F-element
3041	UUCGCUACCAGACGGGAUUACUCUG	1	0	transposon	LTR	ninja-Dsim-like
3042	UCCUGACGAAGAUCGCAAGAAGAG	1	0	transposon	LTR	opus
3043	UCCGAGGAACCAAUUGAUCCAGCUCC	1	1	Su(Ste)		Su(Ste)
3044	UAUGCUCUCGAAUGUAUGGCCCGAUA	1	no match			
3045	UCUCAUCGUCGUAGAACAAGCCCCGAGCUGUAACGUCGGGUACAA	1	no match			
3046	CAAAAGUUGACCGAACUUGAUUAUU	1	0	ncRNA		rRNA
3047	UAAAAAGAUAAAAACCGAAUUACAAGAA	1	2	repeat		trf
3048	UGCUCGAUGAGAAGAAUGGCCCGUAA	1	1	transposon	LINE	RR49783
3049	GCGACUCCAAGAGCGAUGUGCCGAA	1	0	coding gene		CG13913-RA|CG13913
3050	ACCCUGUAGAUCCGAAUUUGUU	1	0	ncRNA		mir-10-RA
3051	AAAGGUAAUAUCCGAGUCGUUCCC	1	0	transposon	LTR	opus
3052	UCCAUCUUCUUCUCGACGUUGACCCA	1	no match			
3053	GAAUAUCGUCUCGGGAGUGAAACUGUGAGGGGAUCCUCAA	1	no match			
3054	UUGUGACUGAUCUAGAGUACAAUGU	1	0	transposon	LTR	DM88
3055	UCUGGUUGAACGUGUCCUGAAAG	1	0	Su(Ste)		Su(Ste)
3056	UAUCAUCAUCGUAGAACAAGCCCGU	1	2	Su(Ste)		Su(Ste)
3057	UCGGCCUGGAUCACGGACAUCUG	1	0	transposon	LTR	ninja-Dsim-like
3058	AUCGUCUCGGAUGAAAUGUUUCAUCGUUAGACGGCGUCGGGC	1	no match			
3059	UAGGCUUACCCGGGGAUACAUAGU	1	0	transposon	LINE	Doc
3060	UAGAUCCACCGUUGACUUGGCCC	1	no match			
3061	UUCAUCUUCUUCUCGACGUUGACCAG	1	0	transposon	LTR	ninja-Dsim-like
3062	GCCCUUGAUUCCGAGGAGCCAAUU	1	1	Su(Ste)		Su(Ste)
3063	AUCUCAUCGUCGUAGAACUAGCCCCGU	1	no match			
3064	UCCUCAACUGUGAUGGAUCCUCAA	1	no match			
3065	CGAAAUCUCAUCUCGUAGAACAAGCCCGA	1	no match			
3066	CACUGUGAGGGGAUCCUCAA	1	no match			
3067	CCAAUUGAUCCAGCUGCUGUUGUU	1	0	Su(Ste)		Su(Ste)
3068	UGAAAUGGACCUGGAAGACGCCCCACC	1	0	coding gene		CG4266-RA|CG4266
3069	GGAUCCUCAACUGUAAGGAA	1	no match			

3070	ACCCACCCUGAUGUGCUUGACGUU	1	no match			
3071	UAGGAUCUUCACCGUUGACUUGGCCCCG	1	no match			
3072	UGUUUUCAUCCGUUAGACGGCUCGGGUCA	1	no match			
3073	AUCCUCAACUGUGAGGGAUCCUCAACUGUAACUCGGGGUCAAU	1	no match			
3074	UCUCAUCGUCGUAGAACUAGCCCCA	1	1	Su(Ste)		Su(Ste)
3075	GAAGAAUCCUGUGAGACCCAGAAAUCC	1	0	transposon	LTR	flea
3076	GAUGUCCACCAAAGGCAUGCCCGA	1	0	transposon	LTR	opus
3077	UCUCAUCGUCUUAGAACAAGCCCGAG	1	1	Su(Ste)		Su(Ste)
3078	AAAAUACUACAUCCGUACGUUAGAACCAAGCCCGA	1	no match			
3079	CAGCACACAGAGGAGAGCAGAUGAGACGCAC	1	no match			
3080	UUAUCGACUCGUGUACCUGGUUA	1	0	transposon	LTR	micropia
3081	UCGAAAUUCUCAUCGUCGUAGAACUAGCCCGACUGUAACUCGGGG	1	no match			
3082	UUCUCGAUCGUCGUAGAACAAGCCCGACG	1	no match			
3083	UGGCAUUCGCGUCGAGGCCCAGGUAUA	1	no match			
3084	UUCGAACUGAAGGAGUACGAAUGAGUC	1	no match			
3085	CCUGCUGACGGGGCCCGAAAACACG	1	0	transposon	LTR	opus
3086	UGUUUCAUCGUUAGACGGCUCCGGG	1	2	AT-chX		AT-chX-B
3087	GCACACAGAGGAGAGCAAGAUGAGAGCA	1	no match			
3088	UCCGAGGAACCCAUUGAUCCAGCUGC	1	0	no annotation		
3089	CAGCACACAGAGGAGAGAAGAUGAGAGCA	1	no match			
3090	CGAAAUCCAUCGUCGUAGAACUACGCCCGA	1	no match			
3091	UAGAUCUUCACCGUUGACUACGCCCCC	1	no match			
3092	AAUAUGCCAGAAUUUUUAUUGGGGGUUC	1	no match			
3093	UGUUUCAUCGUUAGAAGGCUCGGG	1	1	AT-chX		AT-chX-B
3094	UUGUUUCGGAGCGUUGGCCAGUUUC	1	1	transposon	Low_complexity	F-element
3095	CAUGGAUUGGCUUGUUUUGUUCAGUUG	1	no match			
3096	CUCUCAUCGUCGUAGAACUAGCCCGA	1	1	Su(Ste)		Su(Ste)
3097	ACUGCUCAUUCCGAACAUACACCGA	1	0	transposon	LINE	Doc
3098	UCCUGAAGCCGGUGAUCGACAGUUA	1	1	coding gene		CG42398-RA|SteXh:CG42398
3099	GCUUCCGGGAUUUUGACGCCGUUAAU	1	0	transposon	LTR	opus
3100	AAGGAUCAUGAUACCGACUAAAGGCAA	1	1	no annotation		
3101	CGAAAUCUCAUCGUCGUAGAACAAGCCCG	1	no match			
3102	GAGAGUGAUGCCGAGUAUAAAAGUAA	1	0	repeat		trf
3103	UAACAUCAUCAUCUGACUUGCAAUU	1	1	transposon	DNA	1360
3104	UCCUAAUCGCGAGCUUAUUGUUGUACC	1	0	transposon	LTR	diver2
3105	GGUCCUCAACUGUGAGGGGAUCCUCAA	1	no match			
3106	UCUGAACGUACCACGGUGCCCCAGG	1	2	transposon	Low_complexity	F-element
3107	AAUAGGGACGGUCGGGGGCAUCAGU	1	no match			
3108	CUCUCAUCGUCGUAGAAGAACGCCCG	1	no match			
3109	CAGCACACAGAGGAGAGAUGAGAGCA	1	no match			
3110	UCUCCAUCGUCGUAGAAUUAGCCCGA	1	no match			
3111	UGGUUCAUCGUUAGACGGGACUACGGGCCCG	1	no match			
3112	UUUCACAUCGUCGUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
3113	UGAUCUUCACCGUUGACUUGGCC	1	1	Su(Ste)		Su(Ste)
3114	CUCUCAUCGUCGUAGAAACAACGCCCGUA	1	no match			
3115	UCCGAUUUGGCGCAAAUAGAUGGAC	1	2	transposon	LTR	ninja-Dsim-like
3116	UAAAGGUAUUCCUUGUCGAACGUCGUCUG	1	no match			
3117	UGAUACCGAGAAUUACAAUGGUUGA	1	1	transposon	LTR	gypsy3
3118	AAACCUGAAGGUGAUCCUGAGUCGCA	1	1	transposon	LINE	Helena
3119	UCUCAUCGUCGUAGAACAAAGCCCCGA	1	no match			
3120	UUAUACCCAGGUGAUUCCCACUAU	1	0	Su(Ste)		Su(Ste)
3121	GGACUUUUUUCCGAUGAGCCAGU	1	0	transposon	LINE	G2
3122	UGUUUCAUCAGACGGCUCGGGC	1	no match			
3123	GAUCUCAACUGUGAGGGGAUUCUCAA	1	no match			
3124	GUUCACCGGACGGGCCUAGGGAACU	1	no match			
3125	UCUCAUCUUCGUAGACAAGCCCGA	1	no match			
3126	UUUUUCAUCGUUAGACGACUCGGGC	1	2	AT-chX		AT-chX-B
3127	UAGAUCUUCACGUUGACUUGGCCCU	1	no match			
3128	AUCGUCUCGGGAGUGAAAUGUUUCAUCGUUAGACGGCUCGGGC	1	no match			
3129	CAAGGUGGAAAUGAAGAAAGUACA	1	no match			
3130	CCCAGGCAGUGGGGACUCCAGAUUCC	1	0	transposon	LTR	Tabor
3131	UCGGUGGGCGAAGAGUUGAGUGG	1	0	transposon	Low_complexity	F-element
3132	UGACGAGAACUCGAGCCGCGUAGCC	1	0	no annotation		
3133	GUUCCGGGGGAAGUAUGGUUGCAAAG	1	1	ncRNA		rRNA
3134	UCCUAGAAAAAACGAGGUUCACCCAA	1	0	transposon	Low_complexity	F-element
3135	AAAGACACCGUUUCCCGUCCGAUCAA	1	no match			
3136	UAACCGUUCGUUGGCGUUGAUGGCA	1	0	transposon	LTR	HMS-Beagle
3137	AAGAAAGCACCGUUUCCCGUCCGAUCAA	1	no match			
3138	CAACCGGACGGGCCUAGAGUUAU	1	0	coding gene		CG40635-RA|CG40635
3139	UUCUGACGAGAACUCGAGCCGCGUAGC	1	0	no annotation		
3140	UGUUUUCUAUCGUUAGACGGCUCGGGC	1	no match			
3141	CUCGGGAUCAAGGGCAACGAGUAA	1	2	Su(Ste)		Su(Ste)
3142	CUAUAACUCGGGUCA	1	1	transposon	LTR	gypsy8
3143	AUUGGCAUUUAUUUAUGCAUCCCGUU	1	1	transposon	LTR	Stalker
3144	CAGGCACGACAGAGGAGAGCGAGAUGAGAGCA	1	no match			
3145	UCUGGAUACUAGGUAUCGACGCCGAUA	1	0	no annotation		
3146	UCUGAAGACCCCGACGGUUAUGCA	1	0	transposon	LTR	297
3147	UUCUUUUGAAGAUGGCCCACAUAAAG	1	2	repeat		trf
3148	UUGAUACGAAUGAGACCCGCAGUA	1	0	repeat		trf
3149	ACCAGGACGGGUGACGCAUAAGUUU	1	0	transposon	LINE	R1-element
3150	UCUCAUCGUCGUAGAAGAGCCCCGA	1	no match			
3151	UAAUGCCUAACGAGGUAGAUGACUC	1	0	transposon	LINE	Cr1a
3152	UCGGUUGCCCACUUACAGGCCCAAA	1	0	transposon	LTR	ninja-Dsim-like

3153	UCCUGAAGCCGGUGAUCGAUAGUUCC	1	0	repeat		trf
3154	GAAGAAUUGUAGGUUAUAGCCGUAGAG	1	1	ncRNA		pseudogene
3155	UUCGAUUACCUGGUUCGAUCCAAU	1	0	transposon	LTR	Dm88
3156	UUCUCGAGGAACAGUUUGACGAGCUA	1	0	coding gene		CG9279-RA|CG9279
3157	UUUGUAGGUGUACGGCCCAGUCGG	1	no match			
3158	UCGUCAGUCGUCGUAGAACAAGCCCGGACU	1	no match			
3159	UUUAUCCGAAAUCGUGUUCGAGUU	1	0	transposon	LTR	micropia
3160	AUCGCUCGGGAUGAAAUGUUUCAUCGUUAGACGGCUCGGGC	1	no match			
3161	UAGGUUAUGGGCCCAGUGAACAUU	1	1	ncRNA		alphagamma-element
3162	UCCAGUCGUCGUAGAACAAGCCCGACUG	1	no match			
3163	UCGAGUUGCAGAAUUGAGGACCCCGUC	1	1	transposon	LTR	Quasimodo
3164	GCCCCAGUUGCCUAAUGGAUAAGGC	1	1	ncRNA		tRNA
3165	AAAUAGGGGACGGGUCGGGGGGCAUCAGU	1	no match			
3166	GAAUAUCGUCUCGGGAGUGAAAACUUACGGGAGUAAGGAUUGGCU	1	no match			
3167	UUAUAAGGAUGAUGAUACCGAUUAAAG	1	0	no annotation		
3168	AUCGUCUCCGGGAUCGAAAUAGAUCCUUCACCGUUGACCUUGGCC	1	no match			
3169	AUCCUUCAACUGUGAGGGGAUCCUCAA	1	no match			
3170	AUCUCAUCGUCGUAGAACAAAGCCCCGAGG	1	no match			
3171	UUGAUCUUCACCGUUGACUAGCCCA	1	2	Su(Ste)		Su(Ste)
3172	UUGGACUUUUUUCCGAUGAUCCAGU	1	1	transposon	LINE	G2
3173	CACGUCAGUUCGGAUGCCUACUACCGG	1	no match			
3174	UAUACACCGGCAAGAACAAAACGA	1	0	repeat		HETRP_DM
3175	UCUCGUCGUCGUAGAACAAGCCGA	1	no match			
3176	AUUCUCAUCAUCGUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
3177	CAACUGUGAGGAGAUCCUCAA	1	no match			
3178	UUCUCAUCAUCGUAGAACAAGCCCG	1	0	Su(Ste)		Su(Ste)
3179	UGUUUUCAUCGUUUAGACGGCUCGGGU	1	no match			
3180	UAGAUCUUCUUCGUUGACUAGCCCCA	1	2	Su(Ste)		Su(Ste)
3181	GAAAAGCACCGUUUCCCGUCCGAUCAA	1	no match			
3182	UAGCUGGCCGAGUAUUUGAUAUUCUUU	1	0	Su(Ste)		Su(Ste)
3183	UCAACUGUGACUCAACUGUGACUCAA	1	no match			
3184	GAAGACCCUGUUGAGCUUGACUCU	1	1	repeat		LSU-rRNA_Dme
3185	CUCUCAUCAUCGUAGAACUAGCCCGA	1	2	Su(Ste)		Su(Ste)
3186	UAGAUCUGCACCGUUGACUAGCCCC	1	1	Su(Ste)		Su(Ste)
3187	AAAUCUUCGUCGUAGAACAAGCCCGA	1	no match			
3188	UAAAUGCCAGAACUUUAUUGGGGUUC	1	0	transposon	LINE	baggins
3189	UUGUAUGUGGGGUAGGGGAGCGUU	1	no match			
3190	UGUUUCAUCGUUAGACGGCUCGGGCCUA	1	2	AT-chX		AT-chX-B
3191	UAAAGAAAUCCUGACAAGAUCCCGG	1	no match			
3192	GUUCUUUGAAGAUGGCCACAUAAAG	1	no match			
3193	UCUCAUCGUCGUAACAGCCCGA	1	no match			
3194	UCCGAGGAACCAAUUGAUUCAGCUG	1	0	Su(Ste)		Su(Ste)
3195	CUCUCAUCCGUCGUAGAACAACGCCCGU	1	no match			
3196	CAGCACACAGAGGAGAGCAGAUGAGAACA	1	no match			
3197	GCAUAUAAUGCCCGAAGGUUGCCC	1	2	transposon	LINE	R1-element
3198	GAUGCCAUCAAGCGCACUGAUUAACCGA	1	0	no annotation		
3199	GAAAUCUCAUCGUCGUAGAACAAGUCCCGAACUC	1	no match			
3200	GUAGAUCUUCACCGUUGACUAGCCCCG	1	1	Su(Ste)		Su(Ste)
3201	CGAAAAAACCUGAAGAUGAUCCUCGAGUCGC	1	no match			
3202	GGAUCCUCAACUGUGAGGA	1	no match			
3203	UCAUGACCGAAAUUCGAAAAAACGAA	1	1	repeat		SAR_DM
3204	UUCUCAACGUCGUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
3205	CUCGCUGAGAGCUGAGAGCCCGCA	1	0	ncRNA		alphagamma-element
3206	UGUCUCAUCGUUAGACGGCUCGG	1	1	AT-chX		AT-chX-B
3207	AUCGUCGUCGGGAUAGAAAUCUUAUCGUCGUAGAACAAGCCCGA	1	no match			
3208	AAAGGUGGGAAUGAAGAAAGUACA	1	2	no annotation		
3209	UAGAUUUUCACCGUUGACUUGGCCCA	1	1	Su(Ste)		Su(Ste)
3210	GUAUGACCCCUGGUACUGAAAUG	1	0	transposon	DNA	Bari1
3211	UCAACUGUGAGGUGAUCCUCCAA	1	no match			
3212	UGCCUAUCUUGAGCGGCGAGGACC	1	0	transposon	LINE	Cr1a
3213	CCUGGAUGAUGAUAAGCAAAUGCUGA	1	no match			
3214	UAACAAGAUGCAUACGAUCGUGGC	1	no match			
3215	UUCCGCAUGUGAAUGUUGGCCCCAGA	1	0	no annotation		
3216	UCCGAACGUACCACGGUGCCUCAGU	1	1	transposon	Low_complexity	F-element
3217	AUCUUCAUCGUCGUAGACAGCCG	1	no match			
3218	AUCGCAUCGUCGUAGAACAAGCCCGAGG	1	1	Su(Ste)		Su(Ste)
3219	UCUCAUCGUCGUAGAACAAGCCCGCUGUAACUCGGUCAAU	1	no match			
3220	AGAUCUUCACCGUUGACUUGGCCCAU	1	1	Su(Ste)		Su(Ste)
3221	UGUCUCAUCGUUAGACGACUCGGGC	1	2	AT-chX		AT-chX-B
3222	CAGGCACGACAGAGGAGAGCGAGAGUGAGAGCA	1	no match			
3223	UGUUUCAUCGUUAUACGGCUCGGA	1	2	AT-chX		AT-chX-B
3224	CGCGACGACGCCGAGUGGAAUCCAGG	1	0	transposon	LTR	ninja-Dsim-like
3225	CAACACAGAGGAGAGCAGACGAGAGCA	1	no match			
3226	AGCACACAGAGGAGAGCAGAUGAGAGCAA	1	no match			
3227	UUGUACUACCCCUUCAUGACUCAAAGA	1	0	transposon	LINE	Fw2
3228	CAAACAUUAGCUCCGAUCUCCUGAGU	1	0	transposon	LINE	R1-element
3229	CAGCACACAGAGGAGACAGAUGGAGCA	1	no match			
3230	CCCGAUGAGAAGAAUGGCCCGUAAA	1	0	transposon	LINE	RR49783
3231	GGUGGAAAUGAAGAAAGUACAAA	1	no match			
3232	GAGGAGGAUCCUCAA	1	1	coding gene		CG13685-RA|CG13685
3233	GGAUCCUCAAUUGUGGGGGGAUCCUCAA	1	no match			
3234	UGGACUUUUUUCUGAUGAGCCAGU	1	1	transposon	LINE	G2
3235	AUCUCAUCGUCGUAAAAACAAAGCCCGAC	1	no match			

3236	CGAGCAAUCAUCCAAGGCUGAGAUCCGG	1	0	transposon	LTR	mdg3
3237	UAGAUUGGACCCAGUACACUGCC	1	1	transposon	LINE	Doc
3238	UGUUUCAUCGUUACGGCUCGGGC	1	no match			
3239	GCCCGAUCAGAAGAAUGGCCCGUAAAG	1	1	transposon	LINE	RR49783
3240	CGUUCCCGUCUCGUGCUAAAGGGUUGAC	1	no match			
3241	UGUUUUCAUCGUUUAGACGGCUCGGG	1	no match			
3242	CGAAAUAGAUCUUCACCGUUGACCUAGCCCC	1	no match			
3243	ACUACCGAGCCCGAGCCUUACUCGAC	1	0	transposon	LTR	Rt1c
3244	UCUCAUCUUCGUAGAACAGCCCGACUGUAACUCGGGGUCAAU	1	no match			
3245	AUCGGGCUUUCGUGACGAGAACUCGAGGC	1	no match			
3246	UAGGAUCUUCACCGUUGACUAGCCCCG	1	no match			
3247	CUGACGAGAACUCGAGCCGCGUAG	1	0	no annotation		
3248	UGUUCAUCGUUAGACCGGCUCGGG	1	no match			
3249	UAAGCGUUCGUUGACGUUGAUGGCUG	1	0	transposon	LTR	HMS-Beagle
3250	UGCUCCGCUAUGCUAGCUCCCGUA	1	0	coding gene		CG5827-RA|RpL37A
3251	UCUCAUCGUCGUAGAACUAGGCCCGA	1	no match			
3252	UACUUACAGUGGGUUGUCCAGGG	1	0	Su(Ste)		Su(Ste)
3253	UCUCAUCGUCGUAGAACAAGCCCGCUA	1	no match			
3254	UCUCAUCGUCGUAGAAACAAGCCCGAGG	1	no match			
3255	CUCGGGAUGAAAAUCCUCAA	1	2	no annotation		
3256	AUCUCAUCGUCGGUAGAACAAGCCCGA	1	no match			
3257	UGUUUCAUCGUUAGACGGCGUCGGGCG	1	no match			
3258	AAUAAGAAGUUGACCGAACUUGAU	1	no match			
3259	GCACACGAGAGGAGAGCAGAUGAGAGCA	1	no match			
3260	UGCCCCGUACUCUGUGGUGACCGA	1	1	transposon	LINE	baggins
3261	UGUUUCAUCGUUAGACGGCACGGGCG	1	1	AT-chX		AT-chX-B
3262	UCUCAUCGUCGUAGAAGAAGCCCGACUGUAACUCGGGGUCAAU	1	no match			
3263	AUCUCAUCGUCGUAGAACAAGCCCCGAGG	1	no match			
3264	UGUACUACCCCUUCAUGACUCAAA	1	0	transposon	LINE	Fw2
3265	UCUCAUCGUCGUAGAACAAAGCCCG	1	no match			
3266	UGCAAAGACGAAGUGCGCCCCGAU	1	0	transposon	LTR	ninja-Dsim-like
3267	UCGCAUCGUCGUAGAACAAGCCCA	1	1	Su(Ste)		Su(Ste)
3268	GUCUACGCCGAGGCCAACCUCUU	1	no match			
3269	CUCAACUGUUAGGGAUCCUCAA	1	no match			
3270	AUCGGUCUCGGGGAUGAACAGCACGACAGAGGAGAGCAGAUGAGA	1	no match			
3271	UUCAACGGUUACCCCAGGAUGCUGU	1	no match			
3272	UCCGACGCUCUUGACUCGACCAGG	1	0	transposon	LTR	ninja-Dsim-like
3273	AUCGUCUCCGGGAUGAACAGCACACAGAGGAGAGCAGAUGAGAGC	1	no match			
3274	UCAAGGUAGUAACUAUGACGAUUUGACUGUAACUCGGGGUCAAU	1	no match			
3275	UACUCAUCGUCGUAGAACAAGCCCGC	1	2	Su(Ste)		Su(Ste)
3276	UACUGUCCCUUAAAUCAGCACUCGU	1	0	no annotation		
3277	CUACUAACUGUGAGGGGAUCCUCAA	1	no match			
3278	AAUUAAUGCCGGAAGCGAUCCCACU	1	0	AT-chX		AT-chX-B
3279	AGCCCGAUGAGAAGAAUGGCCCGUAAA	1	1	transposon	LINE	RR49783
3280	UUGACAAUUCCAAGGCUGAGAUCCGG	1	no match			
3281	AGAAAAAGGUGGAAUGAAGAAAGUACA	1	no match			
3282	UACUUGUUCCCCGGAUAGUUUAGUUA	1	0	ncRNA		rRNA
3283	UCUACUUCCCACCUAGACACUCGA	1	0	transposon	LINE	G2
3284	GGAUCCUCAACUGGUGAGGGGAUCUCAA	1	no match			
3285	CCUCAACUCGUUAGGAA	1	0	no annotation		
3286	AUCGUCUCGGGACUGAAAUAGAUCUUCACCGUUCGACUAGCCCA	1	no match			
3287	CGAAGGAAAACCUGACCGGACCUAU	1	1	Su(Ste)		Su(Ste)
3288	UCUCAUCGUCGUAGAACCAAGCCCGA	1	no match			
3289	AUUGAUCCAGCUGCUGUUGUUGU	1	0	Su(Ste)		Su(Ste)
3290	UCUAUAGGAAAAACCUAGUUACUUACG	1	0	Su(Ste)		Su(Ste)
3291	UUCAACUGUGAGGGAUCCUCAA	1	no match			
3292	UCGGUGGCGAAGUUGGCUCAGAGG	1	0	transposon	LTR	297
3293	UCUUGUUUCAGAGCGUUGGCCAG	1	0	transposon	Low_complexity	F-element
3294	UCUCUAUCGUCGUAGAAGAAGCCCGA	1	no match			
3295	UACCGGUAUAGCACACUUUUGUAUAA	1	2	coding gene		CG41580-RA|CG41580
3296	CGUCCGGGAUGAAAAUCCUCAA	1	no match			
3297	AAGGUCAACUGUGAGGGGAUCCUCAA	1	no match			
3298	UUCCGAGGAACCAAUGGAUCCAGCU	1	1	Su(Ste)		Su(Ste)
3299	UCCGUCUUUGAGAGGAAUCAGAUGG	1	0	transposon	LTR	opus
3300	GGUUGUAAACCUCUUUCUAGGGACGAAG	1	no match			
3301	AAAACCUAGUUACUUACGGGCUCGA	1	0	Su(Ste)		Su(Ste)
3302	UUCUCAACUGUGAGGGGAUCCUCAA	1	no match			
3303	AUAAGGGAGUCGGCAAAAUAGAU	1	no match			
3304	AUAGAUCUUCACCGUUGACUAGCCCCU	1	2	Su(Ste)		Su(Ste)
3305	UCAACGUCGUAGAACAAGCCCGAA	1	1	Su(Ste)		Su(Ste)
3306	UCUCUCGUCGUAGAACAAGCCCGA	1	2	no annotation		
3307	AUACUCUGACUCUGUGCUGCUUAGG	1	0	transposon	LTR	DM88
3308	UAACAAGAUGCGUUACGCCAUACGA	1	0	transposon	DNA	1360
3309	AGAUCUUCACCGUUGACUAGCCCCUC	1	1	Su(Ste)		Su(Ste)
3310	UAAAUGCCAGAAUGUUAUUGGGGUUC	1	1	transposon	LINE	baggins
3311	GAUUCCGAGGAAUCUGGGUUCAAGA	1	no match			
3312	CGUAACUUCGGGAUAAGGAUUGGCUCUA	1	1	ncRNA		rRNA
3313	UUUAAUGCCGGAAGCGAUCCCA	1	1	AT-chX		AT-chX-B
3314	AGCUGGCCGAGUAUUUGAUAUUCUU	1	0	Su(Ste)		Su(Ste)
3315	UAACGACCAGACAUACGGCAUUAU	1	1	transposon	LINE	R1-element
3316	UCUCAUCGUCGUAUAACGAGCCCGA	1	2	Su(Ste)		Su(Ste)
3317	UCCGUCGAGGUGGUGGUGCGAUC	1	0	transposon	LINE	R1
3318	GCCGAACGUAGUCUCUCGAGAACG	1	1	transposon	LTR	invader1

3319	AUCUCAUCUUCGUAAGAAACAAGCCCGA	1	no match			
3320	UUCGUCAUCGUCGUAGAACAAGCCCGG	1	no match			
3321	CAGCGACACAGAGGAGAGCAGGAUGAGAGCA	1	no match			
3322	UCUCAUCGUCGUAGAACAACCCCGAG	1	1	Su(Ste)		Su(Ste)
3323	CUGUUUCAUCGUUAGACGGCUCCGG	1	1	AT-chX		AT-chX-B
3324	CAGCGACACAGAGGAGAGCAGAUGAGAGCA	1	no match			
3325	AUAAGAUCUUCACCGUUGACUAGCCCCCUGUAACUCGGGGUCAAU	1	no match			
3326	CAACUGUGGAGGGGAUCCUCAA	1	no match			
3327	UAAUGCCCCGAAGGUUGCCCUCUGUG	1	0	transposon	LINE	R1-element
3328	UCUCAUCGUCGUAGAACAAGCCCCGACUGUAACGUCGGGUCAAUA	1	no match			
3329	GCUAUGAUAUACCGGACAGUGUGCAA	1	0	transposon	LTR	gypsy12
3330	UCCUGUGAGGGGAUCCUCAA	1	no match			
3331	UCUCGAUCGUCGUAGAACAAGCCCGGCG	1	no match			
3332	CCGCCGCCCAGUCCUGCUCGCUUCACC	1	no match			
3333	UUUUCCUGAAAUCCGAGUUAAUAUG	1	1	no annotation		
3334	UUUUCAUCGUUAGACGGCUCGUGC	1	2	AT-chX		AT-chX-B
3335	UGUUUCAUCGUUAGACGGCUCGGGUCAAU	1	no match			
3336	UCUCAUCGUAGUAGAACUAGCCCGGG	1	2	Su(Ste)		Su(Ste)
3337	UUCAUCGUCGUAGAACAAGCCCGACG	1	1	no annotation		
3338	UUCUGACGAGAACUCGAGCCACGU	1	1	no annotation		
3339	UCCAAAGGACUACGAAAAACUCUUGA	1	no match			
3340	CAAGAUAUUACCAAAGCAAAGACAC	1	1	transposon	LINE	Cr1a
3341	UAAAUGCCCCGAAGGUUGCCCUCUG	1	2	transposon	LINE	R1-element
3342	UAAAAUGCCAGAAUUUUAUUGGGUUC	1	no match			
3343	UCCUCAACUGUGAGGGGUUCCUCAA	1	no match			
3344	AUCGUCUACGGGAUGAAAUAGAUCUUCACCGUUGACUAGCCCCA	1	no match			
3345	AUCGUCUCGGGACUGAAACUCUCAUCCGUCGUAGAACAACGCCCG	1	no match			
3346	UCUCAUCGUCGUAGAACAAGCCGUG	1	no match			
3347	UGUUUCAUCGUUAUACGGCUCGGGCA	1	2	AT-chX		AT-chX-B
3348	CAGCACACAGAGGAGAGCAGAUGAGAGCUA	1	no match			
3349	UAGAUCUUCACCGUUGACUAGCCCCUC	1	1	Su(Ste)		Su(Ste)
3350	UGUUUCAUCGUUAGACGGCUCGGGGCUGUAACUCGGGGUCAAUAU	1	no match			
3351	UUGUUUCAGAGCGUUGGCCAGUUUCC	1	0	transposon	Low_complexity	F-element
3352	UCUAUCGAAGAAUCGCGGAUGUGAC	1	0	no annotation		
3353	UGUUUCAUCGUUAGACGGCUCGGGAUGAAAUCUCAUCGUCGUAGA	1	no match			
3354	AUUGACCGAGUUACAGACGGGCGGUUGUUCGUACGAACGGAUGAG	1	no match			
3355	CAGCACACAGAGGAGAGCAGAAGAGAGCA	1	no match			
3356	CAGCACACAGAGGAGAGCAGAUGAAGC	1	no match			
3357	UAAAAAGCGCUUUGCCAUGCUGGAC	1	0	coding gene		CG10843-RA|Cyp4p3
3358	AAAUUAUAAGGAUGAUGAUACCGAUU	1	0	no annotation		
3359	UUUCUGGAGAACUCGAGCCGCGUAG	1	no match			
3360	UCUCAUCCUCGUAGAACAGCCCGA	1	no match			
3361	UCUCCAUCGUCGUAGAACAAGCCCGACUGUAACCUCGGGUCAAU	1	no match			
3362	AUAUCAUUCACGCCACGCUCCGACC	1	0	Su(Ste)		Su(Ste)
3363	UCUCAUCGUCGUAGAAGAAGCCGAU	1	no match			
3364	CAAUGUACUGUUUGUAGGUGUACUGACCCAG	1	no match			
3365	AUCGUCUCGGGGAUGAAAGGACCUCAACUGUGAGGGGAUCCUCAA	1	no match			
3366	CCCCUGAUGAAGAGAGAUGCCCAGUA	1	no match			
3367	GCAUAUCAUUCACGCCACGCUCCGACCU	1	0	Su(Ste)		Su(Ste)
3368	UGUUUCACCUUAGACGGCUCGGG	1	no match			
3369	CUGACGAGAACUCGAGCCGCGUAGC	1	0	no annotation		
3370	UCUCGGGGUGAAAGGAUCCUCAA	1	no match			
3371	UCCUUGCAGGUUGCUGAUCGAUGACCACUGUA	1	no match			
3372	UUCACGCCACGCUCCGACCUGAU	1	0	Su(Ste)		Su(Ste)
3373	UAAAUGUCGUCCCGGAUACAUUGUC	1	0	transposon	LINE	Rt1b
3374	UCCGUUCCCUUGCGAUGCUCGAUU	1	1	transposon	LTR	G4
3375	UCCGAAGUCUGAUACACAGCUGGACG	1	0	repeat		trf
3376	UCUCAUCGCCUAGAACAAGCCCGA	1	no match			
3377	UCUAUGCUUGUUGCUACAGGGCCAA	1	0	transposon	LTR	invader2
3378	UAAUGCCAGAAUUUAUGGGG	1	2	coding gene		CG8256-RD|Gpo-1
3379	AUCUCAUCGUCGUAGAACAAGCUCGA	1	2	Su(Ste)		Su(Ste)
3380	GUCUCGGGGAUGAAAGGAUCCUCAA	1	no match			
3381	UGAGCGAGUGAGAGAGAAGCAA	1	0	coding gene		CG3412-RA|slmb
3382	AACUGAAGGAGUACGAAUGAUCG	1	0	transposon	LINE	R1-element
3383	UGAGAAAGGACAGCCCCGAGGGCCAA	1	0	transposon	LTR	297
3384	GUCUUCUAUCGUCGUAGAACAAGCCCGUCUGUAACCUCGGGUCAA	1	no match			
3385	UAGAUCUUCACCGUUGACCUGCCCCGAC	1	2	Su(Ste)		Su(Ste)
3386	GCCGGUCGAGAGAUAUCGGAAGAGGUU	1	no match			
3387	UUCCGAGGAACCAACUGAUCCAGCUG	1	1	Su(Ste)		Su(Ste)
3388	UGUUUCAUCGUAAGACGGCUCGGGC	1	1	AT-chX		AT-chX-B
3389	UGAUUAAAGCCUUCGCGUCUUCCACGAC	1	0	AT-chX		AT-chX-A
3390	UCUCAUCGUCGUAGAAGAAGCUGA	1	no match			
3391	AAUUCCCAAUGCUUUGCGACGGCGCA	1	no match			
3392	UCUCAUCGUCGUAGAACAAGCCCGCUC	1	no match			
3393	AAUGGAGGGGAUCCUCAA	1	2	no annotation		
3394	CGUUGACCAAUUUCCGAACAUAAAA	1	2	transposon	LTR	Dm88
3395	UCGUCAUCCGUCGUAGAACAAGCCCG	1	no match			
3396	UCUCAUCGCCGUAGAACAACGCCCCGA	1	no match			
3397	AAAACCUAGUUACUUACGGGCUCGACA	1	0	Su(Ste)		Su(Ste)
3398	UGUUUCAUCGUUAGACGGCUCGGGCCU	1	2	AT-chX		AT-chX-B
3399	AUCGUCGUCGGGAUAGAAACCUCAUCGUCGUAGAACAAGCCCGA	1	no match			
3400	UGUUUCAACGUUAGACGGCUCGGG	1	1	AT-chX		AT-chX-B
3401	UGCCCGAUGAGAAGAAUGGCCCGUAA	1	0	transposon	LINE	RR49783

3402	AUUGACUGAACGAUUCCUUUUGGA	1	0	transposon	LINE	G5
3403	UAUCUAGAGACCGAAGAUAGUUGAG	1	0	transposon	LTR	opus
3404	UACCCAACCGAAGACACCGGCCCAA	1	2	no annotation		
3405	ACCACCUUGACGGAGCUCACGUACC	1	0	transposon	LINE	R1-element
3406	UCUCAUCGUCGUAAAAGAAGCCCGA	1	2	Su(Ste)		Su(Ste)
3407	UCUAAUCGUCAUAACAAGCCCGA	1	no match			
3408	UGCAAUGAAGUGUGGCCCUAGUGCU	1	0	transposon	Low_complexity	F-element
3409	AAUUGAAAUGAAGCCGGCAAUGCCCA	1	1	transposon	LINE	baggins
3410	ACAACAGAGGAGAGCAGAUGAGAGCA	1	no match			
3411	GAUUCUUGAGAAUUUGGACGCCGUU	1	0	transposon	LINE	Cr1a
3412	GCUCCGAAAACAAGGCCCGUCGCU	1	0	transposon	LINE	G6
3413	UCUCAUCGUCGUAGAACAAGCCCGACUGUAACCUCGGGUCAA	1	no match			
3414	AAAUGUUUUCAUCGUUAGACGGCUCGGG	1	no match			
3415	CAUAUCAUUCACACCACGCUCCGA	1	0	Su(Ste)		Su(Ste)
3416	UCGCAUCGUCGUAGAACAAGCCCCGCUGUAACUCGGGGUCAAUAU	1	no match			
3417	UCUCAUCGUCGUAGAACAAGCCCGUAA	1	2	Su(Ste)		Su(Ste)
3418	UACUCAUCGUCGUAGAACAAGCCCGAGG	1	1	Su(Ste)		Su(Ste)
3419	UCUCAUCGUCGUAUAACAACCCCGA	1	2	Su(Ste)		Su(Ste)
3420	UCCAUUGGAUGAUCACAGCACAACUGACC	1	0	coding gene		CG32217-RA|Su(Tpl)
3421	UCAAAAUACUUAUGCGAGCACCCGUUC	1	0	transposon	LTR	opus
3422	CUGUUUCAUCGUUAGACGGCUCCGGCC	1	2	AT-chX		AT-chX-B
3423	UCACCCUGAGUGCGACCAACGGCA	1	0	transposon	LTR	opus
3424	CCCGAAAACACGAGAGCCGAGAACUGUAACUCCGGGUCAAU	1	no match			
3425	CAACCGUGAGGGGAUCCUCAA	1	no match			
3426	AAAACCACCGUUUCCCGUCCGAUCAA	1	no match			
3427	CAGCACGACAGAAGGAGAGCAGAUGAAGAGCCA	1	no match			
3428	ACCACCCUGAACGCGCCCGAUCUC	1	no match			
3429	AAAUAGACCUGGGCUUGUCAGCGGA	1	0	transposon	LTR	GATE
3430	CUGUAACUCGGGAUAAGGAUUGGCUCU	1	no match			
3431	UAGAUCUUCACCGUUGACAGCCCCGA	1	no match			
3432	GAAUCCGAAGUUUGAACGUGGGAACG	1	0	transposon	LTR	Idefix
3433	AUCGUCUCGGGACUGAAACUCUCAUCCGUUCGUACGAAACAACGC	1	no match			
3434	UCAUCUUCUUCUCGACGUUAGACCA	1	no match			
3435	UUAAGAAGAUGUCCGGGAAGCUG	1	0	Su(Ste)		Su(Ste)
3436	UAGAUCUUCACCGUUGACUUUCCCCA	1	1	Su(Ste)		Su(Ste)
3437	ACCUCAACUGUGAGGGAUCCCUCAAA	1	no match			
3438	UGUACUGACCCAGUCGGAUCUAUUGUAC	1	0	transposon	LINE	baggins
3439	UCCUCAUCGUCGUAGAACAAGCCCCG	1	no match			
3440	UUUCCGCGAGAAAAACCCGAUGGGA	1	0	no annotation		
3441	ACCACCACCUGUACUCUGACUCCGAGG	1	0	transposon	LINE	R1-element
3442	AACCUGACCGGUAGGCUAAACAAGU	1	0	coding gene		CG2033-RA|RpS15Aa
3443	UCUCAUCGUCUUAGAACAAACCCGA	1	2	Su(Ste)		Su(Ste)
3444	AUCUUAUCGUCGUAGAACAAGCCCGACUAGUAACGUCGGGUCAAU	1	no match			
3445	CUCCAUGCAGCAGAAGCGACAGAGU	1	0	transposon	LINE	baggins
3446	CUGUAACUCGGGGUCAAAUAAAAAAAAAAA	1	no match			
3447	AAUCCUCAACUGUGAGGGAUCCUCAACUGUAACGUCGGGUCAAU	1	no match			
3448	GCGUCGCUACCAGACGGGAUUACUCUG	1	0	transposon	LTR	ninja-Dsim-like
3449	UCAGCAGACGAAGACCGGCCACCC	1	0	transposon	DNA	hopper2
3450	UACAAGAUGCAUACGAUGUGGCCUGUAACUCGGGGUCAAU	1	no match			
3451	CUGUAACUCGGGGUCAAUGGGUCAAU	1	no match			
3452	GCCGAGAUAUGCUGCGAGUUGCCAA	1	0	transposon	Simple_repeat	I
3453	UCUCAUCGUCGUAGAACUAGCCCGGA	1	2	Su(Ste)		Su(Ste)
3454	UGGAUACCGAGAAUGACAAUGGGUUGA	1	no match			
3455	ACUGUGAGGGUAUCCUCAA	1	no match			
3456	UGUUUCAUCGCUAGACGGCUCGG	1	1	AT-chX		AT-chX-B
3457	UCUCACUCGUCGUAGAACAAGCCCGUA	1	no match			
3458	UGAAAGGAAUGCCCGACGUGCCAA	1	0	transposon	LTR	ninja-Dsim-like
3459	GGGGACUGUGAGGGGAUCCUCAA	1	no match			
3460	UACUGUUUGUAGGUGUACUGGCCC	1	0	transposon	LINE	baggins
3461	CGUUUCAUCGUUAGACUGCUCGGG	1	2	AT-chX		AT-chX-B
3462	UGUUUCAUCGUUAUACGGCUCGG	1	1	AT-chX		AT-chX-B
3463	UGUUUCAUCGUUGGACGGCUCGG	1	1	AT-chX		AT-chX-B
3464	ACCUCAACUGUGAGGAUCCUCAA	1	no match			
3465	AUUCUCACUCGUCGUAGAACUAAGCCCCGA	1	no match			
3466	GUGAGGUCUCCGGACGUGAUCACUGUGAC	1	0	ncRNA		rRNA
3467	CUCAACUGUGAGGGGAUCCUCUAA	1	no match			
3468	AUCGUCUCGGGAGUGAAAUCUCUAUCGUCGUAGAACAAGCCCGA	1	no match			
3469	AGGGAAUCCUCAACUGUGAAAA	1	no match			
3470	ACCCCUUAGGCGAGUUAGCGAAGCUGGCC	1	no match			
3471	GAUCUUCACCUUUGACUUGGCCCA	1	1	Su(Ste)		Su(Ste)
3472	UAGACCUUCACCGUUGACUUGGCCC	1	1	Su(Ste)		Su(Ste)
3473	AUCGUCUCGGGGAUGAAAAUCAACUGUGAGGGGAUCCUCAA	1	no match			
3474	AUCGUCUCGGGACUGAAACUCUCAUCGUCGUAGAACAACGCCCGU	1	no match			
3475	UAAGACCCAAUCUAUGUGCUUAACAU	1	0	transposon	LTR	gtwin
3476	CAAGAUAUGACCAAAGCAAAGACACCUG	1	2	transposon	LINE	Cr1a
3477	UCUCGAUCGUCGUAGAACAAGCCCGACGUGUAACGUCGGGUCAAU	1	no match			
3478	UCUCAUCGUCGUAGACAAGCCCG	1	no match			
3479	UAUCAUGUGAAAUUUGUGAACCCUGUU	1	0	transposon	LTR	copia
3480	GUAAUUGGGCCCGAACCUUCUGGUG	1	no match			
3481	UCUCAUCGUCGUAGCAAGCCCGA	1	no match			
3482	UCGAACACCGGCUUCAGGAUCACGUC	1	0	Su(Ste)		Su(Ste)
3483	CCCAAACCCAGACCGGCCACCCCUU	1	0	transposon	LINE	G2
3484	CUCAAUGUGAGGGGAUCCUCAA	1	no match			

3485	UCUCAUCGUCGUAGAAUAAGCCCGACGG	1	no match			
3486	UCUCAUCGUAGAACAAGCCGA	1	no match			
3487	UUCCUGUUGAUGUGAGGGCCCCAAGUC	1	0	transposon	LINE	G2
3488	UACAAUCUCCGAGCACGACCCCCGAUCU	1	no match			
3489	CCUCAUCGUCGUAGAACAAGCCCAA	1	2	Su(Ste)		Su(Ste)
3490	UGAUCCGAUAUUCUUCUGCUGCAACC	1	no match			
3491	AUAGAUCUUCACCGUGGACUAGCCCCC	1	no match			
3492	UAAGGAUAUCCCUGUUGCCCGCCCGA	1	1	transposon	LINE	G2
3493	GAUAACGAGACCGGAUAUUCCUGC	1	0	coding gene		CG9010-RA|CG9010
3494	GGCUCAACUGUGAGGGGAAUCCUCAA	1	no match			
3495	AUUCGAAAGGAAGGAGGCCACAAUCU	1	1	coding gene		CG14560-RA|msopa
3496	AUAACUCCGAGGGAGGGUGAAACCAUC	1	1	transposon	LINE	R1-element
3497	GUGUUUCAUCGUUAGACGGCUCGGGCCU	1	no match			
3498	UACCUUAUUAAUGGAUUGGCUGA	1	0	transposon	LTR	Stalker
3499	GCUUAUCCUUUGCUUGAUGAUUCG	1	0	ncRNA		rRNA
3500	UCUCAUCGUUGUAGAACAAGCCCGACUA	1	no match			
3501	CUAGAUCUUCCACCGUUGACUACGCCCCUA	1	no match			
3502	UCCUGUUGAUGUGAGGGCCCCAAGUCUGUAACUCGGUCAAU	1	no match			
3503	UCCUGACGAAGAGCGCAAGAAGAG	1	0	transposon	LTR	opus
3504	CAAGAUAUGACCAAAGCCAAGACAC	1	1	transposon	LINE	Cr1a
3505	AAAAACCUGAAGAUGAUCCUGAGUAAAA	1	no match			
3506	UCAUAUGAUCUGAGAAUAUUUUC	1	2	transposon	LINE	Cr1a
3507	AUCUCAUCGUCGCAGAACAAGCCCGA	1	2	Su(Ste)		Su(Ste)
3508	AUUUUCAUCGUUAGACGGCUCGGG	1	2	AT-chX		AT-chX-B
3509	UCUCACCGUCGUAGAACAAGCCCGAG	1	1	Su(Ste)		Su(Ste)
3510	CGAAAUCUCAUCGUCGUAGAACAAGUCCCGA	1	no match			
3511	CCCCAACUCCUGGAGCGGUUCGGC	1	0	coding gene		CG1851-RA|Ady43A
3512	UACUUACCCGUUUACAGAAGUGGA	1	0	transposon	LTR	accord
3513	UUUGUUGAUCAGUUUACACAUUAUU	1	0	transposon	LINE	Cr1a
3514	AAUCAACUGUGAGGGGAUCCUCAA	1	no match			
3515	UCAACUGGGAUCCUCAA	1	2	coding gene		CG2216-RA|Fer1HCH
3516	AGAAAUAGAUCUUCACCGUUGACUAGCCCC	1	no match			
3517	UGUAACUCGGGGUCAAUAUCGUCUCGGGGAUGAAAGCACACAGAG	1	no match			
3518	CGCACGACAGAGGAGAGCAGAUGAGAGCA	1	no match			
3519	UAUGCCCGUACUCUGUGGUGACCG	1	no match			
3520	UCUCAUUUCGUCGUAGAACAAGCCCGA	1	no match			
3521	UCCGAAGUCUGAUACACAGCUGGAC	1	0	repeat		trf
3522	CCUAAGUUGCACUUCUGAUGAUGGC	1	0	coding gene		CG1558-RA|Kmn1
3523	AAGGCGAAAAGAACCCCGGGAGG	1	no match			
3524	UCUCAUCUCGUAGAACAAGCCCGAG	1	no match			
3525	UGGACUUUUUUCCGAUGAGCCAGC	1	1	transposon	LINE	G2
3526	UCUUCACCGUUGACUUGCCCCACA	1	0	Su(Ste)		Su(Ste)
3527	AAAACUGCACCCUGAGUAUGGCAUUA	1	0	transposon	LINE	BS
3528	AAACCUCAACUGUCAGGAUCCUCAA	1	no match			
3529	GAUGGAAAUGAAGAAAGUACA	1	2	coding gene		CG1647-RA|CG1647
3530	AUCUCAUCUUCGUAGAACUAGCCCGA	1	2	Su(Ste)		Su(Ste)
3531	UAACUAUGUGACCAAGAAAGUUAGCU	1	0	transposon	LTR	297
3532	GGAUCCAACUAUGAGGGGAUCCUCAA	1	no match			
3533	UGUUUCAUCGUUAGACGGACUCGGGCC	1	no match			
3534	AUGGUUUCAUCGUUAGACGGCUCGGGCCGUGUAACGUCGGGUCAA	1	no match			
3535	UGCACCCUGGAUCGGACUCGCUAGUCA	1	0	transposon	LINE	INE-1
3536	UGUUUCAUCGUUGACGGCUCGGG	1	no match			
3537	AGAAAAAAAAAAAAAGGGAUCCUCAA	1	no match			
3538	GGUGCAUAUCAUUCACGCCACGCUCCG	1	0	Su(Ste)		Su(Ste)
3539	GGCCACUCCCGCCAACUGGACCCUGG	1	no match			
3540	GUUUAAGGCAACUCCCUGGACGCACGA	1	0	transposon	LTR	ninja-Dsim-like
3541	AUUGAUACGAAUGAGACCCGCACGUAA	1	no match			
3542	UCUCAUCGUCGUAGAACAAGCUGA	1	no match			
3543	GCACACAUAGGAGAGCAGAUGAGAGCG	1	no match			
3544	CGAAAUCCUCAUCGUCGUAGAACAAGCCCGACU	1	no match			
3545	CUCAUCGUCGUUGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
3546	UUUAUACCUGAGCCAGUUUUGUUUUG	1	no match			
3547	UCCGGGAAGAAAAACAAUGAUGGCCAAGU	1	no match			
3548	UACUCGAGUAAGCGACUCAGUCAUG	1	0	transposon	LTR	gypsy12
3549	UCUCAUCAUCGUAUAACAAGCCCGAU	1	1	Su(Ste)		Su(Ste)
3550	UCUCAUCGUCGUAGAACAAGCGCGAG	1	1	Su(Ste)		Su(Ste)
3551	GGAUCCUCAACUGUGAGGUCUCAA	1	no match			
3552	CACACAUCGCUGAGGCCCACUGGCA	1	0	Su(Ste)		Su(Ste)
3553	UUCAAGUGCCAUCCUAAGCAUGAACCCG	1	0	transposon	LTR	GATE
3554	AUCUCAUCGUCGUAGAACAAGCCCGACUGUAACGUCGGGUCAAU	1	no match			
3555	AAAAAACCUGAAGAUGAUCCUGAGUCG	1	2	transposon	LINE	Helena
3556	GGCAGACGCAAGUGGACUCGUUG	1	0	transposon	LTR	micropia
3557	AAAGCUCCCCAGGUUGGUCGGUAAGUC	1	no match			
3558	UCUCCAUCGUCGUAGAACUAGCCCGA	1	no match			
3559	UGCAACCCCAACGACCACCCGCUGUU	1	no match			
3560	UGUAUUUAUACUGUUUGGCCGAUGA	1	0	transposon	LTR	flea
3561	UUCUUCUCGACGUUGACCAGCCUCC	1	0	transposon	LTR	ninja-Dsim-like
3562	UGAUACCGAGAAUGACAAUGGCUGAA	1	1	transposon	LTR	gypsy3
3563	GCUGAGAGUGAUGAUCAGAAAA	1	1	coding gene		CG4076-RA|Nufip
3564	UGGACUUUUUCCGAUGAGCCAGUGA	1	no match			
3565	CCGAACUUGAGAUCGGCCCUGGACGACC	1	1	coding gene		CG42398-RA|SteXh:CG42398
3566	UUUCUCAUCGUCGUCGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
3567	UAAACUGUGAGGGGAUCCUCAA	1	no match			

3568	GAUGUUUCAUCGUUAGACGGCUCGG	1	2	AT-chX		AT-chX-B
3569	AUCGUCUCCGGGAUCGAAAUGUUUCAUCGUUAGACGGCCUCGGGC	1	no match			
3570	AUCGUCUCGGGAGUGAAAGCACACAGAGGAGAGCAGAUGAGAGCA	1	no match			
3571	UGUUUCAUCGUUACACGGCUCGG	1	1	AT-chX		AT-chX-B
3572	UGUUUCAUCGUUAGACUGCUCGGC	1	2	AT-chX		AT-chX-B
3573	AUCGUCUCGGGACUGAAAUGUUUCAUCGUUAGACCGGCUCGGGC	1	no match			
3574	CGUUGACCAUUCCGAACAUACGC	1	no match			
3575	AAGACCGAUGUGAAAUCCCCGAGCUU	1	no match			
3576	UUAUGACCCCUGGUACUGAAAUGCA	1	1	transposon	DNA	Bari1
3577	UCGCCCAGAGCGGUUUCCCUGUGUUUCA	1	0	no annotation		
3578	UCGGCAUUACUCCGAUGGUGGCUCC	1	1	transposon	LINE	R1-element
3579	GCCCGAUCUGGUCGAGAACUUCG	1	0	coding gene		CG3644-RA|bic
3580	CAGCACACACGAGGAGAGCAGAUGAGAGCA	1	no match			
3581	AAUCAUCGUCGUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
3582	AUAUCUUCACCGUUGACUAGCCCC	1	1	Su(Ste)		Su(Ste)
3583	AUAAGAUUGGACCCAGUACACUGCCUU	1	0	transposon	LINE	Doc
3584	GUCUCGGGGAUAAAA	1	1	no annotation		
3585	UUCAAACACAGUUCGAACCCCCCCGUG	1	no match			
3586	UCUGGUGGCGAUAGCGAAGAGGACACAC	1	no match			
3587	AUGUUUCAUCGUUAGACGGCUCGGGCU	1	2	AT-chX		AT-chX-B
3588	UACGACUUGAGAGAAACCAGGUG	1	no match			
3589	UCUCUUACUCGUCGUAGAACAAGCCG	1	no match			
3590	AUCUCAUCGUCGUAUCAAGCCCGA	1	no match			
3591	AGAAAAUAAAUGCCAGUAAUUUUUAGUUGGGGUUC	1	no match			
3592	UAUUAUUAAGAUGGGCUUGAAGG	1	0	transposon	LINE	baggins
3593	CAGCCUACGAACAGAGAGAAGACCC	1	no match			
3594	UCUCAUCAUCAUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
3595	UCUCAUCGUCGUAGAACAAGCCCCGCUGUAACCUCGGGGUCAAU	1	no match			
3596	UUAAAUAUGACCCGAUUAAAUGAGAA	1	1	no annotation		
3597	AUCGUCUCGGGGAUGAAACUGUGAGGGGGAUCCUCAA	1	no match			
3598	GGAAUACCGGGUGCUGUAGGCUUU	1	no match			
3599	GCAACGAGAUGGCGACCUUGGAGAAA	1	0	transposon	LTR	297
3600	UGUUUUCAUCGUUAGACGGCUCUCGGC	1	no match			
3601	GAUCCUCAACUGUGAGAGGAUCCUCAA	1	no match			
3602	UCUCGAUCGUCGUAGAACAAGCCCGAGG	1	no match			
3603	CAAGCUGACGCUAUAUGCCCCUGGU	1	0	transposon	LTR	invader6
3604	UUCGCAGAGAGAUAUGGUAGAUGG	1	0	ncRNA		rRNA
3605	UAAACGAGCAAGGAAGAGUCGGAA	1	0	transposon	LINE	Cr1a
3606	GGGCAACUUUGAGGGGAUCCUCAA	1	no match			
3607	UGCCCGAUGAGAAGAAUGGCCC	1	0	transposon	LINE	RR49783
3608	UUGUACACGAGCGCAACGAAAAUACG	1	0	transposon	LINE	R1-element
3609	CAAGAUAUGACCAACGCAAAGACUA	1	2	transposon	DNA	1360
3610	UAGAUCUCCACCGUUGACUUGCCCC	1	1	Su(Ste)		Su(Ste)
3611	AACUCAACUGUGAGGAUCCUCAA	1	no match			
3612	UUUCUUUGAAGAUGGCCCACAUAAAGA	1	0	repeat		trf
3613	UAAAUGCCAGAAUUUUAUUGGG	1	0	transposon	LINE	baggins
3614	AUGGUUGGACUGGGAAUGGCAUC	1	0	no annotation		
3615	CUCACUGUGAGGGGGAUCCUCAA	1	no match			
3616	UCAACUAUCUACGAGACCACCCAAC	1	no match			
3617	CCCCGUACGUGAAAUGACCUGUGUG	1	no match			
3618	UGUUUCAUACGUUAGACGGCUCGGGC	1	no match			
3619	UCCGCGUAGCACUAAAUCACCGAUU	1	1	transposon	LINE	R1-element
3620	AUCUCAUCGUCGUAGAACAAGCCCGACUGUAACUCGGGGUCAAUU	1	no match			
3621	UGUUUCAUCGUUAGACGGCUCGGGCCUG	1	no match			
3622	UAAUAGUGGGUCCGAGGAAUCAAUA	1	0	transposon	DNA	transib2
3623	UACCGGUAUAGCACACUUUUGUAUC	1	0	coding gene		CG41580-RA|CG41580
3624	UCAAAUCCGUAGCUGCAAAUUAAUU	1	0	transposon	DNA	transib2
3625	UGGACUUUUUCCGUAUGAGCCAGUG	1	2	transposon	LINE	G2
3626	UCUGGUUUCAUCGUUAGACGGCUCGGGCG	1	no match			
3627	AUCUCAUCGUCGUAGAACAAGUCCGA	1	2	Su(Ste)		Su(Ste)
3628	UCGAGAGGUCCCCGGUUCAAACCC	1	0	ncRNA		tRNA
3629	UCCUCAUCGUCGCAGAACAACGCCCGA	1	no match			
3630	AUCCUCAACCGUGAGGCAUCCUCAA	1	no match			
3631	UGUUUCCAUCGUUAGACGGCUCGGG	1	no match			
3632	AUCCUCAUCGUCGUAGAACAAGCCCGA	1	2	Su(Ste)		Su(Ste)
3633	UGUUUCAUCGUUAAACGGCUCGGG	1	1	AT-chX		AT-chX-B
3634	AUCGUCGUCGGGAUGAAAUCUCAUCGUCGUAGAACAAGCCCGA	1	no match			
3635	UGUUUCAUCGUUAGACGGCUGGG	1	1	AT-chX		AT-chX-B
3636	UGUUUCAUCGUUUAGACGGCUCGGGCG	1	no match			
3637	GGAUCCUCAACUGUGAGGGGAUUCUCAA	1	no match			
3638	UCUCAUCGUCGUAGAACAUGCCCG	1	1	Su(Ste)		Su(Ste)
3639	GCCUCAGAGACACUGACCGUAGCAC	1	1	transposon	LINE	Rt1b
3640	UCCAGUCUCCCGACUACUAGAUACCCG	1	2	transposon	LINE	INE-1
3641	CCAAGAUAUGACCUAAGCAAAGACAC	1	2	transposon	LINE	Cr1a
3642	AAGGUGGAAAUGAAGGGAUCCUCAA	1	no match			
3643	UUCGAAGCCCGGGUAUUCUGAAGCCAA	1	no match			
3644	GGAUCCUCAACUGUGACUCAA	1	no match			
3645	CAAUCGCCGCUAUGCUUACUUCAGCUA	1	no match			
3646	UAGAUCUUCACCGUUGACUUCGGCCC	1	no match			
3647	GAAAUUCUCAUCGUCGUAGAACAAGCCCG	1	no match			
3648	UACAAUCUCCGAGCAAGACCCCGAUUC	1	2	transposon	LINE	Circe
3649	CAGGCACACAGAGGAGAGCGAGAGUGAGGACA	1	no match			
3650	AAAAGGUGGAAAAUGAAGAAAGUACA	1	no match			

3651	UCCAUCUCUGAAGUUUCAUGUUUGGC	1	0	transposon	LTR	Quasimodo
3652	UAAUGCCCGAAGGUUGCCUCU	1	no match			
3653	UCUCAUCUUCGUAGAACAAGCCCCGACUGUAACUCGGGGUCAAU	1	no match			
3654	UUAAUACCCAGGUGAUUACCACUAU	1	1	Su(Ste)		Su(Ste)
3655	UAUAUAGUGCCGAGAACAAUAUGCC	1	0	transposon	LINE	G5
3656	UCUCAUCGUCGUAGAACAGGCCCUA	1	2	Su(Ste)		Su(Ste)
3657	UCAUCUUCUUCUCGACGUCACCA	1	no match			
3658	UCUCAUCGUCGUAGAACAAGCCCGCU	1	2	Su(Ste)		Su(Ste)
3659	UUCGGGGUUUUGGCACUUUGCG	1	no match			
3660	GCAUGAUGUGAUAGCCGUUACACU	1	no match			
3661	ACAUCCUCAACUGUGAGGAAA	1	no match			
3662	AACUGGUGAGGUGAUCCUCAA	1	no match			
3663	AGUUCAAGCCCGGGUAUUCUGAAGCC	1	0	repeat		trf
3664	UCGUCAUCGUCGUAGAACAAGCGUCGA	1	no match			
3665	UCGUCAUCGUCGUAGAACAAGCCCGGU	1	no match			
3666	UCGGACACCGAACCGGGAAGGCGA	1	0	transposon	LTR	Tirant
3667	UUAUAAGGAUGAUGAUACCGAUGAAA	1	1	no annotation		
3668	AAGAUAUGACCAGAGCAAAGCAAC	1	0	Su(Ste)		Su(Ste)
3669	AGGAUCAACUGUGAGGGGAUCCUCAA	1	no match			
3670	AUCUCAUCGUCGUAGAACUAGCCCCGA	1	no match			
3671	UCCGAACGUACCACGGUCCCCCAGU	1	1	transposon	Low_complexity	F-element
3672	UCUCAUUCGUCGUAGAACAGCCCGA	1	no match			
3673	UCUCAUCGUCGUAGAAGAAGCCCGACUGUAACCUCGGGUCAAU	1	no match			
3674	UCUCAUCGUCGUAGAACAACCCCCGAA	1	no match			
3675	UGUUUCAUCGUUAGACGGCUCGGGCUC	1	2	AT-chX		AT-chX-B
3676	UACUAGCUGGCCGAGGAUCCCGAUU	1	no match			
3677	GCGACACAGAGGAGAGCAGAUGAGAGCA	1	no match			
3678	UCGAUUCCGGCAACGAGAUGACCACG	1	0	coding gene		CG5547-RA|Pect
3679	UGUUUCAUCUUUAGACGGGCUCGGGC	1	no match			
3680	AUUGACCGACGUACAGACGGGCGGUUGUUACGUACCGAACGGACU	1	no match			
3681	UAACAUUGGACCCAGUACACUGCC	1	1	transposon	LINE	Doc
3682	AACUUCGGGAUAAGGAUUGGCUCUGA	1	0	ncRNA		rRNA
3683	UCCGGGAGAAAACAAUGAUGGCCAAA	1	1	coding gene		CG12717-RA|CG12717
3684	AGGGGAUCCUCAACUGUGCCUCAA	1	no match			
3685	GUGAUGUCAACGUUUGAUCCCUG	1	2	transposon	LTR	ninja-Dsim-like
3686	UUUCUCAUCGUCGUAGAACUAGCCCCGA	1	no match			
3687	AGUCUCAUCGUCGUAGAACUAGCCCGA	1	2	Su(Ste)		Su(Ste)
3688	UCCUUCGGGAUGAUUUGUACCGUAG	1	1	transposon	Low_complexity	F-element
3689	UUAAUACCUCGAGUACGUCUGCUCUC	1	0	transposon	LTR	Max
3690	UGUCCGGACAGCUGGGCCCGAA	1	0	Su(Ste)		Su(Ste)
3691	CAACUAGUGAGGGGAUCCUCAA	1	no match			
3692	UAUACCGGACGGUAUGCAACCCUUAUAA	1	0	transposon	LTR	gypsy12
3693	UGUUUCAUCGUUAGACGGCUCGGGGCCUGUAACUCGGGGUCAAUA	1	no match			
3694	UCAACUGUGAGGGGGAUCCUCAAC	1	no match			
3695	CUCUCAUCGUCGUACGAACAACGCCCGAC	1	no match			
3696	UUUCAUCGUCGUAGAACAAGCCCGUA	1	2	Su(Ste)		Su(Ste)
3697	UCGUCAUACGUCGUAGGAACAAGCCCGA	1	no match			
3698	GAUUCCGGAGAGGGAGCCUGAGAAAC	1	0	ncRNA		rRNA
3699	CUGGUUGAACGUGUCCUGAAAGUA	1	0	Su(Ste)		Su(Ste)
3700	UGUAGGUGUACUGACCCAGUCGGA	1	0	transposon	LINE	baggins
3701	GAUCCCCCCUGAAGUCCAAUGCACUG	1	no match			
3702	UGAUUCCGAUCUUGGACAUUCCCGA	1	0	transposon	LTR	roo
3703	GCAAUCAUCCAAGGCUGAGAUCCGG	1	0	transposon	LTR	mdg3
3704	UCGUCAUCGUCGUAGAACAGGCCCG	1	no match			
3705	UCGUCAUCGUCGUAGAACAAGCCCGU	1	no match			
3706	UCGUCAUCGUCGUAGAACAAGCCCGGAA	1	no match			
3707	UUUGGCCAGGUUAUGGAGAAAACCCCGU	1	0	repeat		SAR_DM
3708	UAGAUCUUCACCGUUGACUAGCCCCG	1	1	Su(Ste)		Su(Ste)
3709	UAACACUUCCCAGAAUGCACACCACCCA	1	0	transposon	LINE	Cr1a
3710	GUAGUGACCCCUGGUACUGAAAUGCGA	1	no match			
3711	CCUCAACGUGAGGGAUCCUCAA	1	no match			
3712	AUGUACUACCCCCUUCAUGACUCAAAG	1	no match			
3713	AAGAUAUGACCAAAGCAAAGACAC	1	0	transposon	LINE	Cr1a
3714	UCUACUGACUUCAGUAAGCCCCGUAC	1	0	transposon	LINE	baggins
3715	UGUUUCAUCGUUAGACGGGCUCGGGGC	1	no match			
3716	CGACAACGUUGACCACUGUGCCGGGA	1	2	coding gene		CG8340-RA|128up
3717	CAAGCACCUCGAGAACAGUUCCCUUAG	1	0	coding gene		CG17078-RA|CG17078
3718	CGAAAUGUUUCAUCGUUAGACGCUCGGG	1	no match			
3719	AUGGCACGAAAGACGAACCCGAUUU	1	no match			
3720	UAGACUCGAAAAUUCGUUUUAUG	1	0	Su(Ste)		Su(Ste)
3721	AAACCUGAAGAUGAUCCUGAGUCGCAA	1	1	transposon	LINE	Helena
3722	GUCUUGGGAUGAAAAUCCUCAA	1	no match			
3723	UGUUUCUAUCGUUAGACGGCUCGGCC	1	no match			
3724	CCUGCUGACGGGCCCGAAAACACGA	1	no match			
3725	UACGAAUCGUCCGAGCAAGACCCCGAGUC	1	no match			
3726	UUAAGGCCGAGAGCUGAUGACGAGU	1	no match			
3727	UAGAUCUUCACCGUUGACUUGCUCCACUU	1	no match			
3728	UAAUGCCAGUAAUUUUUAGUUGGGGUUC	1	no match			
3729	UCUCAUCAUCUUAGAACAAGCCCGAA	1	2	Su(Ste)		Su(Ste)
3730	UGUUGCCAGUUCACUCGUUCACAGA	1	0	Su(Ste)		Su(Ste)
3731	UAAAUGCCAGAAUUUUAUUUGGGUUC	1	1	transposon	LINE	baggins
3732	AAAACCUGAAGAAGAUCCUGAGUCGC	1	1	transposon	LINE	Helena
3733	UUUUUCAUCUAGACGGCUCGGGC	1	no match			

3734	GUCUCAAUCGUCGUAGGGAACAACGCCCGA	1	no match			
3735	AACUUACGGGAUAAGGAUUGGCUC	1	2	ncRNA		rRNA
3736	CGAAACCCAAAGGAUACUCGAAGACACCCACGG	1	no match			
3737	GUGGAGGGGAUCCUCAA	1	2	coding gene		CG10675-RA|CHKov2
3738	AAACCUAGAAGAUGAUCCUGAGUCGCA	1	no match			
3739	UCGUCAUCGUCGUAGAACAACGCCCGA	1	no match			
3740	CUAAGACCCAAUCUAUGUGCUUAACA	1	1	transposon	LTR	gtwin
3741	GAGGACUAGCUGGCCGAGGAUCCGA	1	0	Su(Ste)		Su(Ste)
3742	UCUCAUCGUCGUAGAACAAAGACCCGAC	1	no match			
3743	AUCUUCCGACAAACGAAGAAAUGGCA	1	1	transposon	LTR	diver2
3744	UAGAUCUUCCACCGUUGACUAGCCCCCUCGUAACUCCGGGUCAAU	1	no match			
3745	UCCGAGGAAAAAUUGAUCCAGCAGC	1	no match			
3746	UAGAUCUUCACCGUUGAACUACGCCCC	1	no match			
3747	UUCUGCCCGAUGAGAAGAAUGGCC	1	0	transposon	LINE	RR49783
3748	UUCUCAUCAUCGUAGAACAAGCCCGAU	1	0	Su(Ste)		Su(Ste)
3749	UUAAACAUCCUAUGAGGCUACGGC	1	0	transposon	LINE	Cr1a
3750	UUGAUAGAAACCUAUAGAGACAAGGC	1	0	transposon	LTR	opus
3751	UCAACUGUGACGGGAUCCUCAA	1	no match			
3752	UGUUUCAUCUUUACACGGCUCGGG	1	2	AT-chX		AT-chX-B
3753	AUGAUACCGAGAAUGACAAUGUUUGA	1	2	transposon	LTR	gypsy3
3754	UUCCGAGGAACCAACUGAUCCAGCU	1	1	Su(Ste)		Su(Ste)
3755	UUCCGAAGACAAGUAUACCGUGGCA	1	0	transposon	LTR	micropia
3756	UCUCAUCAUUGUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
3757	AACUGUGAGGUGAGGGGAUCCUCAA	1	no match			
3758	ACCUGAAGAUGAUCCUGAGUCGCAA	1	1	transposon	LINE	Helena
3759	UGGUACUUUUUUCCGAUGAGCCAGUGA	1	2	transposon	LINE	G2
3760	UGUUACAAUGAUCUGAUCGGGUUU	1	0	transposon	LINE	Cr1a
3761	UGUUUCAUCGUUAGACCGGCUCGGG	1	no match			
3762	UAAUGCCGGAAGCGAUCCCACUGCU	1	0	coding gene		CG41580-RA|CG41580
3763	UCUCAUCGUCGUAGACAAGCCCGAG	1	no match			
3764	GGACCGAACACUUUGAACCAAGUAU	1	0	Su(Ste)		Su(Ste)
3765	UCUCUACUCGUCGUAGAACAAGCCCGA	1	no match			
3766	CAGCGACACAGAGGAGAGCAGGAUGAGA	1	no match			
3767	UCGCUCGAGAGGGACCGUAUGCGGC	1	0	transposon	LTR	ninja-Dsim-like
3768	GAAUAUCGUCUCGGGAGUGAAAUACUCAUCGUCGUAGAACAAGCC	1	no match			
3769	CUUUUCUCAUCGUCGUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
3770	UAUGUUGGCCUGGAUACGCUUUCUGU	1	0	transposon	LINE	G2
3771	AAUAACUGUAAUUAAUAACUGGCCC	1	0	transposon	LINE	HMS-Beagle2
3772	GUUAUCCUUCCCAAGGCGUCUGGAUC	1	0	transposon	LTR	flea
3773	UUGAACGUGUCCUGAAAGUAGUUGG	1	0	Su(Ste)		Su(Ste)
3774	UGUUUCAUCGUUAGACGCUCUGGC	1	no match			
3775	UACGAGAUGGUCACAGGGACCAGGAGA	1	0	transposon	LTR	ninja-Dsim-like
3776	GUCGGGCGCGGUUAGUACUUAGAU	1	0	ncRNA		5SrRNA
3777	AUCGUCUACGGGACUGAAACUGUGAGGGGAUCCUCAA	1	no match			
3778	AUCUCAUCGUCGUAGAAACAAGCCCGAGG	1	no match			
3779	UAGAUCUUCACCGUUUACUUGCCC	1	1	Su(Ste)		Su(Ste)
3780	UCUCAUCUUCGUAGAACACCCCCGA	1	no match			
3781	GCUGAACGUAGUCUCUCGAGAGCGA	1	1	transposon	LTR	invader1
3782	GGUUUCAUCGUUAGACGGCUCGGGC	1	1	AT-chX		AT-chX-B
3783	UGUCAUCGUCGUAGAACAAGCCCCGA	1	no match			
3784	GCCCCUUACUCUGUGGUGACCAACA	1	1	transposon	LINE	baggins
3785	UCCGAUUUCAAGUACCCGGUGAACAU	1	1	repeat		RSP
3786	CAAGAUAUGACCAAAGCAAAGAC	1	0	transposon	LINE	Cr1a
3787	UAUUUAAGAUCAACGAGAGAUGGCC	1	0	transposon	LTR	opus
3788	GACUUCCGAGAAAGCCCGCAGAAUUU	1	0	transposon	LTR	flea
3789	AUGGCAAACGAGAGCUCUUCCGGGUC	1	0	transposon	LINE	R1-element
3790	UAGAUGAGGUAUAAGGUUGGCCC	1	0	transposon	Low_complexity	F-element
3791	CAACCAGAGAUCUAGAGGGCCCA	1	0	no annotation		
3792	AUUCCGAGGAACCAGUUGAUACCAGCC	1	no match			
3793	UGUUACUGAUCUAGAAGUACAAUGU	1	no match			
3794	AGUGAUGUCAACGUUUGAUCCCCUG	1	1	transposon	LTR	ninja-Dsim-like
3795	UUAUACCCAGGUGAUUACCACUAUUAC	1	0	Su(Ste)		Su(Ste)
3796	GCCGAACGUAGUCUCGAGAGCGAGAU	1	no match			
3797	CAAGAUAUGGCCAAAGCAAAGACA	1	1	transposon	LINE	Cr1a
3798	UACUGGUUUCAUCGUUAGACGGCUCGGGCG	1	no match			
3799	CCUGACGAAGAAGAGAUGCCCAG	1	1	transposon	LTR	ninja-Dsim-like
3800	CGGGAUGAAAUAUGAGGGAUCCUCAA	1	no match			
3801	AAAAGAACCCGAUUCCUCUUCCAUGAGA	1	0	transposon	LINE	Doc4
3802	GAUAUGACCAGCAUGGACCUAACAC	1	0	Su(Ste)		Su(Ste)
3803	UCUCAUCUUCGUAGAACAAGCCCGAGG	1	1	Su(Ste)		Su(Ste)
3804	CUGUAACUCGGUCAA	1	1	repeat		SAR_DM
3805	GUAACUUCGGGAUAAGGAUUGGCUC	1	0	ncRNA		rRNA
3806	CGAAAUGUUUCAUCGUUAGACGGCUCGG	1	no match			
3807	CAAGCCCGGGUAUUCUGAACGCCAAGA	1	no match			
3808	UAUUGUACUACCCCUUCAUGACUCAAA	1	0	transposon	LINE	baggins
3809	AUCAACUGUGAGGGGGAUCCUCAA	1	no match			
3810	CCCCGUACUCUGUGGUGACCGACAC	1	1	transposon	LINE	baggins
3811	AUGUUUUCAUCGUUUAGACGGCUCGGG	1	no match			
3812	UCUCAUCGUCGUAGAACAAGCCCGUCUGUAACCUCGGGUCAAU	1	no match			
3813	ACUCAAACUGUGAGGGAAUCCUCAA	1	no match			
3814	UCUCAUCGUCGUAGAACAAGCCCGAAUGUAACUCGGGUAAAU	1	no match			
3815	UAGCUUGCACCAUCACCAUCACCAU	1	no match			
3816	UUCGUUGUGAUUCUGCGCCCUUACGA	1	0	transposon	LINE	Doc

3817	CUCAACUGUGAGGAUCCCCAA	1	no match			
3818	CCCCGUACUCUGUGGUGACCAACA	1	0	transposon	LINE	baggins
3819	ACUGCAUUCGCGUCGAGGCCAGG	1	1	transposon	LINE	R1-element
3820	ACCCAAGUACAAGCCCGGCGAAAGACC	1	0	repeat		trf
3821	UCAAGUGUGACCGUAGUUUGCUACAA	1	0	transposon	LINE	Circe
3822	UACUGAGUCGUUUGCCGAGAGCUGA	1	0	transposon	LINE	Rt1b
3823	GCGGUGCCGAGAUAUGCUGCGAGUUGCCAA	1	2	transposon	Simple_repeat	I
3824	AAUUCAGCGAUUCCGAUUACUAUC	1	no match			
3825	UCUCAUCUUCGUAGAACAAGCC	1	1	Su(Ste)		Su(Ste)
3826	UUCGAACUGGAAGGAGUACGAAUGAGUC	1	no match			
3827	AAAACCUGAAGAUGAUCCUGAGUCUC	1	1	transposon	LINE	Helena
3828	GCCCCCGUACUCUGUGGUGACCAA	1	1	transposon	LINE	baggins
3829	UGGACUUUUUUUCCGAUGAGCCAGUG	1	1	transposon	LINE	G2
3830	CGUCUCGAGAUGAAUGGAUCCUCAA	1	no match			
3831	UGUUUCAUCGUUAGACGGCUCGUG	1	1	AT-chX		AT-chX-B
3832	GAUUAGAUACCCCGGUAGUCCUGG	1	no match			
3833	UAGAUCUUCACCGUUGACUAACCCCAC	1	1	Su(Ste)		Su(Ste)
3834	UAGAUCUUCACCGUUGACUUGGCAC	1	1	Su(Ste)		Su(Ste)
3835	UGAUACCGAGUAUGACAAUGGUUGA	1	1	transposon	LTR	gypsy3
3836	CCAAGAUCAUCAACCCGUGUUAU	1	0	transposon	LINE	Cr1a
3837	UCGACAAUGACCGAUCAGUAGAGAAC	1	0	transposon	LINE	Rt1b
3838	UCGUCAUCAUCGUAGAACAAGCCCGA	1	2	Su(Ste)		Su(Ste)
3839	AAUCCUCAACUGUGAG	1	2	transposon	LINE	TART-B
3840	UAAAAUGCCAGAAUUUUAUUGGGG	1	2	transposon	LINE	baggins
3841	CACACAGAGGAGAGCAGAUGGAGCA	1	no match			
3842	CGAAAUGUUUCAUCGUUAGACGCCUCGGGC	1	no match			
3843	AUGUUUUCAUCGUUGGACGGCUCGGGC	1	no match			
3844	UUAAGUGCCGGGAAGCGAUCCCACUCG	1	no match			
3845	UCUCAUCGUCUUAGAACAAGCCCGC	1	2	Su(Ste)		Su(Ste)
3846	UCUCAUCGUCGUAGAUCAAGCCCGUA	1	2	Su(Ste)		Su(Ste)
3847	UUCUGUUGAUGUGGGGCCCCAAGU	1	0	transposon	LINE	G2
3848	UCGAACCAGAGGCUGGGUACAUCGC	1	0	no annotation		
3849	CCCAAUCAGUUUUAUCUGCAGAGAUG	1	0	transposon	LTR	gypsy8
3850	CAGCCACAGAGGAGAGCAGAUGAACA	1	no match			
3851	UCUCAUCGUCGUAGAACAAGCCCGACUGUAACUCGGGGUCAAUAU	1	no match			
3852	UAAGGGAAGUCGGCAAAAUAGAU	1	1	ncRNA		rRNA
3853	AUCGAGUUGCAGAAUUGAGGACCGC	1	1	transposon	LTR	Quasimodo
3854	AUCCUCAACUGUGAGGGAUCCUAA	1	no match			
3855	GUCUCGGGUGAAAGGAUCCUCAA	1	no match			
3856	UCUCAUCUCGUAGAACAAGCCCGU	1	no match			
3857	UCUCAUCGUCGUAGAACAAGUCCGAGG	1	1	Su(Ste)		Su(Ste)
3858	CAGCACACAGAGGAGAGCAGGAUGAGAGCA	1	no match			
3859	AGAAAGUCUCAUCGUCGUAGAACAAGGCCCGA	1	no match			
3860	CUCAACUGUGAGGUGAUCCUCAA	1	no match			
3861	AAAAAAAAAAAGGAUCCUCAA	1	2	no annotation		
3862	UAGAUCUUCACCUUUGACUAGCCCC	1	1	Su(Ste)		Su(Ste)
3863	UAGAUCUUCACCGUUGACUACCC	1	1	Su(Ste)		Su(Ste)
3864	AUUCAGCGAUUCCGAUUACUAUC	1	no match			
3865	ACUCCGAACGUACCGGUGCCCCAGU	1	no match			
3866	UAGAUCUUCACGUUGACUUGGCC	1	no match			
3867	UCCAACUGAAGGAGUAGAACAAGCCCGA	1	no match			
3868	UCUCAUCCUCGUAGAACAAGCCCAA	1	2	Su(Ste)		Su(Ste)
3869	GCCGACCGGAGCAGCUUAUACGUAGC	1	1	transposon	LTR	ninja-Dsim-like
3870	UGAUGACCGAAAUUUUGAAAAACGGAU	1	0	repeat		SAR_DM
3871	UCUCAACUGUGAGGGGAUCCUCAA	1	no match			
3872	UUCUCAUCGUCGUAGAGCAAGCCCGA	1	1	Su(Ste)		Su(Ste)
3873	UGUUUCAUCGUUAGGCGGCUCGGGCCG	1	2	AT-chX		AT-chX-B
3874	UCCUCAACUGUGGGGGGAGUCCUCAA	1	no match			
3875	GAAUGUUUCAUCGUUAGACGGCUCGGG	1	no match			
3876	UUCGCCCGCGAGAUGUACCCAAAGACC	1	0	coding gene		CG10627-RA|CG10627
3877	GAUCUUCACCGUUGACUUGCCCCACA	1	0	Su(Ste)		Su(Ste)
3878	UUCGAUCGUCUAGACAAGCCCGACG	1	no match			
3879	CGAAUUGUUAGAGAGCGAGAGAGCGAG	1	0	transposon	DNA	1360
3880	AUCCUCAAUGUGAGG	1	1	coding gene		CG33322-RA|CG33322
3881	AUCCUCAACGUGAGGGGAUCCUCAA	1	no match			
3882	GUCCGUAGUGAUGAAGACAGAGCCA	1	1	transposon	Low_complexity	F-element
3883	UGAUACCGAGAAUGACAAUGGCU	1	0	transposon	LTR	gypsy3
3884	UCCGAGGAAUCUGGGUUCAAGAUUU	1	no match			
3885	UGAUGACCGAAAUUCGGAAAAACGGA	1	0	repeat		ARS406_DM
3886	AUCCUCAACUCGUGAGGGAUCCUCAA	1	no match			
3887	GCAGGACGAAAAGACCCCGGGACCUUCA	1	no match			
3888	AGAAAUCUCAUCGUCGUAGAACAAGCCCGA	1	no match			
3889	AUCUCAUACGUCGUAGAACAAGCCCGA	1	no match			
3890	UCUCAUCAUCGUAGAAUAAGCCCGA	1	1	Su(Ste)		Su(Ste)
3891	UGUUUUCAUCGUUAGACGGCUCGGGCCU	1	no match			
3892	UCGUACAUCGUCGUAGAACAGCCCGA	1	no match			
3893	UCUCAGUCGUCGUAGAAACAAGCCCCUAGCGUGUAACGUCGGGUC	1	no match			
3894	CUCGAAUGAGUGCUUGAUAUGACCAGA	1	1	Su(Ste)		Su(Ste)
3895	UUCCAGUCUCCCGACUAUCAGAUACCCG	1	0	transposon	LINE	INE-1
3896	UUUCCGAAAUCUAGAUGUGUGCAUUUU	1	0	coding gene		CG33664-RA|CG33664
3897	UCCUCAACUGUGAGGGAUCCUAA	1	no match			
3898	UCUCCUCGUCGUAGAACAAGCCCG	1	1	Su(Ste)		Su(Ste)
3899	CGUUAACCAAUUUCCGAACAUACGCC	1	1	transposon	LTR	DM88

3900	UCUCAUCGUCGUAGAACAUCCCCGA	1	2	Su(Ste)		Su(Ste)
3901	UUCCGAAGGAGGUCGGUUGCCUAUGU	1	0	transposon	LTR	gtwin
3902	CAAGAUAUGACCAAAGCAAAGACU	1	0	transposon	DNA	1360
3903	UUGGACCGAACACUUUGAACCAAGUU	1	1	Su(Ste)		Su(Ste)
3904	UCUCAUCGUCGUAGAACAAGCCCGGU	1	2	Su(Ste)		Su(Ste)
3905	UAGAUCUUCACCGUUGACUUUCCCC	1	1	Su(Ste)		Su(Ste)
3906	CGAAAUAGAUCUUUCACCGUUGACCUUGGC	1	no match			
3907	AGCCAGUUACCGAAGCUGCGGAU	1	no match			
3908	GUUCAAGCCCGGGUAUUCUGAAGCCA	1	0	repeat		trf
3909	UCUCAUCGUCGUAGAACAACGCCCC	1	no match			
3910	CCCGUACUCUGUGGUGACCAACAA	1	1	transposon	LINE	baggins
3911	AAAAACCUGAAAGAUGGACUCCUGUAGUCGC	1	no match			
3912	UAAACCGUCAUUCCAGCGCAAGUUGCCC	1	0	transposon	LTR	ninja-Dsim-like
3913	UCCGAACGUACCACGGUCCCCAGU	1	no match			
3914	GAUCCUCAACUGUGAAGGAUCCUCAA	1	no match			
3915	CGCUGACCAAUUUCCGAACAUACGC	1	1	transposon	LTR	DM88
3916	AAUUUACGUAAACCAAAGGCCGUGAGUU	1	no match			
3917	UUCGAACGGUUACCCCAGGAUGCUUGUCGC	1	no match			
3918	AUCGUCAUCGUACGUAGAACAAGCCCGGA	1	no match			
3919	CAACCGGACGGGCCUAGAGUUAUUCCC	1	0	coding gene		CG40635-RA|CG40635
3920	AUGUCACGCGGGAGACCGGGGUCAA	1	no match			
3921	UAUCUGGGAGAGAUCGAUGAAUGACCA	1	0	coding gene		CG3436-RA|CG3436
3922	UUGGGAACACCGCGUGUUGUUGGCCU	1	0	ncRNA		5SrRNA
3923	AUUCUCAUCGUCGUAGAACAAGCCC	1	1	Su(Ste)		Su(Ste)
3924	UUCACCGUUGACUAGCCCCACACA	1	0	Su(Ste)		Su(Ste)
3925	UAAGACCUCUGUUCUGCUUUCAUU	1	0	ncRNA		rRNA
3926	UCAACGUCCCAAGAACUCCCGGU	1	no match			
3927	AUCUCAUCGUCGUAGAACAAGCCCGAGG	1	1	Su(Ste)		Su(Ste)
3928	UAGAUCUUCACCGGUUGACUAGCCCA	1	no match			
3929	AUCGUCUCGGGACUGAAA	1	2	no annotation		
3930	UCUCAUCUUCGUAGAACAAGCCCCGACUGUAAUCGGGUCAAU	1	no match			
3931	AUCGAGAUCAACGACUCACCCGAA	1	no match			
3932	CGCCUGCUGACGGGGCCCGAAAACAC	1	0	transposon	LTR	opus
3933	UACCCACAUCUAACGGAGAGGCCUG	1	0	transposon	LTR	ninja-Dsim-like
3934	UGUUUCAUCGUUAGACGGGCUCGGG	1	no match			
3935	UAGAUCUUCACGUUGACUAGCCCCA	1	no match			
3936	UCUCAUCGUCGUAUAACAAGCCCGUA	1	2	Su(Ste)		Su(Ste)
3937	UUCGAGAGCAUAUAAUCCCCGUAGG	1	no match			
3938	UCAACGGCCCGGACUGCUCCACGCC	1	1	transposon	LTR	ninja-Dsim-like
3939	UAAGUUUACGAAAAUCGAUGACCA	1	0	transposon	LTR	gypsy12
3940	UCUCAUCGUCGUAGAACUACCCCGA	1	1	Su(Ste)		Su(Ste)
3941	CUCUCAUCGUCGUAGAACAACGCCCGU	1	no match			
3942	AUCGUCUCGGGAGUGAAAUACUCUAUCAUCGUAGAACAAGCCCGA	1	no match			
3943	GAAGCACCGUUUCCCGUCCGAUCAACUC	1	no match			
3944	UCUCAUCGUCGUAGAACUAGUCCCGCUC	1	no match			
3945	AGGGGAAUUGGUCCUUAUUAUCCCUC	1	no match			
3946	GAGGAACUCGUUGCCCUUGAUUCCGA	1	0	Su(Ste)		Su(Ste)
3947	CAACCAGCCCGGAGUCUCCAUCCA	1	0	coding gene		CG31359-RA|Hsp70Bb
3948	UCGACGGAGCUCACUUACCUAAUGUA	1	0	transposon	LINE	R1-element
3949	UACUUACAGUGGGUUGUCCAGGGGC	1	0	Su(Ste)		Su(Ste)
3950	CAAGAUAUGACCAAAGCAAAUACA	1	1	transposon	LINE	Cr1a
3951	UUCAAGGUUACCCCAGGAUCCUGU	1	1	repeat		trf
3952	GAAAUCCCGAAGUUUGAACGUGGGAACG	1	no match			
3953	UGCUGACGGGGCCCGAAAACACGAG	1	0	transposon	LTR	opus
3954	CUCUCAUCGUCGUAGAACACGCCGA	1	no match			
3955	UUCACCGUUGACUUGGCCCACACAUC	1	0	Su(Ste)		Su(Ste)
3956	GUUGAUUGACAGACCACUUCUUCU	1	no match			
3957	UCUCAUCGUCGUAGUAACAAGCCCGA	1	no match			
3958	UCGUCUUGGGUGAAAAUCCUCAA	1	no match			
3959	UACUUUUUGUUAGGUGUACUGACCCAG	1	no match			
3960	UAACUUCAGGAUAAGGAUUGGCUCUG	1	1	ncRNA		rRNA
3961	UCUCAUCGUCGUAGAAGAAGCCCCGACUGUAACUCGGGGUCAAU	1	no match			
3962	AUUAAGUGCCGGAAGCGAUCCCGCUG	1	no match			
3963	UCUCAUCGUCGUAGAACAGCCCCGA	1	2	Su(Ste)		Su(Ste)
3964	AGUUUCAUCGUUAGACGGCUCGGGC	1	1	AT-chX		AT-chX-B
3965	UGCAAUUUUCCCGUGGGACUACA	1	no match			
3966	UGUCAUCAACCCGGCGUAUUACCCG	1	0	transposon	LINE	Doc4
3967	AUAGAUCUUCCCCGUUGACUUGGCC	1	2	Su(Ste)		Su(Ste)
3968	UCGUCAUCGUCGUAGAACAAGCCGA	1	no match			
3969	UGUUUCAUCUUUAGACGGCUCGGGUC	1	no match			
3970	UCUCAUCGCGUAGAACAAGCCCG	1	no match			
3971	UCCUAUUACAGGAGAUAUUUGGACAC	1	0	Su(Ste)		Su(Ste)
3972	AUCGGUCUCGGGGAUGAAAUGUUUCAUCGUUAGACGGCUCGGG	1	no match			
3973	AACGGUGCCCAAAAUUUAAACAACUCAUG	1	no match			
3974	UAUAUCUCUCGAGCCUUCAUCCAGU	1	0	transposon	LTR	Tabor
3975	UGGACUUUUUUCCGAUGAGCCAGUGAA	1	0	transposon	LINE	G2
3976	CAAGAUAUGACCUAAGCAAAGACACUA	1	1	transposon	LINE	Cr1a
3977	UCCUCAUCAUCGUAGAACAAGCCCG	1	1	Su(Ste)		Su(Ste)
3978	AAUUAAUACCUCGAGUACGUCUGCUC	1	1	transposon	LTR	Max
3979	UCGUCAUCGUCGUAGAACAAGCCCGAA	1	2	Su(Ste)		Su(Ste)
3980	AUCUCAUCGUCGUAGAACAAGCUCG	1	2	Su(Ste)		Su(Ste)
3981	UCGGCCCGAACCCUUCUGGUGUG	1	0	transposon	LINE	baggins
3982	AUCUCACCGUCGUAGAACAAAGCCCGA	1	no match			

3983	UAGAUCUUCUUCGUUGACUAGCCCC	1	2	Su(Ste)		Su(Ste)
3984	UCCGUCGGGAUGAUUUGUACCGUAGA	1	0	transposon	Low_complexity	F-element
3985	UUUCGAUUGCAAAUUCUCGGGCU	1	0	no annotation		
3986	UCUCAUCGUCGUAGAACAAGUCCCGACUC	1	no match			
3987	AAAACCUAGUUACUUACGGGCUCUAC	1	0	no annotation		
3988	AUAACAAGAUGCAUACGAUGUGGCUC	1	0	transposon	DNA	1360
3989	AUGCACCCGCGAUUGAGAUCAACCAG	1	1	no annotation		
3990	UUCGUCAUCGUCGUAGAACAAGCCCGGA	1	no match			
3991	CUCUCAUCGUCGUAGAAGAAGCCCGA	1	2	Su(Ste)		Su(Ste)
3992	AUCUCAUCGUCGUAGAAACAAAGCCCG	1	no match			
3993	UCCUGUAGUACGAGGUCAUGCCUAGUU	1	2	transposon	LTR	opus
3994	UCGUUGACCAAUUUCCGAACAUACGCCA	1	0	transposon	LTR	Dm88
3995	UAGAUCUUCACCGUUGACUAGCCCGA	1	1	Su(Ste)		Su(Ste)
3996	UCUCAUCGUCGUAGAACAAGCCCGACUGUAACUCGGG	1	no match			
3997	UGUUACAUCGUUAGACGGCUCGGGUC	1	no match			
3998	GACUGCAGAUCGAGAGGUCCCCGGUU	1	0	ncRNA		tRNA
3999	UUCAAGCCCGGGUAUUCUGAAGUCAAG	1	1	repeat		trf
4000	AACUGAGGAGGAUCCUCAA	1	2	coding gene		CG18635-RA|CG18635
4001	CACACAGAGGAUAGCAGAUGAGAGCA	1	no match			
4002	UCUCAUGUCGUAGAACAAGCCCG	1	2	no annotation		
4003	UAAAUACUCAUCAUCGUAGAACAAGCCCGA	1	no match			
4004	UAGAUCUUCACCGUUGACUACCCCCCA	1	2	Su(Ste)		Su(Ste)
4005	UGUUUCAUCGUUAGACGGCUCGGGGCCUGUAACUCGGGGUCAAU	1	no match			
4006	UAAGAGUCUUCACCGUUGACUAGCCCCAACG	1	no match			
4007	UUAAUGCCUGAAGCGAUCCCCACUGC	1	no match			
4008	UCUCAUCGUCGUAGAACACCCGA	1	no match			
4009	CGUAACUUCGGGAUAAGGAUUGCUCU	1	no match			
4010	UUUAGUUACCCGAUUUUUAAACAAA	1	0	transposon	LINE	Cr1a
4011	UUCCGUCGGGAUGAUUUGUACCGUAGU	1	1	transposon	Low_complexity	F-element
4012	UCCGUGGACCUCGAGACCCUCGACC	1	0	no annotation		
4013	UGUGCCCGAGCGUGGGACCUCGAUAA	1	0	transposon	LINE	Cr1a
4014	AAGUUUCAUCGUUAGACGGCUCGGGC	1	2	AT-chX		AT-chX-B
4015	CAAACGGAAAUGGGCCGGCGAGG	1	0	coding gene		CG4625-RA|Dhap-at
4016	GCUACCGGGCGUGAUCAUCCACAAUC	1	0	transposon	LTR	ninja-Dsim-like
4017	AUCGUCAACUGUGAGGGAUCCUCGAA	1	no match			
4018	AUCGUCUCGGGGAUGAAAAUCGAGGAACCAAUCGAUCCAGCUG	1	no match			
4019	CGAAACCUCAACUGUGAGGGGAUCCCUCAA	1	no match			
4020	CACACAGAGGAGAGAGAUGAGUGCA	1	no match			
4021	GCGGCCGGUGACUACAAUGCCAAGCAUA	1	0	transposon	LINE	Doc2
4022	AACAAAAAAAAAAAAAA	1	0	transposon	LTR	INE-1
4023	UUUCGAGUCGAAUAAAUGAUCCGUC	1	0	transposon	LTR	17.6
4024	GAAAAAUCAGGAGAAUGGAGACGGG	1	0	coding gene		CG9899-RA|CG9899
4025	CAGCUCACAGAGGAGAGCAGAUGAGAGCA	1	no match			
4026	ACCUCCGGAAAAGGAUUGGCUCUGA	1	no match			
4027	GCACACAGAGGAGAGCAGAUGAAG	1	no match			
4028	UCUCAUCGUCGUAGAACAAGGCCGA	1	1	Su(Ste)		Su(Ste)
4029	UGACAAACGACGAAACGCUACGGGGAG	1	no match			
4030	UGUUUCAUCGUUAGACGGCUCGGGUUA	1	no match			
4031	UAUCCGUAAGGGUUGACAUAUUGCA	1	0	transposon	LTR	DM88
4032	ACAAGAUAUGACCAAAGCAAAGGCUA	1	1	transposon	DNA	1360
4033	AAACCUCGAAGAACGUCGACCUGGGC	1	no match			
4034	UCGAAGGUAGUAACUAUGACGAUUUGACUUGUAACUCCGGGUCAAU	1	no match			
4035	CGCCCCGUACUCUGUGGUGACCAACAC	1	1	transposon	LINE	baggins
4036	UCUCAUCAUCGUAGAACAAGCCCGACUGUAACUCGGUCAAU	1	no match			
4037	AUCUCAUCGUCGUAGGACAAGCCCG	1	2	Su(Ste)		Su(Ste)
4038	UCAACUGUGAGUCGAUCCUCAA	1	no match			
4039	UAGUAUCUUCACCGUUGACUAGCCCC	1	no match			
4040	GUCUCAUCGUCGUAGAACAAGCCCGUACUGUAACCUCGGGUCAAU	1	no match			
4041	UAGGAAAAACCUAGUUACUUACGG	1	0	Su(Ste)		Su(Ste)
4042	UAGAUCUUCACCGUUGACUUGCCCCGU	1	1	Su(Ste)		Su(Ste)
4043	GCCCGGCUAGCUCAGUCGGUAGAGCAU	1	0	ncRNA		tRNA
4044	CAGCACACAGAGGAGAGCAGAUGAGAGCACUGUAAUCGGGUCAAU	1	no match			
4045	AGAUCUUCACCGUUGACUUGGCCCUCUGUAACUCGGGGUCAAU	1	no match			
4046	AUCUCAUCGUCGUAGAACUAGCCCG	1	1	Su(Ste)		Su(Ste)
4047	UGUUAUCCCUUAUAAGAUCCUCGAAA	1	0	transposon	LTR	flea
4048	UUCAAGCCCGGGUAUUCAGAAGCC	1	1	repeat		trf
4049	UAAUAAUAAUAAAUAAAAGGUGACCGG	1	0	AT-chX		AT-chX-A
4050	AAUUCCGAUCUUGGACAUUCCCCGAUG	1	no match			
4051	AUGAAAAACUGUGAGGGGAUCCUCAA	1	no match			
4052	CAGCACACAGACGGAGAGCACGAUGAGACGCA	1	no match			
4053	UUUUUUAGGCUUUGCAUGCUUUUGCG	1	1	transposon	LINE	G2
4054	AUCGUCAUCGUCCGGUAGAACAAGCCCGA	1	no match			
4055	UGUUUCAUCGUUAAACGGCUCGGGA	1	2	AT-chX		AT-chX-B
4056	CAUCUUCAACGUUUAAAGAUGGAU	1	0	no annotation		
4057	CGUGGCGAAAGCGGCUCUCUGGU	1	no match			
4058	GAGUCCUCAACUGUAAGGGGAUCCUCAA	1	no match			
4059	CCUCGAUAGUAUAGUGGUUAGUAUCAC	1	1	ncRNA		tRNA
4060	UUCGAUUACCUGGUUCGAUCCAAUG	1	0	transposon	LTR	Dm88
4061	GGAUCAACUGUGAGGGUAUCCUCAA	1	no match			
4062	AACUGUGAGGGGAUUCUCAA	1	no match			
4063	AAAAAUACAACUGUGAGGGGAUCCUCAA	1	no match			
4064	GCUCCCAACUGUGAGGGGAUCCUCAA	1	no match			
4065	UGGCCGAGGAUCCGAUUUUCUUUG	1	0	Su(Ste)		Su(Ste)

4066	AGGAAUCAACUGUGAGGGGGAUCCUCAA	1	no match			
4067	CUCGUCAUCGUCGUAGAACAAGCCCGGA	1	no match			
4068	GUGGUUGGGAAGGUCAGGACAUUUA	1	no match			
4069	UGACUCUACGUGACGGAACCGGAGUA	1	0	transposon	LINE	Rt1b
4070	UGAGAAGAUAUUGUACCCGUGAAUU	1	0	coding gene		pseudogene
4071	UGAAUUUGGCCAGUAUUUCAUCU	1	0	transposon	LTR	DM88
4072	AUCGUCGUCGGGAUAGAAAAUCUCAUCGUCGUAGAACAAGCCGA	1	no match			
4073	AGGACUUUUUUUCCGAUGAGCCAGUG	1	2	transposon	LINE	G2
4074	AAAAGUGGAAAUGAAGAAAGUACA	1	2	no annotation		
4075	ACUCACUGUGAGGGGAUCCUCAA	1	no match			
4076	UCUACCCAUGGAAUUCCGAGCUCCUC	1	0	transposon	LINE	G2
4077	UCUCAUCUCCGUAGAACUAGCCCGA	1	2	Su(Ste)		Su(Ste)
4078	UCCUCAACUGUGAGGGGACCCUCAA	1	no match			
4079	UUAAAUAAGACCCACACGCAAAUAAU	1	0	no annotation		
4080	ACCAACGGGAUACCUCAUACUCGAGU	1	0	transposon	LTR	micropia
4081	UACUGUUUCAUCGUUAGACGGCUCGGG	1	2	AT-chX		AT-chX-B
4082	UCUGUGAUGCCCUUAGAUGUCCUGG	1	0	ncRNA		rRNA
4083	AUGGUUUCAUCGUUAGACGCUCGGGCCGUGUAACGUCGGGUCAAU	1	no match			
4084	AAAAUCUCCGAUUGGUAUUAGCGAGUU	1	0	transposon	LTR	opus
4085	UCUUCAUCCAUCGUAGAACAAGCCCGACGUGUAACUACGGGUCAAU	1	no match			
4086	CAACUUGAGGAUCCUCAA	1	2	coding gene		CG10117-RA|ttv
4087	UCCGAAGUGACAGACCGAAUCGCAAUUC	1	0	transposon	LTR	Quasimodo
4088	CCUGACGGACAGACUCAAGACCCUG	1	0	transposon	LTR	F
4089	UCCUGGCGUGCCCGGUACUCAGCUAU	1	1	transposon	LTR	Stalker
4090	GGUGAGGGUGCGAAUCCUCAA	1	no match			
4091	GUAUGACCCCCUGGUACUGAAAUGCA	1	no match			
4092	AGAAAAAAAAAAAAAAAAAAAAGACAUCCUCUAA	1	no match			
4093	CGUUGACCAAUUUCCGAACAUACGCCUG	1	2	transposon	LTR	DM88
4094	CACCCGCGAGUAUAGUUACUUUAGUC	1	0	transposon	LTR	gypsy12
4095	UAGCGUUCGUUGACGUUGAUGGC	1	1	transposon	LTR	HMS-Beagle
4096	UCUCAUCGUCGUAGAACUAGCCUGA	1	1	Su(Ste)		Su(Ste)
4097	UGUUUCAUCUUUUAGACGGCUCGGGC	1	no match			
4098	AACGUUUGCGUUGUCUCUGAGGGAACC	1	1	coding gene		CG4841-RA|CG4841
4099	UUUAACCCGAUAAAAAAGAAUAUC	1	0	transposon	LTR	opus
4100	UGAUGACCGAAAUUUGGAAAAACAGA	1	0	repeat		ARS406_DM
4101	UCGUCUCGGGUGAAACUCAA	1	no match			
4102	UAAAGAAAUCCUGACAAGAUCCCGGGC	1	no match			
4103	AUCGUCUCGGGGAUGAAAUUCCGAGGAACCAAUUGAUCCAGCUG	1	no match			
4104	UCUCAUCGUCGUAGAAGAAGCCCGAU	1	2	Su(Ste)		Su(Ste)
4105	UUACAGUGAAAUGAAGAAAGUACA	1	no match			
4106	CGAAAUGUUUCAUCGUUAGACGGCCUCGGGU	1	no match			
4107	UCGUCAUCGUCGUAGAACAAGCCCGGAGGCG	1	no match			
4108	GACUGCAGAUCGAGAGGUCCCCGUU	1	1	ncRNA		tRNA
4109	UUGGCCAGCAAAUUCGCUCAGUCGAA	1	0	coding gene		CG13391-RA|Aats-ala
4110	UUCUUUGAAGAUGGCCCACAUAAA	1	0	repeat		trf
4111	UGUUUCAUCGUUAGACGGCUCGGCCGAAA	1	2	AT-chX		AT-chX-B
4112	AUCGUCGUCGGGAUAGAAACGUUGACCAAUUUCCGAACAUGCGA	1	no match			
4113	UCUCAUUGUCGUAGAACUAGCCCGA	1	1	Su(Ste)		Su(Ste)
4114	GCCACACAGAGGAGAGCAGAUGAGAGCA	1	no match			
4115	AGAAAUCUCAUCAUCGUAGAACAAGCCCG	1	no match			
4116	AUCGUCUCCGGGAUGAAAACUGUGAGGGGAUCCCUCAA	1	no match			
4117	UCGGAACGUUACCCCGAGGAUGCGGA	1	1	coding gene		CG8357-RA|Rep1
4118	UAAAACAAUUGAAAGAUAAGACCCCAUA	1	no match			
4119	CUCUACUGACUUCAGUAAGCCCCG	1	0	transposon	LINE	baggins
4120	GCCUAACGAGGUAGAUGACUCCUGGAA	1	0	transposon	LINE	Cr1a
4121	GGUGCAUAUCAUUCACGCCACGCUCCGA	1	0	Su(Ste)		Su(Ste)
4122	GCACACAGAGGAGAGCAGAUGAAGA	1	no match			
4123	AUCGUCUCGGGGAUGAAAAUCUCAUCGUCGUAGAACAAGCCCCGA	1	no match			
4124	AUGUUUCAUCGUCGUAGAACAAGCCCGA	1	2	no annotation		
4125	UCUCAUCGUCGUAGAACAAGCCCGCUCGUAACUCCGGGUCAAU	1	no match			
4126	UCAUCUUCUUCUCGACGUUUACCA	1	1	transposon	LTR	ninja-Dsim-like
4127	AUCGUCUCGGGAGUGAAAUAGUUUCUAUCGUUAGACGGCUCGGGC	1	no match			
4128	AUUCGUCAUCGUCGUAGAACAAGCCCGGA	1	no match			
4129	AACAAGAGUCACUUAUUUUUGCUACC	1	0	transposon	DNA	HB
4130	UCUCAUCGUCGCAGAACAAGCCCCGA	1	no match			
4131	UAGAUCUUCACCGUUGACUGCCC	1	2	Su(Ste)		Su(Ste)
4132	AACUUGAGAGUUUGAUCCUGGCU	1	no match			
4133	UGUUUCAUCGUUAGACGCUCGGC	1	no match			
4134	GUGAUGUCAACGUUUGAUCCCCUA	1	1	transposon	LTR	ninja-Dsim-like
4135	UCUUCACCGUUGACUUGCCCGCA	1	1	Su(Ste)		Su(Ste)
4136	AAGAUAUGACCAAAGCAAAAGACAC	1	1	transposon	LINE	F
4137	GAAUAUCGUCUCGGGACGUGAAAUACUCAUCGUCGUAGAACAAGCC	1	no match			
4138	CGCAAAAGUUGACCGAACUUGAA	1	1	ncRNA		rRNA
4139	UCUCAUCGUCGUAGAACAAGCGUCGA	1	no match			
4140	CGUUGACCAAUUUCCGAACAAGCCCGA	1	no match			
4141	AUGUUUCAUCGUUAGACGGCUCGGGA	1	2	AT-chX		AT-chX-B
4142	CCUCAACGUGAGGGAUUCUCAA	1	no match			
4143	GUAGUGGCGAGCGAAAUGGGAUGAGC	1	no match			
4144	CGAAAUGUUUCAUCGUUAGACGCCUCGGG	1	no match			
4145	CCUCAUCGUCUUAGAACAAGCCCGA	1	2	Su(Ste)		Su(Ste)
4146	AUCUCAUCGUCGUAGAACAACCCCGAA	1	2	Su(Ste)		Su(Ste)
4147	GAUUACCGAGGGCCGAGUACAUAUCGAG	1	0	transposon	LTR	G4
4148	UCCGAACCAAUCGAUCCAGCUG	1	no match			

4149	UAAGAUUGCGAUCUGGAGUUGCCCCC	1	2	transposon	DNA	1360
4150	ACACUUCAACUGUGAGGGGAUCCUCAA	1	no match			
4151	AACCUCGAAGAACGUCGACCUGGGC	1	no match			
4152	ACUCAACUGUGAGGGGGAUCCUCAA	1	no match			
4153	CCGAACUUGAGACCCCAGCAGCUC	1	0	no annotation		
4154	CUCGGGAUGAAAGGAUCCUCAA	1	no match			
4155	UAGGCUAAUCGGACAGGUGACUCG	1	0	transposon	LTR	GATE
4156	AUCUCAUCGGCGUAGAACAAGCCCGA	1	2	Su(Ste)		Su(Ste)
4157	UGUUUCAUUGUUAGACGGCUCGGGU	1	2	AT-chX		AT-chX-B
4158	AUGUUUCAUCGUUAGACGGCUCGGGCCU	1	no match			
4159	AUCCUCAACUGUGAGAA	1	1	coding gene		CG1666-RA|Hlc
4160	UUCCGAGGAACCCAUUGAUCCAGCU	1	0	no annotation		
4161	UCGAACCAGAGGCUGGAUCCUCAA	1	no match			
4162	GCUCAUCGUCGUAGAACAAGCCCG	1	1	Su(Ste)		Su(Ste)
4163	AAAAUCUCAUCGUCGUAGAACAAGCCCGU	1	no match			
4164	UGUUUCUUCGUUAGACGGCUCGGGC	1	1	AT-chX		AT-chX-B
4165	UACGUUUCUCCCUAAUGACUUCGUCCGA	1	1	transposon	LTR	qbert
4166	CCUCAACUGUGGGGAUCCUCAA	1	no match			
4167	UAUAAUGCCAGAAUUUUAUUGGGG	1	2	transposon	LINE	baggins
4168	UGAACCUCGGGUACGUCUGCUCU	1	no match			
4169	UUUCAUCGUCGUAGAACUAGCCCGA	1	1	Su(Ste)		Su(Ste)
4170	UAUCCGUAAGGGUUGACAUAUUGCU	1	1	transposon	LTR	DM88
4171	CUGUGAGGGGAUCCCUCAA	1	no match			
4172	AACCAAUCCCAACCAAACCCGUCAA	1	0	transposon	LINE	G2
4173	UUCACCGUUGACUUGGCCCACACAU	1	0	Su(Ste)		Su(Ste)
4174	AUCUCAUCGUCGUAGAACAAGCCCUAG	1	2	Su(Ste)		Su(Ste)
4175	UGCACGUACCUGGAGAACAAUCAGA	1	2	transposon	LTR	ninja-Dsim-like
4176	UCUCAUCGUUAUACGGCUCGGGC	1	2	AT-chX		AT-chX-B
4177	GAAUAUCGUCUCGGAGUGAAUAGAUCUUUACCGUUGACUUGGCC	1	no match			
4178	UCAACUGUGAGGGGACUCCUCAA	1	no match			
4179	CGAAAUACUGACUCGUGUACGUUCGUUGA	1	no match			
4180	GACGUAGCUGGCCGAGGAUCCGAGUU	1	no match			
4181	GGUACUUUUUUCCGAUGAGCCAGU	1	2	transposon	LINE	G2
4182	UGAUGAUGUUACCCCUUACAAAAA	1	0	repeat		SAR_DM
4183	UCUCAUCGUCGUAGAACAAGCCCGACU	1	2	Su(Ste)		Su(Ste)
4184	AUGUUCAUCGUUAGACGGCUCGGG	1	no match			
4185	UAAAGAUAAAUAUGACCGAUUGCGU	1	0	transposon	LTR	mdg1
4186	CAGCACACAGAGGAGAGCAGAUGAGAGAA	1	no match			
4187	CACCGGGUUUUGGCACUUUGCCGCGG	1	0	transposon	LINE	R1
4188	UAAUGCCCCGAAGGUAGCCCUCUGU	1	1	transposon	LINE	R1-element
4189	UGAUAUGACGCGUGACAUGCCACA	1	0	transposon	LINE	G2
4190	GCUGUGCGUUCGUGGAGACGACCCGA	1	0	transposon	LTR	ninja-Dsim-like
4191	AUCGAAAUCUCGGUGGGCAGUAGACCCG	1	no match			
4192	UUAUAAGGAUGAUGAUACCGAUUAAAGG	1	0	no annotation		
4193	UUCGUAGAGAUGUCUCGGGAACCAUAA	1	1	coding gene		CG3338-RA|CG3338
4194	UACGCGCGGCAGGACGGAAAGACC	1	no match			
4195	GAAUAUCGUCUCGGGAGUGAAAUACUCAUCAUCGUAGAACAAGCC	1	no match			
4196	UUAUAAGGAUGAUGAUACCGAUUAAU	1	1	no annotation		
4197	CAAGAUAUGACCAAAGCAAUACUA	1	no match			
4198	UCUCAUCGUCGUAGAACAAGACCCGA	1	no match			
4199	UCGUCCUAAGAUGAUGUGACCCCAGA	1	no match			
4200	AGAUGGUGCGAGAACCCAGAGUA	1	0	coding gene		CG11719-RA|Mst98Ca
4201	UGGGAUCGAUUGGAAAUAACCUGA	1	no match			
4202	AUCGUCAGUCAAGUCGUAGAACAAGGGCCCGACG	1	no match			
4203	UUGGGAACACCGCGUGUUGUUGGCC	1	0	ncRNA		5SrRNA
4204	CAAGAUAUGACCAAAGAAUGACCA	1	0	transposon	DNA	1360
4205	AGGGCCCGCCUGACCUUGAAUCA	1	no match			
4206	GAGGAACUCGUUGCCUUUGAUUCCGA	1	0	Su(Ste)		Su(Ste)
4207	UCUCAUCGUCGUAGAACAAGCCCGACUGUAACCGGGUCAAU	1	no match			
4208	UGUUUCAUCUUUAGACGGCUCGGA	1	2	AT-chX		AT-chX-B
4209	AAUCCUAACUGUGAGGGGAUCCUCAA	1	no match			
4210	AGGAUCCUCAACUGUGAGGGAAUCCUCAA	1	no match			
4211	UAGAUCUUCACCGUUGACUUGCCCCCG	1	1	Su(Ste)		Su(Ste)
4212	UAGUAUCUUCUACCGUUGACUUGGCCC	1	no match			
4213	ACCGGAGAAGACGUUGAAUGGAUC	1	0	no annotation		
4214	GAAGACCCCAGGUAGUGAUUGUAUCC	1	0	coding gene		CG3506-RA|vas
4215	GCAAAAGAAACCAGAGGACAACCUGAA	1	0	repeat		trf
4216	AGUAGAUCUUCACCGUUGACUAGCCC	1	0	Su(Ste)		Su(Ste)
4217	CGUUUCAUCGUUAGACGGCUCGGGA	1	2	AT-chX		AT-chX-B
4218	CGUGGACGAGUUGCUCGAUUUGUACGU	1	0	transposon	Low_complexity	F-element
4219	AAAUGCGAGUGAAGCCCGGAAUAAACUU	1	0	transposon	DNA	transib2
4220	UCAAUGUUGGCCCCAGAAUACGCAG	1	1	no annotation		
4221	AAUCCUCAACUGUGAGGAUCCUCAA	1	no match			
4222	UGAGACCGCCCCUGGACAACCCACUAA	1	2	Su(Ste)		Su(Ste)
4223	UAAGAAAAUGUCCGGACAGCUG	1	0	Su(Ste)		Su(Ste)
4224	AUCUCAACUGUGAGGGAUCCUCAACG	1	no match			
4225	UGUUUCAUACCUUGAGCCAACUGAUC	1	0	transposon	LTR	gypsy6
4226	UCUCAUCGUCGUAGAACAAGCCCGACUGUAACUCGGGGUCAA	1	no match			
4227	UGGACCUUUUUCCGAUGAGCCAGU	1	1	transposon	LINE	G
4228	UCAACUUGAGGGGGAUCCUCAA	1	no match			
4229	UGUUUCAUCGUUAGACUGCUCGGCC	1	2	AT-chX		AT-chX-B
4230	GGAUCCUCAACUGUGAGGGAUCAA	1	no match			
4231	UCUCACCCGUCGUAGAAGAAGCCCCGA	1	no match			

4232	UCCUCAACUGUUGAGGAUCCUCAA	1	no match			
4233	UAUAGACCGAUUAGUCUGCUGACAA	1	0	transposon	LINE	G2
4234	CAUGCUCGAGAUGACAACCUUGGACC	1	0	transposon	LTR	micropia
4235	UAAGUAAGCCCGAAGAUUGUCCAA	1	0	transposon	LTR	opus
4236	UAAUAUUGGACCCAGUACACUGCC	1	1	transposon	LINE	Doc
4237	UAACAUCAUCAUCUGACUCGCAAUU	1	0	transposon	DNA	1360
4238	AUCAUCAACUGUGAGGGGAUCCUCAA	1	no match			
4239	AUGUUUCAUCGUUAGACGGCUCGGGUC	1	no match			
4240	UCUCAUCGUCAGAACAAGCCCGA	1	no match			
4241	UCUCAUCGUCGUUGAACUAGCCCG	1	1	Su(Ste)		Su(Ste)
4242	CAGCACACAGAGGAGAACAGAUGAGAGC	1	no match			
4243	UACAAGCAAGACCACCACACCCGA	1	0	transposon	LTR	gypsy12
4244	UGAAAGAUUACUCAGAGAACAACU	1	0	coding gene		CG12288-RA|CG12288
4245	GCUGGCCGAGUAUUUGAUGUUCU	1	0	Su(Ste)		Su(Ste)
4246	UACUCAUCAUCGUAGAACAAGCCGA	1	no match			
4247	UCUCAUCAUCGUAGAACAAACCCAA	1	2	Su(Ste)		Su(Ste)
4248	AAGAUCUUCACCGUUGACUUGGCCCAU	1	2	Su(Ste)		Su(Ste)
4249	UUUCCUGGCUAGGUUGUCCGCUA	1	0	transposon	LINE	baggins
4250	CUGAUGACCUGAUGACUCUUUGACA	1	no match			
4251	UUGUUACUUCCCGUGUUCAGCUUU	1	0	transposon	LINE	Doc3
4252	CGAAAUGUUUCAUCGUUAUACGGCUCGGGC	1	no match			
4253	UUCCGAGGAACCAAUUGAUCCAGCUCC	1	1	Su(Ste)		Su(Ste)
4254	GUAACUUCGGGAUAAGGAUUGGCUUUAA	1	2	ncRNA		rRNA
4255	CAGCACACAGAGGAGAGCAGAUGAGGGCA	1	no match			
4256	CAAGAUUUGACCAACGCAAAGACUA	1	no match			
4257	UACGGAACGAGAGACGCGACUCGGAU	1	no match			
4258	AAUCCUCAACUGUGAGGGAAUCCUCAA	1	no match			
4259	ACCAAGAUAUGACCAAAGCAAAGAC	1	2	transposon	LINE	Cr1a
4260	AUGUUUCAUCGUUAGACGACUCGGGC	1	2	AT-chX		AT-chX-B
4261	GCCAAUUUACAUUUAGACGAGACGGCCCC	1	no match			
4262	GCACACAGAGGAGGGCAGAUGAGAGCG	1	no match			
4263	UUCCUCGGGAUCAAGGGCAACGAGUA	1	1	Su(Ste)		Su(Ste)
4264	UCCCUACAAUGUUAUCCAGGACCCCA	1	1	transposon	LINE	Rt1b
4265	UUAGCUUCAUCCAAGCGAACUGUCGCCC	1	2	transposon	LINE	R1
4266	CUCAACUGUGAGGGGAUCCUCGAA	1	no match			
4267	UCUCAUCUUCGUAGAACAAGCCCGGA	1	2	Su(Ste)		Su(Ste)
4268	UCCGAUUUGGCGCAAAGAGAUGGAC	1	1	transposon	LTR	ninja-Dsim-like
4269	UUCGCAGAAUAAAUAUACGCCCGGUC	1	0	transposon	LTR	GATE
4270	AUAACGUUCGGGAUAAGGAUUGGCUCUGGA	1	no match			
4271	CAAGAUGUGACCUAAGCAAAGACACU	1	1	transposon	DNA	1360
4272	AACUGUGAGGGGUCCUCAA	1	2	no annotation		
4273	CAGAUCUUCACCGUUGACUUGGCCC	1	1	Su(Ste)		Su(Ste)
4274	AUCCUCAACUGUGAGGGAUCCCUCAA	1	no match			
4275	GGUAGAGCUCGUUUGAUUUUGACCA	1	0	transposon	LINE	X
4276	GAGAGCUCUGAUGACGAACCCGG	1	0	transposon	LINE	G6
4277	GUCGCCUGCUGACGGGGCCCGAAAAA	1	1	transposon	LTR	opus
4278	AGUUUCAUCGUUAGACGGCUCGGG	1	1	AT-chX		AT-chX-B
4279	UGGGCCGAACACUUUGAACCAAGUAU	1	0	Su(Ste)		Su(Ste)
4280	UUCACCGUUGACUAGCCCCACACAUC	1	0	Su(Ste)		Su(Ste)
4281	UCCGAACGUACCACGGUGCCCCAGUGCUGUAAUCGGGUCAAU	1	no match			
4282	UCCUAACUGUGAGGGGAUCCUCAA	1	no match			
4283	AGUCAUGCGAGGCGAAACCGGGUUCA	1	no match			
4284	AUGUACUGUUUGUAGGUGUACUGGCC	1	1	transposon	LINE	baggins
4285	UGACAAAAUGGCCGAAGUUAUCAAA	1	0	transposon	LINE	BS
4286	GAUCCAUGGCAACUUCUGUCGGGUCAAU	1	no match			
4287	GAGUCUCGAACUAGGUGAGGAGAGUCCUCGAA	1	no match			
4288	UAGAAGAAACAUUGAAGCGCUUGAC	1	no match			
4289	UGUACUGUUUGUAGGUGUACUGACC	1	0	transposon	LINE	baggins
4290	ACUCAACUGUGAGGGAUCCUCAA	1	no match			
4291	UGGACUUUUUUUCCGUAUGAGCCAGU	1	no match			
4292	UAUAAGAUUGGACCCAGUACACUGCCUU	1	0	transposon	LINE	Doc
4293	UGUUUCAUCGUUACGACGGCUCGGG	1	no match			
4294	AUUCGGGAUGAAAUCUCAUCGUCGUAGAACAAGCCCGA	1	no match			
4295	CUCAACUGUGAGGAGGAUCCUCAA	1	no match			
4296	AAAAGGUAGGAAAUGAAGAAAGUACA	1	no match			
4297	AUCGUCUCGGGACUGAAAGGUUUCAUCGUUAGACGGCCUCGGGC	1	no match			
4298	GCCCCGGUACUCUGUGGUGACCAACACG	1	no match			
4299	UCUCAUCGUCGUAGAACAAUCCCGAG	1	1	Su(Ste)		Su(Ste)
4300	UCCUGUUGAUGUGAGGGUCCCAAG	1	1	transposon	LINE	G2
4301	GGGGCGACACUCGGGCUCACGGC	1	no match			
4302	UAUAAUCGCCCCGAAGGUUUGCCCUCUG	1	no match			
4303	UGUUUACCCAGAAACAAUG	1	0	transposon	LINE	HMS-Beagle2
4304	CUUGACGGAGCUCACUUACCUAAUGUA	1	1	transposon	LINE	R1-element
4305	AUGUUUCAUCGUUAGACGGCUCGGGCCUU	1	no match			
4306	GGAUCCUUAACUGUGAGGGAUCCUCAA	1	no match			
4307	AUCGUACCAUAGGACCCGGUCCCUAAA	1	0	transposon	LINE	F
4308	UCUUCUUCUCGACGUUGACCAGCCUC	1	0	transposon	LTR	ninja-Dsim-like
4309	UGUUUCAUCGUUAGACGGCUCGAGAUUC	1	no match			
4310	AUCUCAUCGUCGUUAGAACAAGCCCGA	1	no match			
4311	AUCGUCUCGGGGAUGAAAUGUUUCAUCGUUAGACGGCUCGGGCC	1	no match			
4312	ACGUGAGGGGAUCCUCAA	1	no match			
4313	AAUCUCAUCGUUAGAACAAGCCCGA	1	no match			
4314	GACAAUGACGACCAGCCCAGUACC	1	0	coding gene		CG18749-RB|CG18749

4315	UGUCAACGUUUGAUCCCCUGGAUUCC	1	0	transposon	LTR	ninja-Dsim-like
4316	CUUCGGGAUAAGGAUUGGCUCUA	1	1	ncRNA		rRNA
4317	UUCGUUAGCCCGUCCGUUUCUCGAGA	1	1	transposon	LINE	Cr1a
4318	UCUCAUCGUCGUAGAAGAAGCCCGGA	1	2	Su(Ste)		Su(Ste)
4319	UACCAUUAUUUUGGCCAUGCUCGACUC	1	0	coding gene		CG31911-RA|Ent2
4320	ACCGGACGUGCUGUGGCCCGUAUUCCCC	1	0	coding gene		CG5502-RA|RpL4
4321	GGAUCGAACUGUGAGGGGAUCCUCCAACG	1	no match			
4322	GGUCCCAACUGUGAGGGAUCCUCAA	1	no match			
4323	UAAAAUCUCAUCGUCGUAGAACAAGCCCGA	1	no match			
4324	GAAGAUGUCCGGACAGCUGGGCCCGAA	1	0	Su(Ste)		Su(Ste)
4325	CUCUCAUCGUCGUAGAACUACGCCCGUA	1	no match			
4326	AUCCGAACGUACCACGGUGCCCCAGUG	1	1	transposon	Low_complexity	F-element
4327	UAGAUCUUCACCGUUGACUUGCCCCCUC	1	1	Su(Ste)		Su(Ste)
4328	UGCUCUCGAAUGUAAUGCCCGAUAUA	1	no match			
4329	AUGUUUUCAUCGUUUAGACGGCUCGGGCCC	1	no match			
4330	UAAGCCCCGUACUUCUGUGGUGACCAA	1	no match			
4331	GUAUGACCCCCUGGUACUGAAAUGCAGA	1	no match			
4332	GCUGAUCACGUCUCGCGAACCUUGGU	1	0	transposon	LINE	R1-element
4333	UGUUUGUAGGUGUACUGACCCAGU	1	0	transposon	LINE	baggins
4334	GGGAACACCGCGUGUUGUUGGCCU	1	0	ncRNA		5SrRNA
4335	AUCGAUCUUCACCGUUGACUAGCCCC	1	2	Su(Ste)		Su(Ste)
4336	AUCGUCAUCGUCGUAGGAACAACGCCCGA	1	no match			
4337	UACAAUCUCCGAGCAAGACCCCGACUGUAACUCGGGGUCAAU	1	no match			
4338	GAAGCCGUCGAGAGAUAUCGGAAGAGU	1	0	ncRNA		rRNA
4339	CCUCAACUGUGAGGGAUCCUCAAA	1	no match			
4340	CCUCAUCAUCGUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
4341	GAACUCAUCCUACUCGCUAGACGCAC	1	1	ncRNA		alphagamma-element
4342	UCUUGUUUCAGAGCGUUGGCCACG	1	2	transposon	Low_complexity	F-element
4343	UGUUUUCAUCGUUAGACGGCUCGGGCUGUAACUCGGGGUCAAU	1	no match			
4344	AGUCUCGGGUGAAAGGAUCCUCAA	1	no match			
4345	AAAGAAACCCGAUUCCUCUUCCAUGAG	1	2	transposon	LINE	Doc4
4346	UCAACUGUGACCUCAA	1	1	no annotation		
4347	UCAUCUUCUUCUCGACGUCGACCAG	1	1	transposon	LTR	ninja-Dsim-like
4348	GAAAUCGUCAUCGUCGUAGAACGAAGCCCGA	1	no match			
4349	CGAAAUAGAUCUUCACCGUUGACCUAGCCC	1	no match			
4350	GACGCCGAGUGGAAUCCAGGGAGCAUA	1	0	transposon	LTR	ninja-Dsim-like
4351	UUGGUUAGAAAACAACCAGGACUAUA	1	0	Su(Ste)		Su(Ste)
4352	UCUCAUCGUUGUAGAACAAGCCCUAG	1	2	Su(Ste)		Su(Ste)
4353	GAAUUGCUUACCGUGGGUUGUCCAG	1	1	Su(Ste)		Su(Ste)
4354	GCUCCGAUCUCCUGAGUCGAACUCCGA	1	1	transposon	LINE	R1-element
4355	UGUUUUCAUCGUCAGACGGCUCGGGC	1	no match			
4356	AUCGUCUCGGGACUGAAACUAGAUCUUCACCGUUCACUACGCCCA	1	no match			
4357	CAACUGUGAGGGGAGUCCUCAA	1	no match			
4358	UCCUCAACUGUUGAGGGAUCCCUCAA	1	no match			
4359	UAAAAAAAAAAAAAAAA	1	0	transposon	LTR	17.6
4360	UGUUAAUAAGACCCCGUUUGUUUC	1	0	transposon	LTR	297
4361	AGAAAUCUCAUCGUCGUAGAACAACGCCCG	1	no match			
4362	UCAUCUUCUUCUCGACGUUCGACCA	1	no match			
4363	UAGAUCUUCACCGUAGACUUGGCCCA	1	1	Su(Ste)		Su(Ste)
4364	UCUCAUCGUCGUAGAACAAGCCGG	1	1	Su(Ste)		Su(Ste)
4365	UCUCAUCGUCGUAGAACGAGCCCCGA	1	no match			
4366	UACAAUCUUCGAGCAAGACCCCGA	1	0	transposon	LINE	Circe
4367	UCUCAUGUCGUAGAACAAGCCCGA	1	2	no annotation		
4368	UCUCAUCGUCGUAGAACAAGCCCCUA	1	2	Su(Ste)		Su(Ste)
4369	UGUUUUCAUCGUUAGACGGCUCGGGG	1	no match			
4370	GCCCGUAGUGAUGAAGACAGAGCCA	1	0	transposon	Low_complexity	F-element
4371	CAGCACACAGAGGAGAGCAGAUGAGAGGCA	1	no match			
4372	AAAUAUUGGCGCGGUCAUCGGAACUG	1	0	transposon	LTR	ninja-Dsim-like
4373	CAGCACACAGAGGACGAGCAGAUGAGAGCA	1	no match			
4374	AAAACCCUGAAGAUGAUCCUGAGUC	1	2	transposon	LINE	Helena
4375	UUGUUGAUCAGUUUACACAUUAUUG	1	0	transposon	LINE	Cr1a
4376	CAAGCACACAGAGGAGAGCAGAUGAGAGCA	1	no match			
4377	UGUUUCAUCGUUAGACGGCUCGGCCGUGUAACUACGGGUCGAAUA	1	no match			
4378	GUAUCGUCUCGGGACUGAAAUCUCAUCGUCGUAGAACAAGCCCGA	1	no match			
4379	UGACUUGAGGAUGGCUUCGAUGUUG	1	0	transposon	LINE	X
4380	CCCCGUACUCUGUGGUGACCAACACA	1	0	transposon	LINE	baggins
4381	AGAUGACCGAAAUUCGGAAAGACGG	1	1	repeat		SAR_DM
4382	UCACAUCGUCGUAGAACAAGCUCG	1	2	Su(Ste)		Su(Ste)
4383	CAACUCACAGAGGAGAGCAGAUGAGAGCA	1	no match			
4384	UUUUUCAUCGUUAGACGGCUCUGGC	1	2	AT-chX		AT-chX-B
4385	AUCCCAUCAUCGUAGAACAAGCCCGA	1	2	Su(Ste)		Su(Ste)
4386	UAAGAUCGAAUCUGUUACGGCCCUU	1	0	repeat		trf
4387	UACUGAGUCGAUUACCGAGGGCCGA	1	0	transposon	LTR	G4
4388	UCCUCAUCGUCCGUUAGAACCAAGCCCCGACU	1	no match			
4389	UCCAUAUUCAGACAAUUAUUUUAAGU	1	1	repeat		SAR_DM
4390	GGUACGACCCAAGUAGGACUAUUGU	1	0	transposon	LTR	297
4391	UCUCAUCGUCGUAGAACAAAGCCCGUACUGUAACCUCGGGUCAAU	1	no match			
4392	UCUGCUUCAUCGUUAGACGGCUCGGG	1	1	AT-chX		AT-chX-B
4393	UAUAGGACCCCAUUACAAUUGUAA	1	0	transposon	DNA	1360
4394	AGAAAUUGUUUCAUCGUUAGACGGCUCGGGC	1	no match			
4395	CAAGAUAUGACCUAUGCAAAGACACU	1	1	transposon	DNA	1360
4396	UGUUUCAGAGCGUUGGCCAGUUUUCC	1	2	transposon	Low_complexity	F-element
4397	CCUCAUCCUACAGACCAAUCAGUCUCCC	1	1	transposon	LINE	Doc2

4398	UCUCAUCGUCGUAGAACCAGCCCG	1	1	Su(Ste)		Su(Ste)
4399	UCAACUGUGAGGUCCUCAA	1	2	no annotation		
4400	AUCCUCAACUGUGAGGGAUCCUCGAA	1	no match			
4401	AGAACUAUCAACCGUCAGGUGCCCCA	1	0	transposon	LTR	opus
4402	GCUGGCCGAGUAUUUGAUAUUCUUU	1	0	Su(Ste)		Su(Ste)
4403	AGGAAUUCAACUGUGAGGGGAUCCUCAA	1	no match			
4404	AUCUCAACUGUGAGGGAUCCUCAA	1	no match			
4405	UAACUAAGAACCGUGAGAUAUCGCUAA	1	0	no annotation		
4406	GUCUCGGGAUAAAGGAUCCUCAA	1	no match			
4407	CACGGAGUACGGGGCUUACUAAAGUC	1	1	transposon	LINE	baggins
4408	AUGAUUCCGAUCUUGGACAUUCCCGA	1	0	transposon	LTR	roo
4409	AAGUCUCAUCGUCGUAGAACAAGCCCGA	1	no match			
4410	UUGUAAUGACCCCCCAUAAAAAAUUGGA	1	1	repeat		SAR_DM
4411	GAUGUCCACCAAAGGCAUGCCCGAUGU	1	0	transposon	LTR	opus
4412	UAAGAUUGCACCCAGUACACUGC	1	1	transposon	LINE	Doc
4413	UCUCAUCUUCGUAGAACAGGCCCGA	1	2	Su(Ste)		Su(Ste)
4414	UCUCGAUCGUCGUAGAACAAGCCCGUCG	1	no match			
4415	ACACCUUGACGGAGCUCACGUACC	1	1	transposon	LINE	R1-element
4416	AUACUACACUCGUCGUAGAACAAGCCCGA	1	no match			
4417	UUCUCAUCGUCGUAGAACAGGCCCGA	1	1	Su(Ste)		Su(Ste)
4418	UAGAUCUUUACCGUUGACUUGGCCC	1	0	no annotation		
4419	AAUUUAAGCAUAUUAAUUAGGGGAGG	1	0	ncRNA		rRNA
4420	GGAUCCCAACUGGAGG	1	1	coding gene		CG15099-RA|CG15099
4421	GCAUACAUGCCCAGGAGUCUGUUCA	1	1	transposon	LINE	baggins
4422	UUCCCGUGUCUUCCGUGUUAAGGUCU	1	no match			
4423	UAAAUUUUUCAUCGUUAGACGGCUCGGGC	1	no match			
4424	ACAUGUCGAGCUCCCAGAACAACAACA	1	no match			
4425	UGUGUUGACUGGAAGUGCCUUCAUA	1	0	transposon	LTR	mdg1
4426	UUCCCAUCGUCGUAGAACAAUCCCGA	1	2	Su(Ste)		Su(Ste)
4427	ACUAACUUCCAUCCUAAAACUGAUAC	1	no match			
4428	UGUUUCAUCGUUAGACGGGCUCGGGU	1	no match			
4429	GAAUGUAAAAUUCUCGCCUCCUCU	1	0	AT-chX		AT-chX-B
4430	CACGCCACGCUCCGACCUGAUGUA	1	0	Su(Ste)		Su(Ste)
4431	CAAGAAAUGACCUAAGCAAAGACAC	1	1	transposon	DNA	1360
4432	UCUCAUCGUCGUAGAGCAAGCCCUA	1	2	Su(Ste)		Su(Ste)
4433	CAAGUACCGGAGUAUGUGCUUCACA	1	0	transposon	LTR	1731
4434	CGGGAUGAAAUCUCAUCGUCGUAGAACAAGCC	1	no match			
4435	CAGCACACAGACGGAGAGCAGAUGAGACGCCA	1	no match			
4436	UGUUUCAGAGCGUUGGCCAGUUCCG	1	2	transposon	Low_complexity	F-element
4437	AAUGUUUCAUCGUUAGACGGCUCGGG	1	2	AT-chX		AT-chX-B
4438	UCUCCUCGUCGUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
4439	AUCGUCUCGGGACUGAAACUCUCAUCCGUUCGUACGAACAACCGC	1	no match			
4440	UCUCAUCGUCGUAGGACAAGCCG	1	2	Su(Ste)		Su(Ste)
4441	CCCCGUACUCUGUGGUGACCAACACACC	1	0	transposon	LINE	baggins
4442	AUACCUCAACUGUGAGGGAUCUCAA	1	no match			
4443	UCUACUUCCCACCUAGACACUCGACG	1	2	transposon	LINE	G2
4444	CCAGGCCCAUCUGGUUGAACGUGUCCU	1	0	Su(Ste)		Su(Ste)
4445	UUCGCGAUGUACGAGUUGGCCAAGGA	1	0	coding gene		CG8859-RA|Cyp6g2
4446	CCGAACGUACCACGGUGCCCCAGUCUGUAACUCGGGGUCAAU	1	no match			
4447	AUCUUCAUCGUCGUAGAACAAGCCCCGA	1	no match			
4448	UUCUCAUCGUCGUAGAACAAGCCCCG	1	2	Su(Ste)		Su(Ste)
4449	UCGUCUCGGGGAUGAAAAUCUCAUCGUCGUAGAACAAGCCCGA	1	no match			
4450	AUUAAUACCCAGGUGAUUACCACUAUU	1	2	Su(Ste)		Su(Ste)
4451	UUAGGUGUAGGCGCAGCGAAAGCUAGU	1	no match			
4452	UAGAUCUUCACCGUUGACUAGCCCCCUGUAACUCGGGGUCAAU	1	no match			
4453	UCGAUCUCCUGAGUCGGACUCCGAG	1	1	transposon	LINE	R1
4454	AGAAAAAAAAACCUGGAAGAUGAUCCUGAGUCGC	1	no match			
4455	CCCAAUCAGUUUUUAUCUGCAGAGAUG	1	no match			
4456	AUCUCAUCGUCGUAGAACAAGCCCGAAG	1	1	Su(Ste)		Su(Ste)
4457	CUUAUAAGGAUGAUGAUACCCGACUAAA	1	no match			
4458	GAUGGGGGACCGCUUGGGAACA	1	0	ncRNA		5SrRNA
4459	UACUCAUCUUCGUAGAACAAGCCCGA	1	2	Su(Ste)		Su(Ste)
4460	AAUCCUCAACUGUGAGGGGUCCUCAA	1	no match			
4461	GCAAUUUUGGCCGUCACUAAAAAAGU	1	0	transposon	LINE	Fw3
4462	UUCAAGGUUACCCCAGGAUGCUGUGUC	1	1	repeat		trf
4463	UCUCGAUCUUCGUAGAACAAGCCCGACG	1	no match			
4464	UUCAGAUGAAGACUGCGAUACCCCA	1	1	transposon	LINE	Cr1a
4465	CUGUUUCAUCGUUAGACGGCUAGGG	1	1	AT-chX		AT-chX-B
4466	CAGCACACAGAGGAGAGCAGAUGGGAGCA	1	no match			
4467	UAAAUGCCAGAAUUUUAUUCGGGGUUC	1	no match			
4468	UUUAUCUGCAGAGAUGUAUGAACGAA	1	0	transposon	LTR	gypsy8
4469	UUCUCAUCGUCGUAGAACAAUCCCG	1	1	Su(Ste)		Su(Ste)
4470	UAGAUCUUCACCGUUGACUUGCCCCAA	1	1	Su(Ste)		Su(Ste)
4471	UCGUCAGUCGUCGUAGAACAAGCCCGA	1	no match			
4472	AUUCACUUCUUCUCGACGUUGACCAG	1	no match			
4473	GCCCCGUACUCUGUGGUGACCAACAU	1	0	transposon	LINE	baggins
4474	UAGAUCUUCACCGUUGACUAACCCCC	1	2	Su(Ste)		Su(Ste)
4475	GUAGAUCUUCACCGUUGACUUGGCCCA	1	0	Su(Ste)		Su(Ste)
4476	UAUAAUGCCCCGAAGGUUGCCCUCU	1	0	transposon	LINE	R1-element
4477	AUCGUCUCGGAUGAAAUACUCAUCGUCGUAGAACAAGCCCGAC	1	no match			
4478	UGCACGGACCUGGAGAACAACCAGA	1	0	transposon	LTR	ninja-Dsim-like
4479	UCUCAUCGUCGUAGAACAAUACCGA	1	2	Su(Ste)		Su(Ste)
4480	CGCGCAGGUUCAAAUCCUGCAGUUGUCGCC	1	no match			

4481	UAGAUCUUCACCGUUGACUAGCCCCUA	1	2	Su(Ste)		Su(Ste)
4482	UAUGCAAUAUUUGUUGCGCCCCGGG	1	0	transposon	LTR	opus
4483	AUACUCAUCGUCGUAGAACAAGCCCGAA	1	2	Su(Ste)		Su(Ste)
4484	UGUUUCACCGUUAGACGGCUCGGGC	1	1	AT-chX		AT-chX-B
4485	UAUAGAGUGAUGCCGAGUAUAAAAG	1	1	repeat		trf
4486	UAAUUUCCCGGAAAUGUGCAAACGAC	1	1	no annotation		
4487	GCAUCCUCAACUGUGAGGGAUCCUCAA	1	no match			
4488	UGUUUCAGCGUUAGACGGCUCGGGC	1	1	AT-chX		AT-chX-B
4489	UCUCAUCGUCGUAAAACAAGCCCAA	1	2	Su(Ste)		Su(Ste)
4490	AAACCUGAAGAUGAUCCUGAGUU	1	1	transposon	LINE	Helena
4491	UUCUCAUCGUCGUUAGAACAAGCCCGA	1	no match			
4492	CCAGGCCCAUCUGGUUGAACGUG	1	0	Su(Ste)		Su(Ste)
4493	UCUCAUCGUCGUAGAACUAGCCCGAAA	1	2	Su(Ste)		Su(Ste)
4494	AAGGGGAAAUGAAGAGGAUCCUCAA	1	no match			
4495	UUUUUAAAACAAUUGAAAGAUAAGACCCA	1	2	transposon	LTR	copia
4496	UAACGAGGUAGAUGACUCCCUGGAA	1	no match			
4497	UCUCAUCGUCGUAGAACAAGCCCCGACUGUAAACUCGGGGUCAAU	1	no match			
4498	AAAAACUGUGAGGGAUCCUCAA	1	no match			
4499	UUCCGAGGAACCAAUGGAUCAGCU	1	no match			
4500	AUCGUCUCGGGGAUGAAAUAGAUCUCCACCGUUGACUAGCCCCC	1	no match			
4501	UGAUACCGAGAAUGACAAUGUUUGA	1	1	transposon	LTR	gypsy3
4502	UCUGUACAAGCAAGACCACCACACCCG	1	0	transposon	LTR	gypsy12
4503	CGAAACCCAAAGGAUACUCGAAGACACCACGG	1	no match			
4504	UUUGACAAAAUAUGCCCUUCACCAUA	1	0	transposon	DNA	1360
4505	ACUCAUCGUCGUAGAACAAUCCCGA	1	2	Su(Ste)		Su(Ste)
4506	UGCCAAAUCAUUCACCCGGUCAAUAACC	1	0	transposon	LINE	Doc3
4507	AUCUCAUCGUCGUAGAACAAGUCCGAG	1	2	Su(Ste)		Su(Ste)
4508	AAAAACCUAGAAGAUGAUCCUGAGUCGC	1	no match			
4509	GCCCCGGUACUCUGUGGUGACCAACAACG	1	no match			
4510	UGACCCAGUCGGAUCUAUUGUACUACC	1	0	transposon	LINE	baggins
4511	UCUCGAUCGUCGUAGAACAAGCCCGCGUGUAACGUCGGGUCAAU	1	no match			
4512	GAUCUUCACCGUUGACUUUGCCCA	1	1	Su(Ste)		Su(Ste)
4513	AUAGAUCUUCACCGUUGACUUGCCCCC	1	1	Su(Ste)		Su(Ste)
4514	CAGCACACAGGAGGAGAGCAGAUGAGACGCA	1	no match			
4515	CAACACGAAGACCCAGUAUACCGUA	1	0	transposon	LTR	Idefix
4516	AUCUCAUCUCGUAGAACAAGCCCGA	1	no match			
4517	ACCGAUGAGCCAGUGGAGUUGCUUA	1	0	transposon	LINE	G2
4518	AAAAACAACUGUGAGGGAUCCUCAA	1	no match			
4519	AUCUCUACCGUCGUAGAACAAGCCCGA	1	no match			
4520	CGCACACAGAGGAGAGCAGAUGAGAGA	1	no match			
4521	GAAAUGAAGAAAGUACAAGGUUGUAU	1	no match			
4522	UACUUUUUGUAGGUGUACUGACCCAG	1	1	transposon	LINE	baggins
4523	CUCAACUGUGAGGGGUAUCCUCAA	1	no match			
4524	AUGUUUCAUCGUUAGACGGCUCGGAC	1	2	AT-chX		AT-chX-B
4525	UAUCAUCGUCGUAGAACAAGCCCCGA	1	no match			
4526	GACAUCGAGCGGUCAAUAACCCAUAG	1	0	transposon	Low_complexity	F-element
4527	UAGAUCUUCACCGUUGACUAGCCCAAA	1	2	Su(Ste)		Su(Ste)
4528	GGUAACGGGAGGGUAUGAUAUGGAGG	1	0	transposon	LINE	G2
4529	UGUUUCAUCGUUGACGACUCGGGC	1	no match			
4530	CAUCGUCGUAGAACAAGCCCGA	1	0	Su(Ste)		Su(Ste)
4531	AAAGACGUGCAACUCUACGGCCCGG	1	1	transposon	LTR	ninja-Dsim-like
4532	UAUAUCUUCACCGUUGACUAGCCCC	1	1	Su(Ste)		Su(Ste)
4533	UGACGACCCAGCAGCAGCUAUCGAA	1	0	no annotation		
4534	GAUCCUCAACUGUGAGG	1	1	coding gene		CG13506-RA|CG13506
4535	UUCGUGACGAGAACUCGAGCCGACGAAAAA	1	no match			
4536	CCCCGUACUCUGUGGUGACCAAGACC	1	0	transposon	LINE	baggins
4537	UAGAUCUUCACCGUUGACUAGUCCCCA	1	2	Su(Ste)		Su(Ste)
4538	UAGAUCUUCACCGCUGACUUGGCCCA	1	1	Su(Ste)		Su(Ste)
4539	GUCUCGGGAUGAUCUCAA	1	no match			
4540	AUACUGUUUGUAGGUGUACUGACCCAGCUGUA	1	no match			
4541	AGGUGAAUACUCGAGCGCCAUCGAGA	1	1	coding gene		CG7185-RA|CG7185
4542	CUGACGAAUACCAUUGGGGGUAGAGCA	1	no match			
4543	UCUAGGCUGAGCGGAGAACCAGAGGU	1	1	transposon	Low_complexity	F-element
4544	CCUCAACUGUGAACUGUGAGGUCAA	1	no match			
4545	GAGAGGCGCCGUAGUUACCACCUC	1	1	no annotation		
4546	ACCUCACGACAUCGAGCGAUUGC	1	0	coding gene		CG40748-RB|CG40748
4547	CAACUGUGAGGGGUAUCCUCAA	1	no match			
4548	UGUUUUCAUCGUUAGACGGCUCGGCCG	1	no match			
4549	AUCUCAUCGUCGUAGAACAAGCCCCUA	1	no match			
4550	UCGGGAUGAAAUCCUCAACUGUGAGGGAUCCUCAA	1	no match			
4551	GAUUGCCAUAUUGCGGGAUUAUCUC	1	0	coding gene		CG12717-RA|CG12717
4552	UUCGCAAAAGUUGACCGAACUUGAUUA	1	0	ncRNA		rRNA
4553	CCCUUUCCUGGCUAGGUUGUCCG	1	0	transposon	LINE	baggins
4554	CCUCGAACUGUGAGGGAUCCUCAACGUGUAACGUCGGGUCAAU	1	no match			
4555	AUCUCAUCGUCGUAGAACAAGCCCCGAGCUGUAACGUCGGGUACA	1	no match			
4556	UGAUACCGAUAAUGACAAUGGUUGA	1	1	transposon	LTR	gypsy3
4557	AACUCUAAGCGGUGGAUCACUCGGC	1	0	ncRNA		rRNA
4558	UGUUUCAUCGUUAGACGGCUCCAG	1	2	AT-chX		AT-chX-B
4559	ACCUCAUCGUCGUAGAACAAAGCCCCGA	1	no match			
4560	GCGGAACCGGGCGGAACUGGAACAUCUCGUGUAACGUCGGGUCAA	1	no match			
4561	UUCCUGUUGAUGUGAGGGCCCCAA	1	0	transposon	LINE	G2
4562	UGCACCGAAAGUAUAUGAGAGGAGACU	1	0	Su(Ste)		Su(Ste)
4563	GAACUGUCGAACACCGGCUUCAGG	1	0	Su(Ste)		Su(Ste)

4564	UCUCAUCGUCGUAGACCAAGCCCA	1	2	Su(Ste)		Su(Ste)
4565	AGUGAGGGGGAUCCUCAA	1	no match			
4566	UCUCAUUCGUAGAACAAGCCCG	1	no match			
4567	GUUCUCAUCGUCGUAGAACAAGGCCCGA	1	no match			
4568	CUACAAUCUCCAAGCACGACCCCCGAUU	1	no match			
4569	AAAGGUGGAAAUGAAGAAAGUACACUGUAACUCGGGGUCAAU	1	no match			
4570	AUCGUCUCCGGGAUGAAAUCUCAUCGUCGUAGAACAAGCCCG	1	no match			
4571	UCUUGAUCCGAUGAUAUCACUUGGCC	1	0	no annotation		
4572	AUGUUUCAUCGUUAGACGGCCCGGGC	1	2	AT-chX		AT-chX-B
4573	UCCUCAACUGUGAGGGAAUCCUAAA	1	no match			
4574	UGUUUCAUCGUUAGACUUCUCGGGC	1	2	AT-chX		AT-chX-B
4575	AAUCCUCAACUUGAGGGAUCCUCAA	1	no match			
4576	UUCGGCUUUUCUGACGAGAACUCGAGC	1	no match			
4577	AAAACAACUGUGAGGGGAUCCUCAA	1	no match			
4578	UCUCAUAGUCGUAGAACAAGCCGA	1	no match			
4579	UGUUUCAUCCGUUAGACGGCUCCGGGCCUGUAACUCCGGGGUCAA	1	no match			
4580	UUAAAUGCCGGAAGCGAUCCCACUG	1	2	AT-chX		AT-chX-B
4581	AGGGGGCAACUCCAUCUCAGAGCGGA	1	no match			
4582	AACUGGUGAGGGGAUCCUCAA	1	no match			
4583	GAACCUCAACUGUGAGGGAUCCUCAA	1	no match			
4584	UAGAUCUCCCCGUUGACUAGCCCCA	1	no match			
4585	GCAUAUCAUUCACGCCACGCUCCGACC	1	0	Su(Ste)		Su(Ste)
4586	UACUAGCACUACCUAAAAUGACCCCUAC	1	0	coding gene		CG31550-RA|CG31550
4587	GCUGACUCUACGUGACGGAACCGGA	1	0	transposon	LINE	Rt1b
4588	AUCGUCUCGGGGAUGAAACAAGAUAUGACCAAAGCAAAGACUA	1	no match			
4589	ACUCAACUUGAGGGGAUCCUCAA	1	no match			
4590	GAUUGCCGGAGCUCUGUACUACGAUA	1	0	coding gene		CG33129-RA|CG33129
4591	GCACACAGAGGAGAGCAAGAGAGCA	1	no match			
4592	CUCAACUGUCAGGGAUCCUCAA	1	no match			
4593	UUUCUGACGAGAACUCGAGCCACGUA	1	1	no annotation		
4594	UGCAAUUUCCCGUGGGACUACA	1	no match			
4595	UAGGAUCUUCACCGUUGACUAGCCCCGA	1	no match			
4596	UACACACUCAUCAACCAAAGGACGUA	1	0	no annotation		
4597	UCGUCUAUCGUCGUAGAACAAGCCCGA	1	no match			
4598	ACCUCAUCGUCGUAGAACAAAGCCCGAC	1	no match			
4599	UGUUUCAUCGUUAGACGGCUCGGU	1	1	AT-chX		AT-chX-B
4600	CCCAUCGUCGUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
4601	CUGUGGGAAUUGGAUCCUUAUUAUCCCUCC	1	no match			
4602	AUCUCAUCGUCGUAGAACAAAGCCCG	1	no match			
4603	AAUGUAGAUAAGGGAAGGGGCAAAU	1	no match			
4604	UACUAGCUGGCUGAUGAUCCGAUAU	1	no match			
4605	UUACGAGUAAUGGUCUUCGGGGCGU	1	1	coding gene		CG12717-RA|CG12717
4606	UUCGGCUUUCUGACGAGAACUCGACGC	1	2	no annotation		
4607	UUCUUGACAACCCGAUCGGACCUCAC	1	0	transposon	LTR	invader4
4608	AUCGUCGUCGGGAUAGAAAUCAUCGUCGUAGAACAAGCCCGA	1	no match			
4609	UGUUUCAUCGUUAGACGGCUCGGGUG	1	1	AT-chX		AT-chX-B
4610	UUCCAUCGUCGUAGAACAAGCCCUA	1	2	no annotation		
4611	CUCGCAAGAAAACUAAUCCGAGUAU	1	0	transposon	LINE	Doc
4612	UCUCAUCGUCGUAGAACACGCC	1	1	Su(Ste)		Su(Ste)
4613	UCGACAUUUUGCCAGUUCACUCGU	1	0	no annotation		
4614	CAAGAUAUGACCUAAGCAAAAGACAC	1	2	transposon	LINE	F
4615	CGAAAUAGAUCUUCACCGUUGACCUUGGCCC	1	no match			
4616	CUUUCUCAUCGUCGUAGAACAACGCCCGA	1	no match			
4617	GGUGGAAAUGAAGAAAGUACA	1	2	no annotation		
4618	UUCGAGAGCAUAUAAUCCCCGUAGGU	1	no match			
4619	UGAAACCCACUUUUCAGAUCGAGCAC	1	0	transposon	LTR	HMS-Beagle
4620	CAAGAUAUGACCUAAGCAAAGACA	1	1	transposon	LINE	Cr1a
4621	UAAACUGUGAGGGGAGUCCUCAA	1	no match			
4622	UUAAACAUCCUAUGAGGCUACGG	1	0	transposon	LINE	Cr1a
4623	GACUAGCUGGCUGAUGAUCCGAUA	1	2	Su(Ste)		Su(Ste)
4624	UGUACGGACCUGGAGAACAACCAGAG	1	1	transposon	LTR	ninja-Dsim-like
4625	UGUUUUCAUCGUUAGACGGCUCGGGCC	1	no match			
4626	UAGCUGGCCGAGGAUCCGAUUUU	1	0	Su(Ste)		Su(Ste)
4627	GCAACAAGAGCGUGUCCCGGGCAGACCU	1	0	coding gene		CG10229-RA|katanin-60
4628	UCUCAUCGUCGUAGAACAAGCCCCGACUGUAACCUCGGGGUCAAU	1	no match			
4629	AUCGUCCUCGGGAUGAAAUCUCAUCGUCGUAGAACAAGCCCGA	1	no match			
4630	UUAUACCCGGGUGAUUACCACUAUU	1	1	Su(Ste)		Su(Ste)
4631	UGUUUCAUCGUUAGACGGCUCGCC	1	2	AT-chX		AT-chX-B
4632	AUCUCAUCGUCGUAGUCAAGCCCGA	1	no match			
4633	UUCUUCAUCGUUAGACGGCUCGGGCCUGUAACGUCGGGUCAAU	1	no match			
4634	UCCGAGGAACCAAUUGAUACAGCUG	1	1	Su(Ste)		Su(Ste)
4635	UCGUCGGGAUGAAAAUCCUCAA	1	no match			
4636	AUCGUCUCGGGGAUGAAAUCUCAUCGUCGUAGAACAAGCCCA	1	no match			
4637	UGUUUCAUCGUUGACUAGCCCCA	1	no match			
4638	AUCUCAUCAUCGUAGAAACAAGCCCCG	1	no match			
4639	AAAUCUCAUCGUCGUAGAACAAGCCCCGA	1	no match			
4640	CACUGCAUUCGCGUCGAGGCCCAG	1	0	transposon	LINE	R1-element
4641	GUAUGACCCCUGGUACUGAAAUUCAG	1	1	transposon	DNA	Bari1
4642	UGACCGUUGACCAUUAUCGUCUCA	1	0	transposon	LTR	opus
4643	AUCGACCGGCACGAUUCUAUCGAGU	1	2	transposon	LTR	G4
4644	UUCUCAUCUUCGUAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
4645	UUCUCAUCGUCGUAGAACAAGCCCCA	1	1	Su(Ste)		Su(Ste)
4646	UGUUUCAUCGUUAGACGGCUCGGGCCUGC	1	no match			

4647	UGUUUCAUCGUAUACGGCUCGGG	1	no match			
4648	GUUCAAGCCCGGGUAUUCUGAUUUA	1	no match			
4649	UCGUCAGUCGUACGUAGAACAAGCCCGGACUG	1	no match			
4650	UCUCAUCGUCGGAGAACAAGCCCGA	1	1	Su(Ste)		Su(Ste)
4651	UCUACAUCGUCGUAGAAGAAGCCCGACUGUAACGUCGGGUCAAU	1	no match			
4652	UUCUAAGGUUACCCCAGGAUGCUGU	1	no match			
4653	UAUAAUCGCCCCGAAGGUUUGCCCCUCUG	1	no match			
4654	CAGGCUACACAGAGGAGAGCGAGAUGAGAGCA	1	no match			
4655	UUCAAGGUUACCCAGGAUGCUGU	1	no match			
4656	UACUCGAUCGUCGUAGAACAAGCCCGCGUGUAACGUCGGGUCAAU	1	no match			
4657	GUAGAUCUUCACCGUUGACUAGCCCCC	1	1	Su(Ste)		Su(Ste)
4658	UUCGUGACGAGAAACGUCGAGCCGCGAAAAACUG	1	no match			
4659	CGAAAUGUUUCAUCGUUAGACGGCCUCGGC	1	no match			
4660	UCCACCGGGUUUUUGGCACUUUGCCGC	1	no match			
4661	UAGAUCUUCCCCGUUGACUAGCCCCA	1	1	Su(Ste)		Su(Ste)
4662	AUCGUCUCGGAUAAAUGUUUCAUCGUUAGACGGCUCGGG	1	no match			
4663	CCGAAGAAGAUGAACCCCCUUAAAGU	1	0	coding gene		CG1338-RA|hydra
4664	GUGUUUCAUCGUUAGACGGCUCGGGCU	1	2	AT-chX		AT-chX-B
4665	UCGUCAUCGUCGUAGAACAACGCCCGGA	1	no match			
4666	CAACUGUGAGGGGAUCCUCUAA	1	no match			
4667	UUGACCACACGAACGAAUCCGAGAAA	1	1	coding gene		CG12717-RA|CG12717
4668	CUCUCAUCGUCGUAGAACGACGCCCGA	1	no match			
4669	ACAAACUGUGAGGGGAUCCUCAA	1	no match			
4670	GAUUCUGCCGGAACGAGGGACCCAUG	1	0	coding gene		CG7131-RA|CG7131
4671	UUCUCAUCGUCGUAGAACAAGCCG	1	1	Su(Ste)		Su(Ste)
4672	AUCGUCAUCGUCGGUAGAACAAGCCCGGA	1	no match			
4673	CCGAUCUCCUGAGUCGGACUUCGAGC	1	0	transposon	LINE	R1-element
4674	AAGAUAUGACCAAAGCAAAGACACC	1	1	transposon	LINE	Cr1a
4675	UGAAAUCCGAGUUAAUAUGGCACCAA	1	0	no annotation		
4676	UUUUUAAAACAAUUGAAAGAUAAGACCC	1	2	transposon	LTR	copia
4677	AUUUCUAGUCGAAUAAAUGAUCCGUC	1	1	transposon	LTR	17.6
4678	AGAUCUUCACCGUUGACUAGCCCCCA	1	1	Su(Ste)		Su(Ste)
4679	UCCACGGACCUGGAGAACAACCAGA	1	1	transposon	LTR	ninja-Dsim-like
4680	UAGGUUAUGGGCCCAGUAAACAUUUAG	1	0	ncRNA		alphagamma-element
4681	GUUUAUCCAAGACAGGUUGUGC	1	no match			
4682	GAGGACUUGAGCGAUUUAAGUGGC	1	1	Su(Ste)		Su(Ste)
4683	UGUUUCAUCGUUAGACGGCUCGGGCCUGUAACUCGGGGUCAAUAU	1	no match			
4684	UAGACCUUCACCGUUGACUAGCCCCA	1	1	Su(Ste)		Su(Ste)
4685	UUUGAAUGCCCUAAUUACUUCUCCU	1	0	no annotation		
4686	UCUCAUCGUCGUAGAACAAGACCGA	1	1	Su(Ste)		Su(Ste)
4687	UAGUCUAGAGACCGAAGAUAGUUGA	1	no match			
4688	CCCCUCUGGUGAUAAAUUGAUUG	1	0	transposon	DNA	Tc1-2
4689	AAAAUUUCUCAUCGUCGUAGAACAAGCCCGA	1	no match			
4690	AGUAUGACCCCUGGUACUGAAAUG	1	1	transposon	DNA	Bari1
4691	GCCGUCGAGCAUCGCUACCUGGAUU	1	1	coding gene		CG6210-RA|wls
4692	AUCUCAUCGUCGUAAGAACUACGCCCGA	1	no match			
4693	UAGAUCUUCACCGUUGACUUGCCCGA	1	1	Su(Ste)		Su(Ste)
4694	CACACAGAGGAGAGAGAUGAGAGCA	1	no match			
4695	CUUAACUUCGGGAUAAGGAUUGGCUCU	1	1	ncRNA		rRNA
4696	GAAGCGACAUGCUCGAGAUGACAACC	1	0	transposon	LTR	micropia
4697	UAAAAUGCCAGAAUUUUAUUGGGGG	1	2	transposon	LINE	baggins
4698	CUCAACUCGUGAGGGGAUCCUCAA	1	no match			
4699	UUAAGUCCCGUUACUUGUGCCCAG	1	0	transposon	LINE	G6
4700	UAAAAGAAAUCCUGACAAGAUCCCGGG	1	no match			
4701	AAAUAACAAGAUGCAUACGAUGUGGCUCCA	1	1	transposon	DNA	1360
4702	UCUCAGUCGUCGUAGAACAAGCCCGACG	1	no match			
4703	UGUCCACCAAAGGCAUGCCCGAUG	1	0	transposon	LTR	opus
4704	UGAUACCGGGAAUGACAAUGGUUGA	1	1	transposon	LTR	gypsy3
4705	UCUCAUCGUCGUAGAACAAGCCAA	1	2	Su(Ste)		Su(Ste)
4706	UCGUUUCCGGCGAGCGUUGAUUAA	1	no match			
4707	UCUCAUCGUCGUAGAACAAGCCCAACUG	1	no match			
4708	AAAGAGAGUAAUUUUGGCCGUCACC	1	0	transposon	LINE	Cr1a
4709	UAAAUGCCAGAAUUUUUAUUGGGG	1	2	transposon	LINE	baggins
4710	GUAUGACCCCUGGUACUGAAAUGCAU	1	1	transposon	DNA	Bari1
4711	UAGUUUCAUCGUUAGACGGCUCGGGU	1	2	no annotation		
4712	GGCCCGAAAACACGAGAGCAGAGC	1	1	transposon	LTR	opus
4713	GUUCACCGGGUUUUGGCACUUUG	1	0	transposon	LINE	R1-element
4714	UAGAUGUGGCCUGAAGUUGAAUUU	1	0	transposon	LINE	F
4715	CCUCAUCGUCGUAGAACAAGCCCGAGG	1	1	Su(Ste)		Su(Ste)
4716	UUCUCAUCGUCGUAGAACUAGCCCUAA	1	2	Su(Ste)		Su(Ste)
4717	UAAAUGCCAGAAUUUUAUUGGUGUUC	1	1	transposon	LINE	baggins
4718	UCCAAAGGACUCGAAAAAACUCUUGA	1	no match			
4719	UCUCAUCGUCGUAGAACUAGCCCC	1	1	Su(Ste)		Su(Ste)
4720	UAUGCGACGAGUACGCUGUUUGGGC	1	0	transposon	LTR	G4
4721	GGAUCCUCAACUGUGACCUCAA	1	no match			
4722	GACUAGCUGGCCGAGGAUCCGAUA	1	0	Su(Ste)		Su(Ste)
4723	UAGAUCUUCACCGUUGGCUAGCCCCA	1	1	Su(Ste)		Su(Ste)
4724	AUAAAAGAAUCCUGACAAGAUCCCGGG	1	no match			
4725	UCGUCAUCGUCGUAGAACAAGCCCGGAG	1	no match			
4726	UCAAGUCCUUCUGGUGCAACCGAAA	1	1	Su(Ste)		Su(Ste)
4727	AAAACCUGAAGAUGGAUCCUGUAGUCGC	1	no match			
4728	GGAUCCUCAACUGUGAGGGCAUCCUCAA	1	no match			
4729	CCUGCUGACGGGGUCCGAAAACAC	1	1	transposon	LTR	opus

4730	CUAGAUCUUCCACCGUUGACUAGCCC	1	no match			
4731	AUCUCAUCUCGUAGAACAAGCCCG	1	no match			
4732	GUAAUGACCCCUGGUACUGAAAUGCAG	1	no match			
4733	AUCGCUCGGGAUGAAAUUCGCCUUUCUGACGAGAACUCGAG	1	no match			
4734	GAGAUGGCUGCCGGGUACUACCUCA	1	0	transposon	LINE	G2
4735	UCUGAGAUUGUGAAUGAUAAAGAUUGG	1	0	transposon	LTR	invader3
4736	AUCGUCUCCGGGAUCGAAAUGUUUCAUCUUUAGACGGCUCGGG	1	no match			
4737	UUCAACGAGUCGUCACUGGGAGC	1	0	transposon	LINE	accord2
4738	UCUCAUCGUCGUAGAACAACGC	1	2	Su(Ste)		Su(Ste)
4739	UAAAUGACCCGUCAAUAACGAAGACA	1	1	transposon	LTR	RR50207
4740	UCUCAUUGUCGUAGAACAAGCCCUA	1	2	Su(Ste)		Su(Ste)
4741	CGUUUCAUGUUAGACGGCUCGGGC	1	no match			
4742	GGACCCAACUGUGAGGGGAUCCUCAA	1	no match			
4743	AUCUGACGAGAACUCGAGCCACGUAGC	1	2	no annotation		
4744	UAUCAUCGUCGUAGAACAAGCCCG	1	0	Su(Ste)		Su(Ste)
4745	UCGACCGACCAGCUGUGCGAACAGU	1	0	transposon	LTR	invader3
4746	AUUGACCGAGUACAGACGGGCGUUGUUCGUACGAACGGAUGAGAG	1	no match			
4747	UACUCAUCGUCGUAGGAACAAGCCCGA	1	no match			
4748	UAGACCUAGAGCAUCCAAAAAACUUUGUA	1	0	transposon	LTR	gypsy8
4749	UCUCGUCGUCGUAGAAGAAGCCCGA	1	2	Su(Ste)		Su(Ste)
4750	CAAUACUCGUCUCGGGACUGAAAAAAAAAAAAAAACGGAUCCCUCUA	1	no match			
4751	UAUUAUUAUACCCAGGUGAUUACC	1	0	Su(Ste)		Su(Ste)
4752	AUGUUUCAUCGUUAGACGGCUCGG	1	1	AT-chX		AT-chX-B
4753	UUCUCGACGUUGACCAGCCUCCUGUA	1	0	transposon	LTR	ninja-Dsim-like
4754	UAUGACCGAAAUUCGGAAAACGGA	1	2	repeat		SAR_DM
4755	UCCAAAAGAAAUGAUGACCCUUGCAUU	1	no match			
4756	UUUAUACCUGAGCCAGUUUGUUUG	1	1	transposon	Low_complexity	F-element
4757	CAAGAUAUGACCAAAGCAAAACU	1	2	transposon	DNA	1360
4758	AUCAACUGUGAGGGAUCCUCAA	1	no match			
4759	AUCUCAUCGUCGUAGGACAGGCCCGA	1	no match			
4760	UGUUUCAUCGUUAGACGGCUCCGGGA	1	no match			
4761	AAUUUCGAGUCGAAUAAAUGAUCCGUC	1	1	transposon	LTR	17.6
4762	UAAACUCUGGACAGGACGUUGAUGA	1	0	transposon	Simple_repeat	I
4763	GAUUCCGGAGAGGGAGCCUGAGAAA	1	0	ncRNA		rRNA
4764	UCUCAUCAUCGUGAACAAGCCCGA	1	no match			
4765	GGAUCCUCAACUGUGAGGUCAA	1	no match			
4766	UCGGCCCGAACCCUCUGGUGUG	1	no match			
4767	UCCUUCUGAAGAGUUGUUGGCACG	1	0	transposon	LTR	GATE
4768	UAAAUGCCAGAAUUUUUAUUGGGGUUUC	1	no match			
4769	UUCGACCGAACACUUUGAACCAAGUA	1	1	Su(Ste)		Su(Ste)
4770	UCUGUUUCAUCGUUAUACGGCUCGGGC	1	1	AT-chX		AT-chX-B
4771	AACAAGAUGCAUACGAUGUGGCUCCGA	1	2	transposon	DNA	1360
4772	UGUUUCAUCGUUAGACGGCUCGGGCCUGUAACUGGGUCAAU	1	no match			
4773	UCCCAGGCCCAUCUGGUUGAACGUGU	1	2	Su(Ste)		Su(Ste)
4774	UCAAAGCGUUGACUACUUCACGAGGAA	1	2	transposon	LTR	opus
4775	UAAAUGUUUCAUCGUUAGACGGCUCGGGC	1	no match			
4776	AUCGUCUCGGGACUGAAACGUUGACCAAUUUCCGAACACUACGC	1	no match			
4777	UUACAGUGAAAUGAAGAAAAGUACA	1	no match			
4778	UGGUACUUUUUCCGAUGAGCCAGU	1	no match			
4779	AUCUUCAUCGUCGUAGAACAAGGCCCGACUGUAACGUCGGGUCAA	1	no match			
4780	UAUCAUCGUCGUAGAACUAGCCCGA	1	1	Su(Ste)		Su(Ste)
4781	UCCCAGAUCCCAUUUUAGCAUUAGC	1	0	transposon	LINE	G2
4782	AAAGAACCCGAUUCCUCUUCCAUGA	1	0	transposon	LINE	Doc4
4783	UUUCUCAUCGUCGUAGAACAAAGCCCCG	1	no match			
4784	GGCGCCCGAUAAAGCAGCAGAUGGAU	1	0	transposon	LINE	TART-A
4785	UCUCAUCGUCGUAAGAACAAGCCCCGAGG	1	no match			
4786	AUCUCAUCGUCGUAGAACUAGCCCCG	1	no match			
4787	UCUCAUCAUCGUAGAACUAAGCCCGA	1	no match			
4788	UUGAUUCCGAGGAACCAAUUGAUCCA	1	0	Su(Ste)		Su(Ste)
4789	AUCUCAUCGUCUUAGAACAAGCCCGAA	1	2	Su(Ste)		Su(Ste)
4790	GCCCCGUACUCUUGGUGACCAA	1	no match			
4791	GAAUAUCGUCUCGGGAGUGAAAUCUCAUCGUCGUAUAACAAGCCC	1	no match			
4792	CCUCAACUGUGAGGGGACUCCUCAA	1	no match			
4793	CCCAGUAAAACCCAACCAGAUCAAA	1	1	transposon	LTR	copia
4794	AACGAGUGACGAAUUGCACCCAGU	1	0	no annotation		
4795	UCUCAUCGUCGUAGAACUAUCCCGA	1	1	Su(Ste)		Su(Ste)
4796	CCUCAUCUCGUAGAACAAGCCCGA	1	no match			
4797	UCCGAGGAACCCAUCGAUCCAGCU	1	1	no annotation		
4798	UUGAUACGAAUGAGACCCGCAGUAAAG	1	0	repeat		trf
4799	UCUCAUCGUCAUAGAACAAGCCCC	1	2	Su(Ste)		Su(Ste)
4800	UCUCAUCGUCGUAGAGCAAGCCCAA	1	2	Su(Ste)		Su(Ste)
4801	GGUCCCGUAGUGUAGUGGUUAACAUGCCUGC	1	no match			
4802	UCUCAUCGUCGUAGAACAGCCCAA	1	no match			
4803	ACCGAGCUCAAGUCCAUAGGAUAUGC	1	0	coding gene		CG17170-RB|su(f)
4804	AGAAAAUCCUCAACUGUGAGGGGAUCCUCAA	1	no match			
4805	AUCAUCAACUGUGAGGGAUCCUCAA	1	no match			
4806	UCGGAUUUCCGACGCCUCAAGAUCCCGC	1	no match			
4807	CAAGAUAUGACCACAGCAAAGACAC	1	1	transposon	LINE	Cr1a
4808	UCCUCAACUGUGGGGGAUCCUCAA	1	no match			
4809	CGAAAUCUCAUCGUCGUAGAACGAAGCCCGA	1	no match			
4810	UGCUCUCGAAUGUAUGCCCGGAUAUAC	1	no match			
4811	UCUCAUGUCGUAGAACCAAGCCCCGACUGUUAACUCCGGGGUUCA	1	no match			
4812	ACAAUGGGCGCAAGCCUGAUCCAGCCAUG	1	no match			

4813	UCUCAUCGUCGUAGAACAACGCCCGAU	1	no match			
4814	AAAACCUGAAGAUGAUCCUGAGUCGU	1	0	transposon	LINE	Helena
4815	UAGAUCUUCACCGUUGACUUCGCCCA	1	1	Su(Ste)		Su(Ste)
4816	UCUGUCUUCAUCACUACGGGCCCAG	1	0	transposon	Low_complexity	F-element
4817	GUGCCAGGCCCGUAUAACGUUAAUGAUU	1	0	ncRNA		rRNA
4818	CGUUUCAUCGUUAGACGGCUCAGU	1	no match			
4819	CGAAUGUUUCAUCGUUAGACGCCUCGGGC	1	no match			
4820	GGAUCAUCAACUGUGAGGGCAUCCUCAA	1	no match			
4821	GUAGAUCUUCCCGUUGACUUGGCCC	1	no match			
4822	UCUCAGUCGUCGUAGAACAAGCCCGA	1	no match			
4823	ACCUCAACGUGAGGGAUCCUCAA	1	no match			
4824	UCUCAUCAUCGUAGAACAAGCCCGU	1	1	Su(Ste)		Su(Ste)
4825	UCGUCUCGGGUAAGGAUCCUCAA	1	no match			
4826	CUGGUAUAGCACACUUUUGUAUCGGC	1	1	coding gene		CG41580-RA|CG41580
4827	UUCUCAUCGUCGUAGAACAAGCCCGAGG	1	0	Su(Ste)		Su(Ste)
4828	UUCCGAUGUUUUGAAACUGCGUGACCCA	1	0	transposon	LTR	flea
4829	CAACAGAGGAGAGCAGAUGAGAGCA	1	no match			
4830	UCCGAUCUCCUGAGUCGGACUCCGA	1	0	transposon	LINE	R1
4831	UGGACUUUUUCCGUAUGAGCCAGU	1	2	transposon	LINE	G2
4832	AUCUCAUCGUCUUAGAACAAGCCCGA	1	2	Su(Ste)		Su(Ste)
4833	CAGGCACCGAGAAGGGAGAGCAGGAUGAGAGCA	1	no match			
4834	CCCGAAGCGGACCAGCCCAAUUAUU	1	no match			
4835	UGCAAUCUCCGAGCAAGACCCCGA	1	1	transposon	LINE	Circe
4836	AUCCUCAACUGUGAGGGGAUCAUCAA	1	no match			
4837	AUUAUCCGUAAGGGUUGACAUAUUGCA	1	0	transposon	LTR	DM88
4838	GAUGAGCCAGUGAAGUUGCUUAAGG	1	0	transposon	LINE	G2
4839	AUCCUCCUGUGAGGGAUCCUCAA	1	no match			
4840	AUCGUCUCGGGGAUGAAAACGUUUCAUCGUUAGACGGCUCGGGC	1	no match			
4841	AUGAUCCUCAACUGUGAGGGGAUCCUCAA	1	no match			
4842	GAAAAGAGAGCAAUUUUGGCCGU	1	0	transposon	DNA	1360
4843	AGCUACACACGAGGAGAGCACGAUGGAGAGACA	1	no match			
4844	AUGUUUUCAUCGUUAGACGGCUCGGA	1	no match			
4845	UUCUCAUCGUCGUAGAACAAGCCCGGA	1	1	Su(Ste)		Su(Ste)
4846	GACUCACCAGAUGAUAUCAAUGUCCCAA	1	0	transposon	LTR	invader6
4847	UAACAACGUCAACCUGAGGCAUUGGA	1	0	transposon	LTR	diver2
4848	GCCCGGCUAACUCAGUCGGUAGAGCA	1	1	ncRNA		tRNA
4849	UACGGAACGAGAGACGCGACUGGAU	1	0	coding gene		CG15288-RB|wb
4850	UCGUCGUCUACCCAUGGAAUCCCGA	1	0	transposon	LINE	G2
4851	GCCCGUACUCUGUGGUGACCGACA	1	2	transposon	LINE	baggins
4852	UCUCAUCGUCGUAGAAGAACGCCCGUA	1	no match			
4853	AUCGUCAGUCGUACGGUAGAAACAAGCCCGGACUG	1	no match			
4854	UCCGAUGAACCAAUUGAUCCAGCU	1	0	Su(Ste)		Su(Ste)
4855	GUCUCAGGAUGAAAAUCCUCAA	1	no match			
4856	GCCAAAUGAACCUUGGUUGUCUUGAGA	1	0	ncRNA		pseudogene
4857	UGACCUCGUGUGGCCAUACUCUUGG	1	0	transposon	Low_complexity	F-element
4858	UCCAUGCAGCAGAAGCGACAGAGUU	1	0	transposon	LINE	baggins
4859	UCCGAGGAACCAAUUGAUCUAGCU	1	1	Su(Ste)		Su(Ste)
4860	AACUUGAGGGGAUCCUCAA	1	no match			
4861	AUACAAUCUCCGAGCAAGACCCCGA	1	1	transposon	LINE	Circe
4862	GUCUCGGGAUAAAAAUCCUCAA	1	no match			
4863	UCCUCAACUGAGGGGAUCCUCAA	1	no match			
4864	ACAAGAUAUGACCUAAGACAAAGACCA	1	no match			
4865	UAUCAUGUGAAAUUUGUGAACCCUGU	1	0	transposon	LINE	Cr1a
4866	UUCUCGACGUUGACCAGUCUCCUGU	1	1	transposon	LTR	ninja-Dsim-like
4867	AAAACCUGAAGAUGAUCCUGAGUCAC	1	1	transposon	LINE	Helena
4868	UGAACCCGGCAAUGCCCCGUAAGAAGG	1	1	transposon	LINE	INE-1
4869	AUCGUCUCGGGGGCG	1	1	coding gene		CG14194-RA|CG14194
4870	AGAUAAGGGAAGUCGGCAAAAUAGAU	1	2	ncRNA		rRNA
4871	ACACAUGAUCACUUCAGACGUACCCGU	1	no match			
4872	AUCGUCUCGGGAGUGAAAUACUCAUCGUCGUAGAACAAGCCCG	1	no match			
4873	UAGCUUGCACCAUCACCAUCACCA	1	2	coding gene		CG11505-RA|CG11505
4874	UCGUCGGCCUGGAUCACGGACAUU	1	0	transposon	LTR	ninja-Dsim-like
4875	UGGAUGCCGUGAGACAGUUCCGAUGU	1	0	transposon	LTR	ninja-Dsim-like
4876	UAGAUCUUCUUCGUUGACUAGCCC	1	2	Su(Ste)		Su(Ste)
4877	ACAAGAUAUGACCAAAAGCAAAAGACAA	1	no match			
4878	UGUUUUCACCGUUAGACGGCUCGGGC	1	no match			
4879	UCGGCAUCGUCGUAGAACAAGCCCG	1	2	Su(Ste)		Su(Ste)
4880	UACAAUCUCCGAGCACGUCCCCGAUAU	1	1	transposon	LINE	Circe
4881	AUGGUUCAUCGUUAGACGGCUCGGGC	1	2	AT-chX		AT-chX-B
4882	GUCUCGAGAUGAAAAUCCUCAA	1	no match			
4883	CAGCACACAGAGGAGAGCAGAUGGAGCACUGUAACUCGGGGUCAA	1	no match			
4884	AACCGGAGAAGACGUUGAAUGGAUCGGC	1	1	no annotation		
4885	UAGAUCUUUCACCGUUUGACUAGCCCCCA	1	no match			
4886	UCUCAGUCAUCGUAGAACAAGCCCGAU	1	no match			
4887	ACAAGAUAUGACCAAAGCAAAGACUA	1	2	transposon	DNA	1360
4888	UCUCAUCGUCGUAGAACAACGCCCGAA	1	no match			
4889	UCUUCCGACAGACGAAGAAGUGGC	1	0	transposon	LTR	diver2
4890	AUGCUCUCGAAUGUAUGCCCGAUAUA	1	0	transposon	LINE	R1-element
4891	UAUGACCAGCUUGGACCGAACACU	1	0	Su(Ste)		Su(Ste)
4892	AUCUCAUCGUCGUAGAACAAGCCCU	1	2	Su(Ste)		Su(Ste)
4893	UAACGAUAUCCACAUUCGCUGAUGACAC	1	0	transposon	LINE	Doc3
4894	AGAAAAAGGUGGAAAUGAAGAAAGUACA	1	no match			
4895	CUCUCAUCGUCGUAGAACAACGUCCGA	1	no match			

4896	GUUUCAUCGUUAGACGGCUCGGGUC	1	2	AT-chX		AT-chX-B
4897	UCUCUCUCGUAGAACAAGCCCGA	1	no match			
4898	GGUAUAGCACACUUUUGUAUCGGC	1	0	coding gene		CG41580-RA|CG41580
4899	UCAGAUACCCGUUACUCAGCUAGUA	1	0	transposon	LINE	INE-1
4900	UUUUACGUGGACGAGUUGCUCGAU	1	0	transposon	Low_complexity	F-element
4901	UUCGUCAUCGUCGUAGAAGAAGCCCGG	1	no match			
4902	CUUUGACCAAUUUCCGAACAUACG	1	1	transposon	LTR	DM88
4903	CUCAACUGUGAGAUCCUCAA	1	no match			
4904	UUGUACACACCGCCCGUCGCUACU	1	0	ncRNA		rRNA
4905	CCCAACUAUGAGGGGAUCCUCAA	1	no match			
4906	UGCAUGAAUUCCUCUGGGCCGAUAGC	1	0	transposon	LINE	R1-element
4907	UGUUUCAUCGUUAGACGGCUCGGGCCUGUAACGUCGGGUCAAU	1	no match			
4908	GCACACAGAGGAGAGCACGAUGAGACGA	1	no match			
4909	CGAAAUCAACUGUGAGGGGAUCCUCAA	1	no match			
4910	UCUCAUCGUCGUAGAAGCAGCCCG	1	2	Su(Ste)		Su(Ste)
4911	CAGACACAGAGGAGAGCAGAUAGAGCA	1	no match			
4912	GCAACUGUGAGGGGAUCCUCAA	1	no match			
4913	UAAAGAAUCCUGACAAGAUCCGG	1	1	transposon	LTR	invader1
4914	CAGGCACCAGAGGAGAGCAGGAUGAGAGCA	1	no match			
4915	CUGGUGAACGAGUGAACUGGCAACAA	1	no match			
4916	GGACUCAACUGUGAGGGGAUCCUCAACUGUAACUCGGG	1	no match			
4917	GCUGAGAAAAAAACGAAGUAUAACGAA	1	0	transposon	LINE	Doc
4918	UAGAUCCUCACCGUUGACUAGCC	1	1	Su(Ste)		Su(Ste)
4919	UAAGGAUGAUGAUACCGAUUAAAGG	1	0	no annotation		
4920	AUCCUCAACUGUGAGGGAUCCUCUAA	1	no match			
4921	AUGUUUCAUCGUUAGACGGCUCGGGCG	1	1	AT-chX		AT-chX-B
4922	UCCGAGAAUGACUUGAGGUAGUUUU	1	0	transposon	LINE	G2
4923	UCUGGUUGAACGUGUCCUGAAAGU	1	0	Su(Ste)		Su(Ste)
4924	UCUCAUCGUCGAGAACGAGCCCGA	1	no match			
4925	UCUCAUCGUUUAGAACAAGCCCGA	1	no match			
4926	GGAUGAAAUGAUACCGAGAAUGACAAUGGUUGA	1	no match			
4927	CCUCAACUGUGAGGGGAUCCCAA	1	no match			
4928	AGAUAUGACCAAAGCAAAGACAC	1	0	transposon	LINE	Cr1a
4929	GGAUCCUCAACUGCAA	1	1	coding gene		CG4356-RA|mAcR-60C
4930	UUCGAGAAGCGUUUGAUCGAGUGUG	1	0	transposon	LINE	BS3
4931	GUCCUAGCCUCCAAGACGCGCC	1	0	coding gene		CG5174-RA|CG5174
4932	ACGAAGAUGAUUCUGGCCCAGAAAA	1	0	transposon	LTR	invader4
4933	UCUCAUCGUCGUAGGAGAAGCCCGA	1	2	Su(Ste)		Su(Ste)
4934	AUCGUCAUCGUCGUAGAACAAGCCCCGA	1	no match			
4935	UCGAACUGUGAGGGGAUCCUCAA	1	no match			
4936	ACUCAUCGUCGUAGAACAAGCCCA	1	2	Su(Ste)		Su(Ste)
4937	UUAACUACAUGGCGACCGUGACAGGACCCUA	1	1	transposon	LTR	mdg1
4938	UAUGCAAUAUUUGUUGCGCCCUGGG	1	1	transposon	LTR	opus
4939	UUAUAUUGGCUUAGUGCGCUCACAG	1	0	transposon	LTR	Dm88
4940	UCUCAUCGUCGUAGAACUAGUCCCGA	1	no match			
4941	AGUAGAUCUUCACCGUUGACUUGGCC	1	0	Su(Ste)		Su(Ste)
4942	AUCGCUCGGGAUGAACAGCACACAGAGGAGAGCAGAUGAGAGCA	1	no match			
4943	GCACGACAGAAGGAGAGCAGAUGAGAGCA	1	no match			
4944	UGUCCUACUGAUGACGAUGACCCUU	1	0	coding gene		CG14269-RA|CG14269
4945	GCCCCGUACUCUGUGGUGACCAACUA	1	2	transposon	LINE	baggins
4946	UCUCAUCGUCGUAGAACAAUCCCCGA	1	2	Su(Ste)		Su(Ste)
4947	UAAAGUAAGCCCGAAGAUUGUCCAAU	1	2	transposon	LTR	opus
4948	ACACAUCUUCACCGUUGACUUGGCCA	1	no match			
4949	CGUUGACCAAUUUUCCGAACAUACGC	1	no match			
4950	UAACAAGAUGCAUACGAUGUGGCUCCC	1	1	transposon	DNA	1360
4951	AGACUAGCUGGCCGAGUAUUUGAU	1	1	Su(Ste)		Su(Ste)
4952	UAUUGUAAGCCGAAGUAGUCAGU	1	0	transposon	LTR	diver2
4953	UCCUCAACUGUGAGAUCCUCAA	1	no match			
4954	UCUCAUCAUCGUAGAACAAGCCCGUA	1	2	Su(Ste)		Su(Ste)
4955	UCUCAUCGUCGAGAACAGCCCGA	1	no match			
4956	UCCGGGGUCGAAAUGACCUUGACCU	1	no match			
4957	UGAUGACCGAAAUUUGGAAAACCAGAU	1	0	repeat		SAR_DM
4958	UUGUUUCAGAGCGUUGGCCAGUU	1	0	transposon	Low_complexity	F-element
4959	GAACUUGAGACCCCAACCGCUCC	1	1	no annotation		
4960	GGAUAACUGUGAGGCCUCAA	1	2	no annotation		
4961	ACUCUCAUCGUCGUAGAACAAGCCCGA	1	2	Su(Ste)		Su(Ste)
4962	AUCGGUCUCGGGAAUGAAAUAAAUGCCAGAAUUUUAUCGGGGUUC	1	no match			
4963	UAGGAGACGUGGACACGGAAGAGCUG	1	1	transposon	LINE	R1-element
4964	UGAUUCCGAUCUUGGACAUUCCCG	1	0	transposon	LTR	roo
4965	GUAUGACCCUUACUGAAAUGCAG	1	no match			
4966	GGAGUCCUCAACUGUGAGGGGAUCCUCAA	1	no match			
4967	CUCAUCUUCUUCUCGACGUUCGACCAG	1	no match			
4968	UGACCCAAUUGCGCGCACAUAAGCUGC	1	0	transposon	LTR	flea
4969	UAGAUCUUCGCCGUUGACUUGGCCC	1	1	Su(Ste)		Su(Ste)
4970	CAUUCGAUCUCCGACUAGGCCCGCG	1	0	transposon	LTR	GATE
4971	GAGUCUCACUAGUGAGGAGAGUCCUCGAA	1	no match			
4972	AUCGUCUGGGAUGAAAUCGUAAACCAAAGGCCGUGAGUUCA	1	no match			
4973	UCUCAUCGUCGUAGAACAAGCCCGAGU	1	1	Su(Ste)		Su(Ste)
4974	UACCCCAGAUUGUUGACUCUUUGCC	1	2	no annotation		
4975	CUGUGAGGAGAUCCUCAA	1	1	coding gene		CG13800-RA|CG13800
4976	GCCGACGAGAUGCCCAGGGCGUCCA	1	1	transposon	LTR	ninja-Dsim-like
4977	ACUUCGGGAUAAGGAUUGGCUCC	1	1	ncRNA		rRNA
4978	UAGAUCUUCCCCGUUGACUUGCCC	1	1	Su(Ste)		Su(Ste)

4979	UGUUUCAUCGUUAGACGGCUCGGGGCG	1	2	AT-chX		AT-chX-B
4980	AUCGUCUCGGGGAUGAAAAUCUCAUCGUCGUAGAACAAGCCCG	1	no match			
4981	AUCCUCAACGUGAGGGAUCCUCAA	1	no match			
4982	UGUUUCAUCGUUAGACGGCUCGCGC	1	1	AT-chX		AT-chX-B
4983	CUCAACUGUGAGGGGGAUCUUCAA	1	no match			
4984	AUCGUCUCGGGGAUGAAAUCCUCGGGAUCAAGGACAACGAGUU	1	no match			
4985	UGUUUCAUCGUUAGACGGCUCGGGAAA	1	no match			
4986	UCUCAUCGUCGUAGAACAAGCCCAU	1	2	Su(Ste)		Su(Ste)
4987	UCGGCUUUCUGACGAGAACUCGACG	1	2	no annotation		
4988	UAGAUCUUCACGUUGACUUGGCCC	1	no match			
4989	UGCUCUCGAAUGUAUCCCCGAUA	1	1	transposon	LINE	R1-element
4990	AGGUGAGGGUGCGAAUCCUCAA	1	no match			
4991	UACAAUACCCAGUUACUCGUGCCAA	1	1	no annotation		
4992	AGAAAAAAAGGUGGAAAUGAAGAAAGUACA	1	no match			
4993	UCUCAUCGUCGUAGAACAAGCCCGACUGUAACUCGGGGUCAAUAU	1	no match			
4994	AUCGUCAGUCGUACGGUAGAACAAGCCCGGACUG	1	no match			
4995	AUAGAAGAACAUUGAAGCGCUUGAC	1	1	Su(Ste)		Su(Ste)
4996	UCGGCUCGGAGAAACGCUGUAUGCGC	1	0	transposon	LTR	flea
4997	UAGAUUCGGCUAGUUACUCUUCUA	1	0	transposon	LINE	Circe
4998	UUAAAUACCUCGAGUACGUCUGCUC	1	1	transposon	LTR	Rt1c
4999	AAAACCUAGAAGAUGAUCCUGAGU	1	no match			
5000	GCAACUCUACGGCCCGGACUGCUC	1	0	transposon	LTR	ninja-Dsim-like
5001	AUCGUCUCCGGGAUGAAAUUCACGCCACGCUCCGACCUGAUG	1	no match			
5002	UUAUACCCAGGUGAUCACCACUAU	1	1	Su(Ste)		Su(Ste)
5003	UGUUUCGUCUUUAGACGGCUCGGGC	1	2	AT-chX		AT-chX-B
5004	UCUCAUCGUCGUAGAACAAGCUCGAGG	1	1	Su(Ste)		Su(Ste)
5005	UAAUGUCCCUUAAAUCAGCACUCA	1	1	no annotation		
5006	UCUCAUCGUCGUAGAACAAGCCCGUCUGUAAUCGGGUCAAU	1	no match			
5007	CAAAGCCUUUCAGACGUUUGACUC	1	0	transposon	DNA	transib4
5008	AUGUUUCAUCGUUAGACGGACUACGGGC	1	no match			
5009	CGUUGACAAUUCCGAACAUACGC	1	no match			
5010	UGGGUCUUCACCGUUGACUUGCCC	1	2	Su(Ste)		Su(Ste)
5011	AUCAAGGUGGAAAUGAAGAAAGUACA	1	no match			
5012	UCUCAUCGUCGUAGAACAAAGCCCGAGG	1	no match			
5013	GAGUGUGUCUCGGAUCUACCCAG	1	0	transposon	LTR	GATE
5014	UCUCAUCUUCGUAGAACAUGCCCGA	1	2	Su(Ste)		Su(Ste)
5015	UCGCAUCGUCGUAGAACAAGCCCGUA	1	2	Su(Ste)		Su(Ste)
5016	AGAAAAGUCUCAUCGUCGUAGAACAGUCCCG	1	no match			
5017	AUCGUCUCGGGGAUGAAAAAAAGGUGGAAAAUGAAGAAAAGUACA	1	no match			
5018	CGUCAACUGUGAGGGGAUCCUCAA	1	no match			
5019	CUCUCUAUCGUCGUAGAACAACGCCCGA	1	no match			
5020	UUCUCAUCGUCGUAGAACAAGCCCGUA	1	1	Su(Ste)		Su(Ste)
5021	ACCGGACGGGCCUAGAGUUAUUCCCCC	1	0	coding gene		CG40635-RA|CG40635
5022	UCGUCAUCGUACGUAGAACAACGCCCGGA	1	no match			
5023	ACCGAAAAGUAUGACCAGCUUGGACC	1	no match			
5024	UAGAUCGUUCACCGUUGACUAGCCCCA	1	no match			
5025	CUCUCAUCUCGUAGAACAACGCCCGUA	1	no match			
5026	AUUGACCCGAGGUUACAGGACCCGAGCCGUCUAACGAUGAAACAU	1	no match			
5027	AAACAUAUGAUUCCUGUGACCCCGA	1	0	transposon	LTR	springer
5028	UCCAAGCGAACUGUGCCCGAGGCUAUA	1	no match			
5029	UCUCUCGUCGUAGAACAAGCCCGU	1	no match			
5030	ACAAGAUAUGACCAAAAGACAAAGACCA	1	no match			
5031	AUCGUCUCGGGAGUGAAAUACUCAUCGUCGUAGGAACAACGCCCG	1	no match			
5032	UGGACGCCGUUUAUGAACGAAAAAGUU	1	0	transposon	LINE	Cr1a
5033	UUGGUAUGGGUUACGAGAUAUCUGA	1	0	transposon	LTR	flea
5034	UGUUUCAUCGUUAGACGGCUCGGGCUCUGUAACGUCGGGUCAAU	1	no match			
5035	AUCUGAGGAACCAAGUGAUCCAGCUG	1	no match			
5036	UCUCAUCGUCUUAGAACAAGCCCCGA	1	no match			
5037	UCUCAUAGUCGUAGAACAAGCCCG	1	1	Su(Ste)		Su(Ste)
5038	UCCGAGGAACCAAGUGAUCCAGCUG	1	1	Su(Ste)		Su(Ste)
5039	AGUUCAAGCCCGGGUAUUCUGAAGC	1	0	repeat		trf
5040	UAUAAUGCCCCGAAGGUUGCCCGA	1	2	transposon	LINE	R1-element
5041	UUCCGAAGAGGAUGGAUACCGCGGC	1	0	transposon	LTR	GATE
5042	AUAAUGCCCCGAAGGUUGCCCUCU	1	0	transposon	LINE	R1-element
5043	UCUCAUCGUCGUAGAAGAGCCCGA	1	no match			
5044	UAGUACAUUAAUUACUCGAUGACGC	1	0	transposon	LTR	invader4
5045	AUAGGAUCUUCACCGUUGACUAGCCCGA	1	no match			
5046	AUCGUCUCGGGAGUGAAAGGAUCCUCAA	1	no match			
5047	GAAAACCUGAAGAUGAUCCUGAGU	1	0	transposon	LINE	Helena
5048	AUCGUCUCGGGAGUGAAAUACCGAUCUCCUGAGUCGGACUCCGAG	1	no match			
5049	GCACACAGAGGAGAGCGGAUGAGAGCA	1	no match			
5050	UCUCAUCGUCGUAGAACAAGCCCGACUGUACUCGGGUCAAU	1	no match			
5051	UCUCAUCUCGUAGAACAAGCCCGACUGUACUCGGGUCAAU	1	no match			
5052	AUCGUCUCCGGGAUGAAAUGUUUCAUCGUUAGACGGCUCGGA	1	no match			
5053	CCGUGGAGACGGCUUUAGGACCUGGU	1	0	transposon	LTR	ninja-Dsim-like
5054	AGAUAAGGGAAGUCGGCAAAAAUAGAU	1	no match			
5055	AUCUCAACGUCGUAGAACAAGCCCGA	1	2	Su(Ste)		Su(Ste)
5056	AGCCCCGUUACUCUGUGGUGACCACGA	1	no match			
5057	GUCUCGGGAGAAAGGAUCCUCAA	1	no match			
5058	UCAAUGGCGUCUCCUCG	1	1	no annotation		
5059	UGUUCAUCGUACGUACGGCUCGGC	1	no match			
5060	GUUCAACCGGACGGGCCUAGAGU	1	0	coding gene		CG40635-RA|CG40635
5061	UGCAUGACCCAACAUCCUCCCCCCA	1	1	transposon	LTR	rooA

5062	UGCUUGAUAUGACCAGCUUGGACCA	1	1	Su(Ste)		Su(Ste)
5063	AAAAAACUGUGAGGGGAUCCUCAA	1	no match			
5064	CAGCACACAGAGGAGAGCAGAUGGAGAGCA	1	no match			
5065	CAGCAACAGAGGAGAGCAGAUGGAGCA	1	no match			
5066	GACUAGCUGGCCGAGGAUCCGAUUUUU	1	1	Su(Ste)		Su(Ste)
5067	UGUUUCAUCGUUAGACGGCUCGGCCG	1	1	AT-chX		AT-chX-B
5068	CGAAAGCACACCAGAGGAGAGCAGAAUGUAGAGCCA	1	no match			
5069	AGAAAUGUUUCAUCGUUAGACGGCUCCGGGC	1	no match			
5070	GAAUGUUUCAGUCGUUAGACGGCUCGGGUCCUGUAACCUACGGGU	1	no match			
5071	UCUUUAGGUCCGGGUUCGUCAUCAG	1	0	transposon	LINE	G6
5072	GUCCCAACUGUGAGGGGAUCCUCAA	1	no match			
5073	GGGGAUAUAGCUCAGUGGUAGAGCA	1	0	ncRNA		tRNA
5074	GAGUCGUCACCGCGCGCAAUCCCGUG	1	no match			
5075	CGAAAAUGUUUCAUCGUUAGACGGCCUCGGGC	1	no match			
5076	UCCAAACGUACCACGUGCCCCAGU	1	no match			
5077	UACGUGGACGAGUUGCUCGAUUUG	1	0	transposon	Low_complexity	F-element
5078	CUCAUCGUCGUAGAACAAGCCCCGA	1	2	Su(Ste)		Su(Ste)
5079	GGAUCCUCAACUGUGAGGGAUCUCAA	1	no match			
5080	UGUUUCAUCGUUAGACGGCUCGGGCCUGUAACUC	1	no match			
5081	AAAUGUUUCAUCGUUAGACGGCUCGGG	1	no match			
5082	AUCGUCUCGGGACUGAAAUCAACUGUGAGGGGACUCCUCAA	1	no match			
5083	GAUAAAAACCGAAUUACACGUACAAG	1	1	repeat		trf
5084	UAAGAACCGUGAGAUAUCGCUAAUC	1	0	no annotation		
5085	UGAUACCGAGAGUGACAAUGGUUGA	1	1	transposon	LTR	gypsy3
5086	UACAGUGCGAGGAGGCCCGUAUUAC	1	0	transposon	LTR	ninja-Dsim-like
5087	UGAGCUGGGGUUAGACCGUCGUGAGAC	1	1	ncRNA		rRNA
5088	UAGGCUGAGCGGAGAACCAGAGU	1	0	transposon	Low_complexity	F-element
5089	AUCCUCAACUGUGAGGGACCUCAA	1	no match			
5090	UAUGCAAUAUUUGUUGCGCCCCGGGA	1	0	transposon	LTR	opus
5091	AAAAUGUUUCAUCGUUAGACGGCUCGGG	1	no match			
5092	UGUUUCAUCGUUAGACGGCUCGAG	1	1	AT-chX		AT-chX-B
5093	UCUCAUCGUCGUAGAACGACGCCCGA	1	no match			
5094	CAGCACACAGAGGAGAGGAGAUGAGAGCA	1	no match			
5095	UUCCGAGGAACCAAUUUAUCCAGCU	1	1	Su(Ste)		Su(Ste)
5096	UAAAUAGACUAAUCGGACAGGUGACUCGG	1	no match			
5097	GUCUCUAUCGUCGUAGAACAAGCCCGUA	1	no match			
5098	UCUUCUUCUCGACGUUGACCAGCCU	1	0	transposon	LTR	ninja-Dsim-like
5099	UAACAAGAUGCAUACGAUGUGCUC	1	no match			
5100	UCUCAUCGUCGUAGAACAAGCCCGAGCUG	1	no match			
5101	GGCUCCGGGGGAAGUAUGGUUGCAAAG	1	0	ncRNA		rRNA
5102	UAAACAAUUGGAAGAUAAGACCCA	1	1	transposon	LINE	Cr1a
5103	AUCUCAUCAUCGUAGAACAAGCCCCGA	1	no match			
5104	UGUCUUCAUUUAAGCUCUGACUCGAA	1	0	repeat		trf
5105	UCUCAUCGUCGUAGACAAGGCCCGACUGUAACGUCGGGUCAAU	1	no match			
5106	AGGAGUCCUCAACUGUGAGGGAUCCGUCAA	1	no match			
5107	AUAAAAGAAAUCCUGACAAGAUCCCGGGC	1	no match			
5108	AUUCGUGACGAGAAACGUCGAGCCGCGAAAAA	1	no match			
5109	UCUUCACCGUUGACUCGGCCCACACA	1	1	Su(Ste)		Su(Ste)
5110	AUCUCAACUGUGAGGGGGAUCCUCAA	1	no match			
5111	GUCUAACGCCGGAGACAGUUUGA	1	0	transposon	LTR	micropia
5112	UCUCAUCGUCGUAGAACAACGCCCUAA	1	no match			
5113	UUCUCAUCUCGUAGAACAAGCCCGA	1	no match			
5114	UCAGCAAGCACCCCAGACUGCACCCGG	1	0	coding gene		CG1472-RA|sec24
5115	GAAACCCGAUGGAACGAAAUGUCAUU	1	0	coding gene		CG3354-RA|Mst77F
5116	UGUAACGUCGGGUCAAU	1	2	no annotation		
5117	GCUUGGGAACACCCCGUGUUGUUGGCC	1	1	ncRNA		5SrRNA
5118	UCUCAUCGUCGUAGAACAAGCCGAG	1	no match			
5119	CGAAAUCUCAUCGUCUAGAACAGCCCGACUGUAACUCGGGGUCAA	1	no match			
5120	UUGGAUGCCCAGAUACGAAAAACAUA	1	0	no annotation		
5121	UCUCAUCGUCCGUAGAACAAGCCCG	1	no match			
5122	UGAUACCGAGAAUGACACUGGUUGA	1	1	transposon	LTR	gypsy3
5123	GGCACACAGAGGAGAGCGAGAUGAGAGCAA	1	no match			
5124	GCGAGCAGGACUGGGCGGCGG	1	no match			
5125	UGGUACCUUUCCCGAUGAGCCAGU	1	no match			
5126	CUCAACUGUGGGAUCCUCAA	1	no match			
5127	UCUGGUUGAACGUGUCCUGAAAUUAG	1	0	Su(Ste)		Su(Ste)
5128	UAGAUCUUCACCAUUGACUUGCCCC	1	1	Su(Ste)		Su(Ste)
5129	UGUUUCAUCGGUAGACGGCUCGGGC	1	1	AT-chX		AT-chX-B
5130	UCUCAUCGUCGUAGAAGAAUCCCGA	1	2	Su(Ste)		Su(Ste)
5131	AUCUCAUCGUGUAGAACAAGCCCGA	1	no match			
5132	CCUCAACUGUAGGAUCCUCAA	1	no match			
5133	UUCCGAGGAACCAAUUGAUUCAGCUG	1	0	Su(Ste)		Su(Ste)
5134	GGAACUGUGAGGGGAUCCUCAA	1	no match			
5135	GCCAGGCCCGUAUAACGUUAAUGAU	1	0	ncRNA		rRNA
5136	UUGACCUGCUCACAGUAAUUAAGAA	1	0	no annotation		
5137	CAGCACACAGAGGAGAGCGGAUGAGAGA	1	no match			
5138	UACUUGACCCCCGAUCUGUGGAAGGA	1	0	coding gene		CG5920-RA|sop
5139	AUCGUCUCGGGGAUGAAAAUGUUUUCAUCGUUAGACGGCUCGGGC	1	no match			
5140	AUCUCAUCGUCGUAGAACAAAGCCCCA	1	no match			
5141	AUAGAUCUUCACCGUUGACUUGGCCCC	1	2	Su(Ste)		Su(Ste)
5142	UGGACCUUUUUCCGAUGAGCCAGUG	1	1	transposon	LINE	G
5143	UGUUUCAUCGUUAGGACGGCUCGGGC	1	no match			
5144	UGUUUCAUCGAUAGACGGCUCGGGC	1	1	AT-chX		AT-chX-B

5145	UACAAUACCCAGUUACUCGUGCCA	1	0	no annotation		
5146	UUAAUGCCGGAAGCGAUCCCAGUU	1	2	AT-chX		AT-chX-B
5147	UAUUACAUAGAUGUAGAUGGGUCGGAG	1	0	no annotation		
5148	UCUCAUCGUCGUAAAGAAGAAGCCCG	1	no match			
5149	UAACCAAUAACGGCCAGAGAAAUUUC	1	1	ncRNA		alphagamma-element
5150	CCUCAACUGUGAGGGGAUCCUCUAA	1	no match			
5151	CUUGGACCGAACACUUUGAACCAAGU	1	0	Su(Ste)		Su(Ste)
5152	CUGUAACUCCGGGUC	1	1	no annotation		
5153	CAAGAUAUGACCAACGCAAAGACACU	1	1	transposon	LINE	Cr1a
5154	UUUCUACCAAAUAAAUAGGACCUGAAG	1	0	transposon	LTR	Quasimodo
5155	UGUGCCGGAGCGUGGGACCCCGAAGA	1	0	transposon	LTR	roo
5156	UUCCUCGGGAUCAAGGACAACGAGUUC	1	0	no annotation		
5157	GCCCCGUACUCUGUGGUGACCAG	1	1	transposon	LINE	baggins
5158	AUCUCAUCGUCGUAGAACAGGCCCG	1	2	Su(Ste)		Su(Ste)
5159	GACACAGAGGAGAGCAGAUGAGACA	1	no match			
5160	UCGUCAGUCGUCGUAGAACAAGCCCGG	1	no match			
5161	UCUCAUCGUCGUAGAACAACGCCCGACUGUAACGUCGGGUCAAU	1	no match			
5162	UUCAUCGUCGUAGAACUAGCCCGA	1	1	Su(Ste)		Su(Ste)
5163	CAAGAUAUGACCAAAGCAAAAACAC	1	0	transposon	DNA	1360
5164	CGGGCUCAACCUGGGAACUGCGUU	1	no match			
5165	AAGACACCGUUUCCCGUCCGAUCAA	1	no match			
5166	UUCACCGUUGACUAGCCCCA	1	0	Su(Ste)		Su(Ste)
5167	CUAGAUCUUCACCGUUGACUAGCCC	1	1	Su(Ste)		Su(Ste)
5168	CAAGAUAUGACCAAAGCAAAGAUAA	1	1	transposon	DNA	1360
5169	CCCAUCUCACCAGACUCUUUGGCC	1	0	transposon	LTR	diver2
5170	GCUUCUGAGCGCGUAAACCGAU	1	0	transposon	LTR	G4
5171	UAGUUUCGAUCGUUAGACGGCUCGGGC	1	no match			
5172	GUGCAUAUCAUUCACGCCACGCUCCGA	1	0	Su(Ste)		Su(Ste)
5173	GGUUCCAUGGUGUAAUGGUUAGCA	1	0	ncRNA		tRNA
5174	UUGAUGGUGGCAACCGACUCGACGA	1	0	transposon	LINE	Doc4
5175	GAUGUCCACCAAAGGCAUGCCCGAUG	1	0	transposon	LTR	opus
5176	UGUUUCAUCGUUAGACGGCUCAGG	1	1	AT-chX		AT-chX-B
5177	UGUUUCAUCGUUAGACGGCUCGGGUUU	1	no match			
5178	CGAAAUCUCAUCGUCGUAGAACAGCCCG	1	no match			
5179	CCUCAACUGUGAGGGGAGUCCUCAA	1	no match			
5180	AACCACUGUGAGGGGUAUCCUCAA	1	no match			
5181	AAACUUCGGGAUAAGGAUUGGCUCUGA	1	1	ncRNA		rRNA
5182	AUACUGAGUCGUUUACCGAGGGCC	1	0	transposon	LTR	G4
5183	UGAUACCGAGAAUGACAACGGUUGA	1	1	transposon	LTR	gypsy3
5184	UUUGUAGGUGUACUGACCCAGUCGG	1	0	transposon	LINE	baggins
5185	UAGAUCUUCCCGUUGACUUGGCC	1	no match			
5186	UGUUUCAUCGUUAGACGGCUCCGGGCG	1	no match			
5187	AUCGUCAUCGUACGUAGAACAAGCCCGG	1	no match			
5188	UCUCAUCGUCGUAGAACAAGCCCGACCUGUAACUACGGGUCGAAU	1	no match			
5189	CGAAAUCCGAACGUACCACGGUGCCCCAGU	1	no match			
5190	UGUUUCAUCGUUAGACGGCCCCGGC	1	2	AT-chX		AT-chX-B
5191	UAGAUCUUCACCGUGACUUGGCCCCG	1	no match			
5192	UGGUUUCAUCGUUAGACGCUCGGGCCG	1	no match			
5193	AGGCAACUGUGAGGGGGAUCCUCAA	1	no match			
5194	UAAUAUUGGACCCAGUACACUGCCU	1	1	transposon	LINE	Doc
5195	AGUCCUCAACUGUGAGGGAUCCUCAA	1	no match			
5196	UCUCAUCGUCGUAGAACAAGCCAG	1	1	Su(Ste)		Su(Ste)
5197	AGUUCAAGCCCGGGUAUUCUGAAG	1	0	repeat		trf
5198	UGUUUCAUCGUUAGACGACUCGGGUC	1	no match			
5199	UAGAUCUUCACCGUUGACUAGCCCCACUGUAACUCGGGGUCAAU	1	no match			
5200	CCAAGAUAUGACCAAAGCAAAGACACUA	1	1	transposon	LINE	Cr1a
5201	CCUCUUCGUCGUAGAACAAGCCCGA	1	2	Su(Ste)		Su(Ste)
5202	UCCUCAACUGUGAGGGGUCCUCAA	1	no match			
5203	UUAUAAGGAUGAUGAUACCGACUAAGG	1	1	no annotation		
5204	AUCGUCUCGGGACUGAAACCUCUCAUCCGUUCGUACGAACAACCG	1	no match			
5205	AAUCUCAUCGCCGUAGAACAAGCCCGA	1	no match			
5206	UUAAUUCCGAUAACGAACGAGAGACUC	1	no match			
5207	GACUAGCUGGCCGAGUAUUUGAUAA	1	1	Su(Ste)		Su(Ste)
5208	UCGUCAUCGUGUAGAACAAGCCCGGA	1	no match			
5209	AGUGUUUCAUCGUUAGACGGCUCGGGC	1	2	AT-chX		AT-chX-B
5210	UGUAUAGAUGAGGUAUAAGGUUGGCC	1	0	transposon	Low_complexity	F-element
5211	GCACACAGACGGAGAGCAGAUGAGACGCA	1	no match			
5212	UCCAAUUAAAAAAUGACCUGGUGU	1	0	transposon	LTR	roo
5213	GGGGCCCGAAAACACGAGAGCCGAGC	1	0	transposon	LTR	opus
5214	CCUCAUCGUCGUAGAACAAGCCCGACUGUACUAAACGGGUCAAU	1	no match			
5215	AGAAAGGUGGAAAUGAAGAAAGUACA	1	no match			
5216	UCUCCUGGAGACACCAAUGCUGGCC	1	1	coding gene		pseudogene
5217	UACUCAUCGUCGUGAACAAGCCCGA	1	no match			
5218	CAGCACACAGAGGAGAGCAGAUGAAAGCA	1	no match			
5219	UGUUCAUCGUUAGACGGCUCGGGU	1	no match			
5220	AUACUCAUUCGUCCGUUAGAACAAGCCCCGA	1	no match			
5221	UAAAGAAUCCUGACAAGAUCCCGGA	1	1	transposon	LTR	invader1
5222	UGUCCACCAAAGGCAUGCCCGAUGU	1	0	transposon	LTR	opus
5223	UUUGUGUACCCCAGAUUGUUGACUCU	1	1	no annotation		
5224	UCAAACUGAAGGAGUACGAAUGAUC	1	1	transposon	LINE	R1-element
5225	AUCGUCUCGGGAGUGAAAUAACAUACACGCCCAGGAGUCUGUUCC	1	no match			
5226	AGGCUCAACUGUGAGGGGAUCCUCAA	1	no match			
5227	UGGAAUUUUUCCGAUGAGCCAGU	1	no match			

5228	AGGGAGUUUGACUGGGGCGGUAC	1	0	ncRNA		rRNA
5229	CGCACACGAGAGGAGAGCAGAUGGAGAGCA	1	no match			
5230	AUAAGAUCUUCACCGUUGACUAGCCCCA	1	2	Su(Ste)		Su(Ste)
5231	CUCUCAUUCGUCGUACGAACAAGCCCGACUGUAACCUCGGGUCAA	1	no match			
5232	AUAAAUGCCAGAAUUUUUAUUGGGGGUUC	1	no match			
5233	AUGUUUUCAUCGUUAGACGGCUCGGGAAA	1	no match			
5234	UUCCGUACGUCCAGAUUGGCUUGAG	1	0	transposon	LINE	G2
5235	GCUUCACCUUGAAGACCUUAAGUCCCG	1	0	transposon	LINE	BS
5236	UCUCAUCGUCGUAGAACAAGCCA	1	1	Su(Ste)		Su(Ste)
5237	ACGCCACACUGCGUGUUAACGACCUGAC	1	0	coding gene		CG7008-RA|Tudor-SN
5238	UAGAUCUUCACCGUUGACUAGCCCCACUGUAACUCGGG	1	no match			
5239	AAAACCUAGUUACUUACGGGCUCAA	1	0	no annotation		
5240	GAACGUACCUUGAGCAUAUAUG	1	0	ncRNA		rRNA
5241	AAAAAACUUGAAGAUGAUCCUGAGUCC	1	no match			
5242	UCGACGGAGCUCACUUACCUAAUGU	1	0	transposon	LINE	R1-element
5243	UUCGAAUUGCCCUAACGGUGUAACA	1	0	transposon	LTR	flea
5244	UCAACAGUGAGGGGAUCCUCAA	1	no match			
5245	CGAAAGCACACCAGAGGAGAGCAGAUGUAGAGCCA	1	no match			
5246	UCCGAACGUACCACGGUGCCCCAGUA	1	1	transposon	Low_complexity	F-element
5247	CAAGGUAUGACCUAAGCAAAGACAC	1	2	transposon	LINE	Cr1a
5248	UGAUCUUCACCGUUGACUAGCCC	1	1	Su(Ste)		Su(Ste)
5249	UAAUGUUGGCCAAUUUCGAGUAUAUC	1	0	transposon	LINE	Cr1a
5250	UAGAUCCUCACCGUUGACUAGCCCC	1	1	Su(Ste)		Su(Ste)
5251	AUUUCUCAUCGUCGUAGAACAAGCCCCGA	1	no match			
5252	UAGGUUAUGGACCCAGUAAACAUU	1	1	ncRNA		alphagamma-element
5253	UGUUUCAUCGUCGUAGAAGAAGCCCGA	1	2	no annotation		
5254	UCUCACCUCGUAGAACAAGCCCG	1	2	no annotation		

Supplementary Table 2
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Su(Ste)-4 (common)

Alignment	Mismatches	AGO3	Aub
----UCUCAUCGUCGUAGAACAAGCCCGA----------	0	1670	43207
----UCUCAUCGUCGUAGAACAAGCCCG----------	0	151	6212
----UCUCAUCGUCGUAGAAGAAGCCCGA----------	1	10	1183
---UUCUCAUCGUCGUAGAACAAGCCCGA----------	0	24	1138
---AUCUCAUCGUCGUAGAACAAGCCCGA----------	1	3	768
----UCUCAUCGUCGUAGAACAAGCCCGAG----------	0	14	616
----UCUCAUCUUCGUAGAACAAGCCCGA----------	1	1	436
----UCUCAUCGUCGUAGAACAAGCCCGAGG----------	0	94	278
----UCUCAUCGUCGUAGAACAAGCCCAA----------	1	4	263
----CCUCAUCGUCGUAGAACAAGCCCGA----------	1	5	169
--UUUCUCAUCGUCGUAGAACAAGCCCGA----------	0	8	129
----UCUCAUCGUCGUAGAACAAGCCCGU----------	1	17	128
----UCUCAUCGUCGUAGAAGAAGCCCG----------	1	2	117
----UCUCAUCGUCUUAGAACAAGCCCGA----------	1	1	102
----UCUCAUCGUCGUAGAACGAGCCCGA----------	1	2	80
----UCUCAUCGUCGUAGAACAACCCCGA----------	1	1	68
----UCUCAUCGUCGUAGAGCAAGCCCGA----------	1	1	66
----UCUCAUCGUCGUAGAACAGGCCCGA----------	1	4	62
----UCUCAUCGUCGUAGAACAAGCCCGG----------	1	27	58
----UCUCAUCGUCGUAGGACAAGCCCGA----------	1	1	58
----UCUCAUCGUCGUAGAACAAGCCC----------	0	3	55
----UCUCAUCGUCGUAGAACAAGUCCGA----------	1	1	48
----UCUCAUCGUCGUAGAACAAGCCCUA----------	1	3	47
----UCUCGUCGUCGUAGAACAAGCCCGA----------	1	1	46
----UCUCAUCGUCGUAGAACAAGCCGA----------	2	2	44
----UCUCAUCGCCGUAGAACAAGCCCGA----------	1	2	35
-----CUCAUCGUCGUAGAACAAGCCCGA----------	0	1	35
---UACUCAUCGUCGUAGAACAAGCCCGA----------	1	1	32
----UCUCAUCGUCGUAGAACAAGCCCCG----------	2	1	32
----UCUCAUCGUCGUAGAACAAGCCUGA----------	1	13	31
----UCUCAUCGUCGUAGAAUAAGCCCGA----------	1	2	27
----UCUCAUCGUCGUGGAACAAGCCCGA----------	1	1	27
----UCUCACCGUCGUAGAACAAGCCCGA----------	1	1	26
----UCUCAUCGUCGUAGAACAAGCCCGAC----------	1	1	25
---GUCUCAUCGUCGUAGAACAAGCCCGA----------	1	1	21
----UCUCAUCGUCGUAGAACAAGCCCGAU----------	1	20	19
----UCUCAUCGUCGUAGAUCAAGCCCGA----------	1	1	18
----UCUCAUCGUCCUAGAACAAGCCCGA----------	1	1	16
----UGUCAUCGUCGUAGAACAAGCCCGA----------	1	1	16
----UCUCAUCGUCGUAGAACAAACCCGA----------	1	1	13
----UCUCAUAGUCGUAGAACAAGCCCGA----------	1	2	11
----UCACAUCGUCGUAGAACAAGCCCGA----------	1	2	11
----UCUCAUCGUCGUAGAACAAGCCG----------	1	1	11
----UCUCAUCGUCGUAGAACAAGCCCGC----------	1	6	10
----UCUCAUCGUCGUAGAACAAGCCCGGG----------	1	2	7
----UCCCAUCGUCGUAGAACAAGCCCGA----------	1	1	5
---CUCUCAUCGUCGUAGAACAAGCCCG----------	1	1	5
----UCUCAUCGUCGUAGAACAAGCCCGAGC----------	1	2	3
----UCUCAUCGUCGUAGAACAAGCCGGA----------	1	2	3
----UCUCAUCGUCGUAGAAGAAGCCCGAGG----------	1	1	3
----UCUCAUCGUAGUAGAACAAGCCCGA----------	1	6	2
----UCUCAUCGUCGUAGAACAAGCCCU----------	1	2	2
----UCUCAUCGUCGUAGAACAAGCCCGGGG----------	1	1	2
---AUCUCAUCGUCGUAGAACAAGCCCGAGG----------	1	1	1
	Total	**2,128**	**55,827**

Su(Ste)-4 (AGO3 unique)

Alignment	Mismatches	AGO3	Aub
----UCUCAUCGUCGUAGAACAAGCCCGAGGA---------	0	12	0
----------UAGAACAAGCCCGAGGAACUGUCGAA	0	3	0
----------CGUAGAACAAGCCCGAGGAACUGUCGC-	1	2	0
----UCUCAUCGUCGUAGAACGAGCCCGAGG----------	1	2	0
----UCUCAUCGUCGUAGAACAAGCCCGCGG----------	1	1	0
---AUCUCAUCGUCCUAGAACAAGCCCGA----------	2	1	0
----UCUCAUCGUCGUAGAACAAGCCCGAUU----------	2	1	0
----UCUCAUCGUCGUAGAACAAGCCCGGGGU----------	2	1	0
----UCUCAUCGUCGUAGAACAAGCCCGAUC----------	2	1	0
----UCUCAUCGUCGUAGAACAAGCCCGAUGA---------	1	1	0
	Total	**25**	**0**

Su(Ste)-4 (Aub unique)

Alignment	Mismatches	AGO3	Aub
---UUCUCAUCGUCGUAGAACAAGCCCG----------	0	0	204
---AUCUCAUCGUCGUAGAACAAGCCCG----------	1	0	118
----UCUCAUCGUCGUAGAACAAUCCCGA----------	1	0	77
---UCGUCAUCGUCGUAGAACAAGCCCGA----------	2	0	68
----UCUCAUCGUCGUAUAACAAGCCCGA----------	1	0	66
----UCUCAUCGUCGUAGAACAAGCCCUAG----------	1	0	59
----UCUCAUCCUCGUAGAACAAGCCCGA----------	1	0	56
----UCUCAUCGUCAUAGAACAAGCCCGA----------	1	0	53
----UCUCAUCGUCGUAGAACAAGCCCA----------	1	0	51
----UCUCAUCGUCGCAGAACAAGCCCGA----------	1	0	50
--UUUCUCAUCGUCGUAGAACAAGCCCG----------	0	0	49
----UCUCAUCUUCGUAGAACAAGCCCG----------	1	0	44
--AAUCUCAUCGUCGUAGAACAAGCCCGA----------	2	0	43
----UCUCAUCGUUGUAGAACAAGCCCGA----------	1	0	43
----UCUCAUCGUCGUAGAACAAGCCCC----------	1	0	42
----UCUCAUCGUCGUAGGACAAGCCCG----------	1	0	36
--AUUCUCAUCGUCGUAGAACAAGCCCG----------	1	0	33
----UCUCAUCGUCGUAGAACAUGCCCGA----------	1	0	33
----UCUCAUCGUCGUAGAAAAAGCCCG----------	1	0	31
---CUCUCAUCGUCGUAGAACAAGCCCGA----------	1	0	30
----UCUCAUCGUCGUAGAACAGGCCCG----------	1	0	28
------UCAUCGUCGUAGAACAAGCCCGA----------	0	0	27
----UCUCAUUGUCGUAGAACAAGCCCGA----------	1	0	24
----UCUCAUCGUCGUAGAACAAGCC----------	0	0	23
----UCUCAUCGUCGUAGAACAAGCUCG----------	1	0	22
----UCUCAUCGUCGUAGAACAAGCUCGA----------	1	0	21
----CCUCAUCGUCGUAGAACAAGCCCG----------	1	0	20
----UCUCAUCGUCGUAGAAGAAGCCCGAG----------	1	0	19
----UCUCAUCGUCGUAGAACAAGCCCCA----------	1	0	18
----UCUCAUCGUCGUAGAACAAGCCAGA----------	1	0	18
---AUCUCAUCGUCGUAGAAGAAGCCCGA----------	2	0	15
-----CUCAUCGUCGUAGAACAAGCCCGAGG----------	0	0	14
---UUCUCAUCGUCGUAGAAGAAGCCCGA----------	1	0	13
----UCUCAUCGUCGUUGAACAAGCCCGA----------	1	0	12
----UCUCAUCGUCGUAGAAGAAGCCCAA----------	2	0	12
----UCUUAUCGUCGUAGAACAAGCCCGA----------	1	0	12
----UCUCAUCGUCGUAGCACAAGCCCGA----------	1	0	12
----UCUCAUCGUCGUAGUACAAGCCCGA----------	1	0	11
--AUUCUCAUCGUCGUAGAACAAGCCCGA----------	1	0	11
----UCUCAUCGUCGUAGAACAAGC----------	0	0	11
----UCUCAUCGUCUUAGAACAAGCCCG----------	1	0	11
----UCUCAUCGUCGUAGAAUAAGCCCG----------	1	0	11
---UACUCAUCGUCGUAGAACAAGCCCG----------	1	0	10
----UCUCAUCGUCGAAGAACAAGCCCGA----------	1	0	9
----UCUCAUCGCCGUAGAACAAGCCCG----------	1	0	9
---UUCUCAUCGUCGUAGAAGAAGCCCG----------	1	0	9
----UCUCAUCGACGUAGAACAAGCCCGA----------	1	0	9
----UCUCAUCGUCGUAGAACCAGCCCGA----------	1	0	9
---UCGUCAUCGUCGUAGAACAAGCCCG----------	2	0	9
---UUCUCAUCGUCGUAGAACAAGCCCAA----------	1	0	9
---AUCUCAUCUUCGUAGAACAAGCCCGA----------	2	0	9
------UCAUCGUCGUAGAACAAGCCCG----------	0	0	8
----UCUCUUCGUCGUAGAACAAGCCCGA----------	1	0	8
----ACUCAUCGUCGUAGAACAAGCCCGA----------	1	0	8
----UCUCAUCGUCGAAGUACAAGCCCGA----------	2	0	8
----UCUCAUCGUCGUAGAACGAGCCCG----------	1	0	7
---UUCUAAUCGUCGUAGAACAAGCCCGA----------	1	0	7
----UCUCAUCGUCGUAGAACAAUCCCG----------	1	0	7
----UCUCAUCGUCGUAGAGCAAGCCCG----------	1	0	7
----UCUCAUCGUCGUAGAACAAGCCCUAGG----------	1	0	7
--GUUCUCAUCGUCGUAGAACAAGCCCGA----------	1	0	7
----UCUCAUCUUCGUAGAACAAGCCCAA----------	2	0	7
-----CUCAUCGUCGUAGAACAAGCCCG----------	0	0	6
----UCUCAUCGUCGUAGAACAAGCCCGACG----------	1	0	6
---UCCUCAUCGUCGUAGAACAAGCCCGA----------	1	0	6
----UCUAAUCGUCGUAGAACAAGCCCGA----------	1	0	6
--UUUCUCAUCGUCGUAGAAGAAGCCCGA----------	1	0	6
--UUUCUCAUCUUCGUAGAACAAGCCCG----------	1	0	6
----UCUCAUCGUCGUUGAACAAGCCCG----------	1	0	6
----UCUCAACGUCGUAGAACAAGCCCGA----------	1	0	6
----UCUCACCGUCGUAGAACAAGCCCG----------	1	0	6
----UCUCAUCGUCGUAGAACAAGUCCGAG----------	1	0	6
----UCUCAUCGUCGUAGAUCAAGCCCG----------	1	0	6
-UUUUCUCAUCGUCGUAGAACAAGCCCGA----------	0	0	5
----UCUCAUCGUCGUAUAACAAGCCCG----------	1	0	5
----UCUCAUCGUCGUGGAACAAGCCCG----------	1	0	5
----UCUCAUCAUCGUAGAACAAGCCCGAG----------	1	0	5
----UCUCAUCGUCGCAGAACAAGCCCG----------	1	0	5
----UCUCAUCGUCGUAGACCAAGCCCGA----------	1	0	5
------UCAUCGUCGUAUAACAAGCCCGA----------	1	0	5
----UCUCAUCGUCGUGGAAGAAGCCCG----------	2	0	5
----UCUCAUCUUCGUAGAAGAAGCCCGA----------	2	0	5
----UCUCAUCGUCGUAGAACAACCCCG----------	1	0	5

---GUCUCAUCGUCGUAGAACAAGCCCG----------	1	0	5
----UCUCAUCGUCGUAAAACAAGCCCG----------	1	0	5
---UUCUCAUCGUCGUAGAACAAGCCC----------	0	0	5
----CCUCAUCGUCGUAGAACAAGCCCGAG----------	1	0	4
----UCUCAUCUUCGUAGAACAAGCCCGAG----------	1	0	4
----UCUCAUCGUCGUAGAACAAGCCUGAG----------	1	0	4
----UCUCAUCGUUGUAGAACAAGCCCG----------	1	0	4
----UCUCGUCGUCGUAGAACAAGCCCG----------	1	0	4
----UCUCAUUGUCGUAGAACAAGCCCG----------	1	0	4
---UUCUCAUCGUCGUAGAACAAGCCCGU----------	1	0	4
--AAUCUCAUCGUCGUAGAACAAGCCCG----------	2	0	4
---AUCUCGUCGUCGUAGAACAAGCCCGA----------	2	0	4
----UCUCAUCGUCGUAGAACAAGCCUG----------	1	0	4
---AUCUCAUCGUCGUAGAACAAGCCCGU----------	2	0	4
---UUCUCGUCGUCGUAGAACAAGCCCGA----------	1	0	4
----UCUCAUCGUCGUAGAACAAGCCCGACUA---------	2	0	4
----UCUCAUCGUCGUAGAACAAGUCCG----------	1	0	4
---UUCUCAUCGUCGUAGAACAGGCCCG----------	1	0	4
----UCUCAUCGUCGUAGUACAAGCCCGC----------	2	0	4
----UCUCAUCUUCAUAGAACAAGCCCG----------	2	0	3
--AGUCUCAUCGUCGUAGAACAAGCCCGA----------	2	0	3
----UCUCAUCGUUGUAGAACAAGCCCGAGG----------	1	0	3
--GGUCUCAUCGUCGUAGAACAAGCCCGA----------	2	0	3
----UCUCAUAGUCGUAGAACAAGCCCGAGG----------	1	0	3
--UUUCUCAUCGUCGUAGAACAAACCCGA----------	1	0	3
----UCUCAUCGUCGUAGAACAAGCCCGUG----------	1	0	3
----UCUCAUCGUCGUAGAACAAGCCAGU----------	2	0	3
----UCUCAUCGUCGCAGAACAAGCCCGC----------	2	0	3
---UUCUCAUCGUCGUAGAACAAGCCCGG----------	1	0	3
----UCUCAUCGUCGUAGAACAAGCUCGAG----------	1	0	3
----UCUCAUCGUCGUAGAACAAGCACGA----------	1	0	3
---ACCUCAUCGUCGUAGAACAAGCCCGA----------	2	0	3
--AUACUCAUCGUCGUAGAACAAGCCCGA----------	2	0	3
----UCUCAUCCUCGUAGAACAAGCCCG----------	1	0	3
--CUUCUCAUCGUCGUAGAACAAGCCCGA----------	1	0	3
---AUCUCAUCGUCGUAGAACAAGCCCAA----------	2	0	3
---AUCUCAUCGUCGUAGAACAAGCCCGAG----------	1	0	3
----UCUCAUCGACGUAGAACAAGCCCG----------	1	0	3
----UCUCAUCGUCGUAUAACAAGCCCAA----------	2	0	3
---UUCUCAUCGUCGUAGAACAAGCCCC----------	1	0	3
----UCUCAUCGUCGUAAAACAAGCCCGA----------	1	0	3
---UUCUCAUCGUCGUAGAACAAGCCCGAG----------	0	0	3
----UCUCAUCGUCGUAGAACAAGCCCCC----------	2	0	3
---UUCUCAUCGUCGUAGAACAAGCCAGA----------	1	0	3
----UCUCAUCGUCGUACAACAAGCCCG----------	1	0	3
---UUCUCAUCGUCGUAGAACGAGCCCGA----------	1	0	3
---UUCUCAUCGACGUAGAACAAGCCCGA----------	1	0	2
---UUCUCAUCGUCGUAGAACAAGCCCA----------	1	0	2
---AUCUCAUCGUCGUAGAACAAGCCUGA----------	2	0	2
-----CUCAUCGUCGUAGAAGAAGCCCGAG----------	1	0	2
----UCUCAUCGUCCUAGAACAAGCCCG----------	1	0	2
---AUCUCAUCGUCGUAGAACAACCCCGA----------	2	0	2
-AUUUCUCAUCGUCGUAGAACAAGCCCGA----------	1	0	2
--UUCGUCAUCGUCGUAGAACAAGCCCGA----------	2	0	2
---AUCUCAUCGUUGUAGAACAAGCCCGA----------	2	0	2
---UUCUCAUCGUCAUAGAACAAGCCCGA----------	1	0	2
----UCUCAUCGUCAUAGAACAAGCCCG----------	1	0	2
----UCUCAUCGUCGCUGAACAAGCCCGA----------	2	0	2
----UCUCAUCGUUUUAGAACAAGCCCGA----------	2	0	2
---AUCUCAUCGUCGUAGAACGAGCCCGA----------	2	0	2
---UUCUCAUCCUCGUAGAACAAGCCCGA----------	1	0	2
--UUCGUCAUCGUCGUAGAACAAGCCCG----------	2	0	2
----UCUCAUCGUCGUAGAAGAAGCCCGU----------	2	0	2
---AUCUCAUUGUCGUAGAACAAGCCCGA----------	2	0	2
---AUCUCAUCGUCGUAGAAUAAGCCCGA----------	2	0	2
-AAUUCUCAUCGUCGUAGAACAAGCCCG----------	2	0	2
---AUCUCAUCGUCGUAGAACAAGCCCUA----------	2	0	2
----UCUCAUCGUCGUCGAACAAGCCCGA----------	1	0	2
----UCUCAUCGUCGUAGUAGAAGCCCG----------	2	0	2
----UCUCAUCGUCGUAGGACAAGCCCUA----------	2	0	2
------UCAUCGUCGUAGAACAAGCCCAA----------	1	0	2
---UUCCCAUCGUCGUAGAACAAGCCCG----------	1	0	2
----UCUCAUCUUUGUAGAACAAGCCCGA----------	2	0	2
---AUCUCAUCGUCGUAGAACAAUCCCGA----------	2	0	2
-----CUCAUCGUCGUAGAAGAAGCCCGA----------	1	0	2
----UCUCAUCGUCGUAGAACAAGCCCAG----------	2	0	2
----UGUCAUCGUCGUAGAACAAGCCCG----------	1	0	2
---AUCUCACCGUCGUAGAACAAGCCCGA----------	2	0	2
-----CUCAUCGUCGUAGAACAAGCCCGAG----------	0	0	2
----UCUAAUCGUCUUAGAACAAGCCCGA----------	2	0	2
----UCUCAUCGUCGUGGAAGAAGCCCGA----------	2	0	2
----UCUUAUCGUCGUAGAACAAGCCCGAG----------	1	0	2
----UCUCAUCUUCGUAGAACAAGCCCGU----------	2	0	2
---UUCUCAUCGUCGUAGAACAAUCCCGA----------	1	0	2
----UCUCAUCGUCGUAGAACAAGCCCUC----------	2	0	2

---UUCUCAUCGUCGUAGAACAACCCCGA----------	1	0	2
----UCUCAUCGUCGUACAACAAGCCCGA----------	1	0	2
----UCUCAUCGUCGUAGAACAAACCCG----------	1	0	2
--UUUCUCACCGUCGUAGAACAAGCCCGA----------	1	0	2
---UUCUCAUUGUCGUAGAACAAGCCCGA----------	1	0	2
---UUCUCAUCGUCGUAGAACAAGUCCGA----------	1	0	2
----UCUCAUCUCCGUAGAACAAGCCCGA----------	2	0	2
----UCUCAUCGUCGUAGAACAAGCGCGA----------	1	0	1
--AUCCUCAUCGUCGUAGAACAAGCCCGA----------	2	0	1
----UCUCAUCGUCGUAGAAGAAACCCGA----------	2	0	1
---AUCUCAUCGUCCUAGAACAAGCCCG----------	2	0	1
----UCUUAUCGUCGUAGAACAAGCCCG----------	1	0	1
----UCUCAUCGUCGUAGAGGAAGCCCGA----------	2	0	1
---UACUCAUCUUCGUAGAACAAGCCCGA----------	2	0	1
----UCUCAUCGUCGUAGAGCAAGCCCAA----------	2	0	1
----UCUCAUCGUCGUAGAACAAGCCAGAG----------	1	0	1
----UCUCAUCGUCGUAGAACAAGCGA----------	2	0	1
---AUCUCAUCGUCGUAGAACAAGCCCUAG----------	2	0	1
--UUUCACAUCGUCGUAGAACAAGCCCGA----------	1	0	1
---UUCUCAUCGUCGUAGAACAAGCCCGCG----------	1	0	1
----UCUCAUCGUCGUAGAACAAGUCCGAGG----------	1	0	1
---UUCUCAUCGUCGUAGAGCAAGCCCGA----------	1	0	1
---AUCUCAUCGUCGUAGAACAAGCCCGC----------	2	0	1
---AUCUUAUCGUCGUAGAACAAGCCCGA----------	2	0	1
----UCUCAUCGUCGUAAAAGAAGCCCGA----------	2	0	1
----UCUCAUCGUCGUAGAACAUCCCCGA----------	2	0	1
----UCUCAUCGUCGUAGAACAAGCCCAU----------	2	0	1
----UCUCAUCUUCGUAGAACAAGCCCGAGG----------	1	0	1
----UCUCAUCGUCGUAUAACGAGCCCGA----------	2	0	1
----UCUCAUCUUCGUAGAACAGGCCCGA----------	2	0	1
----UCUCAUCGUCGUAUAACAAGCCCGAGG----------	1	0	1
---CUCUCAUCGUCGUAGAAGAAGCCCGA----------	2	0	1
----UCUCAUCGUCGUAGAACACGCC----------	1	0	1
----UCUCAUCGUCGUAGACCAAGCCCA----------	2	0	1
----UCUCAUCGUCGUAGAACAAGCCA----------	1	0	1
---UUCUCAUCGUCGUAGAACAAGCCCCA----------	1	0	1
----UCUCAUCGUCGUAGAACAAGCCCGACU----------	2	0	1
---AUCUCAUCGUCGUAGAACAAGUCCGA----------	2	0	1
---UUCUCAUCUUCGUAGAACAAGCCCGA----------	1	0	1
---AUCUCAUCGUCGUAGAACAAGCUCG----------	2	0	1
----UCUCACCGUCGUAGAACAAGCCCGAG----------	1	0	1
----UCUCAUCGUCGUAGAACAAGCCCGCU----------	2	0	1
----UCUCCUCGUCGUAGAACAAGCCCG----------	1	0	1
----UCUCAUAGUCGUAGAACAAGCCCG----------	1	0	1
----UCUCAUCGUCUUAGAACAAGCCCGC----------	2	0	1
---UUCUCAUCGUCGUAGAACAAGCCG----------	1	0	1
---UACUCAUCGUCGUAGAACAAGCCCGAG----------	1	0	1
----ACUCAUCGUCGUAGAACAAUCCCGA----------	2	0	1
----UCUCAUCGUCGUAGAACAGGCCCUA----------	2	0	1
----UCUCAUCGGCGUAGAACAAGCCCAA----------	2	0	1
----UCUCAUCGUCGUAGCACAAGCCCG----------	1	0	1
----UCUCAUCGUCGUAGAACAAGCCAA----------	2	0	1
----UCUCAUCGUCGUAGAACAAGCCCGGU----------	2	0	1
--AUUCUCAUCGUCGUAGAACAAGCCC----------	1	0	1
----CCUCAUCGUCGUCGAACAAGCCCGA----------	2	0	1
----UCUCAUCGUCGUAGAACAACGC----------	2	0	1
----UCUCAUCUUCGUAGAACAAGCC----------	1	0	1
----UCUUAUCGUCGUAGAACAAGCCCGAC----------	2	0	1
----GCUCAUCGUCGUAGAACAAGCCCG----------	1	0	1
---AUCUCAUCGUCGUAGAACAAGCCCU----------	2	0	1
----UCUCAUCGUCGUAGAACCAGCCCG----------	1	0	1
----UCACAUCGUCGUAGAACAAGCUCG----------	2	0	1
----UCUCAUCCUCGUAGAACAAGCCCAA----------	2	0	1
----UCUCAUCGUCGUAGAACAGCCCCGA----------	2	0	1
----ACUCAUCGUCGUAGAACAAGCCCA----------	2	0	1
---AUCUCAUCGUCGUAGAACAGGCCCG----------	2	0	1
----UCUCAUCGUCGUAGAACAAGCACGAG----------	1	0	1
---CUCUCAUCGUCGUAGAACAAGCCG----------	2	0	1
----CCUCAUCGUCGCAGAACAAGCCCGA----------	2	0	1
----UCAAAUCGUCGUAGAACAAGCCCGA----------	2	0	1
---UUCCCAUCGUCGUAGAACAAUCCCGA----------	2	0	1
---UUCUCAUCGUCGUAGAACAAGCCCCG----------	2	0	1
----UCUCAUCGUCGUAGAACAAGCGAG----------	2	0	1
----UCUCAUCGUCGUAGAAGAAUCCCGA----------	2	0	1
--ACUCUCAUCGUCGUAGAACAAGCCCGA----------	2	0	1
---UACUCAUCGUCGUAGAACAAGCCCGAGG----------	1	0	1
----UCUCAUCUUCGUAGAACAUGCCCGA----------	2	0	1
---UUCUCAUCGUCGUAGAACAAGCCUGA----------	1	0	1
----UCUCAUCGUCGUAGAAGCAGCCCG----------	2	0	1
----UCUCAUCGUCGUAGAACAAGACCGA----------	1	0	1
---UUCUCAUCGUCGUAGAACAAUCCCG----------	1	0	1
----UCUCCUCGUCGUAGAACAAGCCCGA----------	1	0	1
----CCUCAUCGUCGUAGAACAAGCCCGAGG----------	1	0	1
----UCUCAUCGUUGUAGAACAAGCCCUAG----------	2	0	1
---UUCUCAUCGUCGUAGAACAAGCCCGC----------	1	0	1

---UUCUCAUCGUCGUAUAACAAGCCCGA----------	1	0	1
----CCUCAUCGUCGUAGAAGAAGCCCGA----------	2	0	1
--UUUCUCAUCCUCGUAGAACAAGCCCG----------	1	0	1
----UCUCAUCGUCUUAGAACAAGCCCGAG----------	1	0	1
----UCUCAUCGUCGUAGAACAAGCCGG----------	1	0	1
----UCUCAUCGUCGUAGAACAAGCCAG----------	1	0	1
--UGUCUCAUCGUCGUAGAACAAGCCCGA----------	1	0	1
---UUCUCAACGUCGUAGAACAAGCCCGA----------	1	0	1
----UCUCAUCGUCGUAGAACAAGCGCGAG----------	1	0	1
----UCUCGUCGUCGUAGAAGAAGCCCGA----------	2	0	1
----UCUCAUCGUCGUAGAACACGCCCGA----------	1	0	1
---AUCUCAUCGUCUUAGAACAAGCCCGA----------	2	0	1
----UCUCAUUGUCGUAGAACAAGCCCUA----------	2	0	1
--UUUCUCAUCGUCGUAGAAGAAGCCCG----------	1	0	1
----UCUCAUCGUCGUAGGAGAAGCCCGA----------	2	0	1
----UCUCAUCGUUGUAGAGCAAGCCCGA----------	2	0	1
---AUCUCAUCGUCGUAGGACAAGCCCGA----------	2	0	1
-----CCCAUCGUCGUAGAACAAGCCCGA----------	1	0	1
---UUCCCAUCGUCGUAGAACAAGCCCAA----------	2	0	1
---AUCUCAUCGUCGUAGAACAAGCUCGA----------	2	0	1
---GUCUCAUCGUCGUAGAACAAGCCCGU----------	2	0	1
----UCUCAUCUUCGUAGAACAAACCCGA----------	2	0	1
----UCUCAUCGUCGUAGAACAAGGCCGA----------	1	0	1
---UUCUCAUCGUCGUAGAACAGGCCCGA----------	1	0	1
----UCUCAUCGUCGUAGCAGAAGCCCGA----------	2	0	1
----CCUCAUCGUCGUAGAACAAGCCCAA----------	2	0	1
---AUCUCAUCGUCGUAGAACAAGUCCGAG----------	2	0	1
---AUCUCAUCGGCGUAGAACAAGCCCGA----------	2	0	1
----CCUCAUCGUCUUAGAACAAGCCCGA----------	2	0	1
----UCUCAUCGUCGUAGAACAAGCCCGAGU----------	1	0	1
CUUUUCUCAUCGUCGUAGAACAAGCCCGA----------	1	0	1
---AUCUCAACGUCGUAGAACAAGCCCGA----------	2	0	1
----UCUCAUCGUCGUAGAACAAUACCGA----------	2	0	1
----UCUCAUCGUCGUAGAGCAAGCCCUA----------	2	0	1
-----CUCAUCGUCGUUGAACAAGCCCGA----------	1	0	1
----UCUCAUCGUCGUAAAACAAGCCCAA----------	2	0	1
---UACUCAUCGUCGUAGAACAAGCCCGC----------	2	0	1
----UCUCAUCGUCGUAUAACAACCCCGA----------	2	0	1
---UUCUCAUCGUAGUAGAACAAGCCCGA----------	1	0	1
----UCUUAUCGUCGUAGAACAAGCCCUA----------	2	0	1
---UUCUCAUCGUCGUAGAACAAGCCCGAGG----------	0	0	1
----UCUCAUCGUCAUAGAACAAGCCCC----------	2	0	1
---UUCUCAUCGUCGUAGAACAAGCCGGA----------	1	0	1
------UCAUCGUCGUAGAACAAGCCCGAGG----------	0	0	1
---AUCUCAUCGUCGUAGAACAAGCCC----------	1	0	1
----CCUCUUCGUCGUAGAACAAGCCCGA----------	2	0	1
----UCUCAUCGUCUUAGAACAAGCCCAA----------	2	0	1
-------CAUCGUCGUAGAACAAGCCCGA----------	0	0	1
----UCUCAUCGUCGUAGAACAAGCCCGAGA----------	1	0	1
--GAUCUCAUCGUCGUAGAACAAGCCCG----------	2	0	1
----UCUCAUCGUCGAAGAACAAGCCCG----------	1	0	1
----UCUCAUCUUCGCAGAACAAGCCCGA----------	2	0	1
----UCUCAUCGUCGUAGAACAAUCCCGAG----------	1	0	1
----UCUCAUCGUCGUAGAACAAGCCCAAG----------	1	0	1
---AUCUCAUCGUCGCAGAACAAGCCCGA----------	2	0	1
--UUUCUCAUCGUCGUCGAACAAGCCCGA----------	1	0	1
----UCUCAUCGUCGGAGAACAAGCCCGA----------	1	0	1
---AUCUCAUCGUCGUAGAACCAGCCCGAG----------	2	0	1
----UCUCAUCGUCGUAGAACAUGCCCG----------	1	0	1
----UCUCAUCGUCGUAGAAGAAGCCCGAU----------	2	0	1
----UCUCAUCGUCGUAGGACAAGCCG----------	2	0	1
----UCUCAUCGUCGUAGAACAACCCCGAG----------	1	0	1
----UCUCAUCGUCGUAGAACAAGCUCGAGG----------	1	0	1
-AAUUCUCAUCGUCGUAGAACAAGCCCGA----------	2	0	1
---AUCUCAUCGUCGUAGGACAAGCCCG----------	2	0	1
---UUCUCAUCGUCGUAGAACAAGCC----------	0	0	1
---UACUCAUCGUCGUAGAACGAGCCCGA----------	2	0	1
----UCUCAUCGUCUUAGAACAAACCCGA----------	2	0	1
	Total	**0**	**2,228**

Pair	antisense Aub	antisense AGO3	Alignment	sense Aub	sense AGO3
1			5' UUCCUCGGAAUCAAGGGCAACGAGU 3'	4	1
			5' UUCCUCGGAAUCAAGGCCAACGAGU 3'	0	1
	22	4	3' GUCGACCUAGUUAACCAAGGAGCCUU 5'		
	8	1	3' UCGACCUAGUUAACCAAGGAGCCUU 5'		
	4	0	3' GUCGACCUGGUUAACCAAGGAGCCUU 5'		
	3	0	3' UCGACUUAGUUAACCAAGGAGCCUU 5'		
	1	2	3' UCGACCUAGUCAACCAAGGAGCCUU 5'		
	1	0	3' UCGACCUAUUUAACCAAGGAGCCUU 5'		
	1	0	3' GUCGACCUAGUCAACCAAGGAGCCUU 5'		
	1	0	3' UCGACCUAGGUAACCAAGGAGCCUU 5'		
	1	0	3' CGACCUAGUUGACCAAGGAGCCUU 5'		
	1	0	3' GUCGACUUAGUUAACCAAGGAGCCUU 5'		
	1	0	3' CCUCGACCUAGUUAACCAAGGAGCCUU 5'		
	0	22	3' CGUCGACCUAGUUAACCAAGGAGCCUU 5'		
	0	2	3' UCGUCGACCUAGUCAACCAAGGAGCCUU 5'		
	0	2	3' CGUCGACCUGGUUAACCAAGGAGCCUU 5'		
	0	2	3' CGUCGAUCUAGUUAACCAAGGAGCCUU 5'		
	0	1	3' GUCGACCUAGAUAACCAAGGAGCCUU 5'		
2			5' UGGGCCUGGAGUACUUCAGCCAGA 3'	0	3
	6	0	3' GUGCAAGUUGGUCUACCCGGACCU 5'		
	4	1	3' UGUGCAAGUUGGUCUACCCGGACCU 5'		
	2	0	3' CUGUGCAAGUUGGUCUACCCGGACCU 5'		
3			5' UCGACAGUUCCUCGGGCUUGUUCUAC 3'	0	3
	1	0	3' GGACUUCGGCCACAAGCUGUCAAG 5'		
4			5' UGUUCUUCUAGGGAUGGCGUGAA 3'	0	1
	15	0	3' CAGUUCGCGAAGUUACAAGAAGAU 5'		
	2	0	3' UCAGUUCGCGAAGUUACAAGAAGAU 5'		
	1	0	3' UCAGUUCGCGAACUUACAAGAAGAU 5'		
5			5' UUCAACCAGAUGGGCCUGGAGUACUUC 3'	0	1
			5' UUCAACCAGAUGGGCCUGGAGUA 3'	0	1
	1	1	3' UGAAAGUCCUGUGCAAGUUGGUCU 5'		
	1	0	3' GAAAGUCCUGUGCAAGUUGGUCU 5'		
	1	0	3' GAUUAAAGUCCUGUGCAAGUUGGUCU 5'		
	78	38		4	11

Su(Ste) pair4 piRNA is highlighted with green underline.

Supplementary Table 4
Nagao et al.

AT-chX-1 (common)

Alignment	Mismatches	AGO3	Aub
---UGUUUCAUCGUUAGACGGCUCGGGC----	0	2,642	13,615
---UGUUUCAUCGUUAGACGGCUCGGG-----	0	875	3,013
--AUGUUUCAUCGUUAGACGGCUCGGGC----	1	6	190
---UGUUUCAUCGUUAGACGGCUCGGCC----	1	11	183
---UGUUUCAUCGUUAGACGGCUCGGGU----	1	39	172
---UGUUUCAUCGUUAGACGGCUCGG------	0	308	134
---UGUUUCAUCGUUAGACGGCUCGGGCG---	0	693	91
---UGUUUCAUCGUUAGACGACUCGGGC----	1	15	88
---UGUUUCAUCUUUAGACGGCUCGGGC----	1	7	79
---UGUUUCAUCGUUAGACGGCUCGGC-----	1	11	55
---UGUUUCAUCGUUAGACGGCUCGGGA----	1	5	45
---UGUUUCAUCGUUAGACGGCUCGGGCC---	1	1	43
---UGUUUCAUCGUUAGACGGCUCGGA-----	1	5	42
---UUUUUCAUCGUUAGACGGCUCGGGC----	1	2	36
---UGUUCCAUCGUUAGACGGCUCGGGC----	1	4	30
---UGUCUCAUCGUUAGACGGCUCGGGC----	1	12	29
---UGUUUCAUCGUUAUACGGCUCGGGC----	1	1	29
-UCUGUUUCAUCGUUAGACGGCUCGGG-----	0	1	29
---UGUUUCAUCGUUAGACGGCUCGGGUU---	2	1	29
--GUGUUUCAUCGUUAGACGGCUCGGGC----	1	4	26
--AUGUUUCAUCGUUAGACGGCUCGGG-----	1	2	26
---UGCUUCAUCGUUAGACGGCUCGGGC----	1	6	25
-UCUGUUUCAUCGUUAGACGGCUCGGGC----	0	1	25
---CGUUUCAUCGUUAGACGGCUCGGGC----	1	7	24
---UGUUUCAUCGUUAGACGUCUCGGGC----	1	1	22
---UGUUUCAUCGUUAGACGACUCGGG-----	1	1	22
----GUUUCAUCGUUAGACGGCUCGGGC----	0	2	17
---UGUUUCAUCGUUAGACGGCUCGGGCA---	1	13	16
---UGUUUCAUCGCUAGACGGCUCGGGC----	1	2	16
----UUUUCAUCGUUAGACGGCUCGGGC----	1	2	16
----UGUUCAUCGUUAGACGGCUCGGGC----	2	4	15
---UGUUUCAUUGUUAGACGGCUCGGGC----	1	3	15
--CUGUUUCAUCGUUAGACGGCUCGGG-----	0	2	15
---UGUUUCAUCGUUAGACGGCUCGGGUC---	2	24	14
----UGUUCAUCGUUAGACGGCUCGGG-----	2	3	13
---UGUUUCAUCGUUAGGCGGCUCGGGC----	1	3	13
---UGUUACAUCGUUAGACGGCUCGGGC----	1	5	12
---UGUUUCGUCGUUAGACGGCUCGGGC----	1	4	12
---UGUUUCAUCGUUAGACGACUCGGGCG---	1	3	11
---UGUUUCAUCGUUAGACGGCUCGGGAA---	2	1	11
---UGUUUCAUCGUCAGACGGCUCGGGC----	1	1	11
---UGUUUCAUCGUUAGACGGCACGGGC----	1	4	10
---UGUUUCAUCGUUAGACGGCUCGUGC----	1	2	10
---UGUUUCAUCGUUAGACGGCCCGGGC----	1	1	10
---UGUUUCAUCGUUAGACGGCUCUGG-----	1	1	10
---UGUUUCAUCGUUAGACAGCUCGGGC----	1	1	10
---UGUUUCAUCGUUGGACGGCUCGGGC----	1	3	9
---UGUUUCAUCGUUAGACGGCUCG-------	0	3	9
---UGUCUCAUCGUUAGACGGCUCGGG-----	1	7	8
---UGUUUCAUCGUUAGACGGCUCGGGG----	1	3	8
---CGUUUCAUCGUUAGACGGCUCGGG-----	1	2	8
---UGUUUCAUCGUUAGACGGCUCUGGC----	1	1	8
---UGUUUCAUCGUUAGACGGCUCGGUC----	1	1	8
---UGUUUCAUCGUUAGACGGCUCAGGC----	1	10	7
--CUGUUUCAUCGUUAGACGGCUCGGGC----	0	7	7
---UAUUUCAUCGUUAGACGGCUCGGGC----	1	1	7
---UGUUUCAUCGUUAGACGGCUCGGGGC---	2	5	6
---UGUUUCAUCGUUAGGCGGCUCGGG-----	1	1	6
---UGUUUCAUCGUUAGACGGCUCGGAC----	1	10	5
---UGGUUCAUCGUUAGACGGCUCGGGC----	1	4	5
---UGUUUCAUCGUUAGACGGCUGGGC-----	2	1	5
---UGUUCCAUCGUUAGACGGCUCGGG-----	1	3	4
---UGUUUCAUCGUUAGACGGCUCGGCU----	2	3	4
---UGUUUCAUCGUUAGACCGCUCGGGC----	1	2	4
---UGUUUCAUCGUUAAACGGCUCGGGC----	1	2	3
---UGCUUCAUCGUUAGACGGCUCGGG-----	1	2	3

---UGUUUCAUCGUUAGACGCCUCGGGC----	1	1	3
---UGUUUCAUCGUUAGUCGGCUCGGG-----	1	1	2
---UGUAUCAUCGUUAGACGGCUCGGGC----	1	1	2
---UGUUUCAUCGUUAGACGGCUCGGGCGU--	1	35	1
---UGUUUCAUCGUUAGACGGCUCGGU-----	1	9	1
--AUGUUUCAUCGUUAGACGGCUCGG------	1	2	1
---UGUUUCACCGUUAGACGGCUCGGGC----	1	2	1
---UGUUUCUUCGUUAGACGGCUCGGGC----	1	1	1
---UGUUUCAUCGUUAGACGGCUCGCGC----	1	1	1
---UGUUUCAUCGUUAGACGGCUCGGGUG---	1	1	1
---AGUUUCAUCGUUAGACGGCUCGGG-----	1	1	1
---GGUUUCAUCGUUAGACGGCUCGGGC----	1	1	1
--AUGUUUCAUCGUUAGACGGCUCGGGCG---	1	1	1
	Total	**4,870**	**18,500**

AT-chX-1 (AGO3 unique)

Alignment	Mismatches	AGO3	Aub
---UGUUUCAUCGUUAGACGGCUCGGGCGG--	0	175	0
---UGUUUCAUCGUUAGACGGCUCGGGUCG--	2	43	0
---UGUUUCAUCGUUAGACGGCUCGGGCGGA-	0	21	0
---UGUUUCAUCGUUAGACGACUCGGGCGG--	1	10	0
---UGUUCCAUCGUUAGACGGCUCGGGCGG--	1	7	0
---UGUUUCAUCGUUAGACGGCUCGGGCGC--	1	7	0
--AUGAUUCAUCGUUAGACGGCUCGGGC----	2	5	0
---UGUUUCAUCGUUAGACGACUCGGGCGU--	2	5	0
---UGUUUCAUCGUUAGACGGCUCGGGCAU--	2	4	0
----UGUUCAUCGUUAGACGGCUCGGGCG---	2	4	0
---UGUAUCAUCGUUAGACGGCUCGGC-----	2	3	0
---UGCUUCAUCGUUAGACGGCUCGGGCG---	1	3	0
---UGUUUCAUCGCUAGACGGCUCGGGCGG--	1	3	0
---UGUUUCAUCGUUAGACGGCUCGGCG----	2	2	0
---UGCUUCAUCGUUAGACGGCUCGGCC----	2	2	0
---UGUUUCGUCGUUAGACGGCUCGG------	1	2	0
---UGUUUCAUCGUUAGACGGCUCGGGCGA--	1	2	0
---UGUUUCAUCGUUAGACGGCUCGU------	1	2	0
---UAUUUCAUCGUUAGACGGCUCGGG-----	1	1	0
---UGUUUCAUCGUUAGAUGGCUCGGGCG---	1	1	0
---UGUUUCAUCGUUAGACGUCUCGGGCG---	1	1	0
---UGUUUCAUCGUUAGACUACUCGGGC----	2	1	0
---UGUUUCAUCGAUAGACGGCUCGGG-----	1	1	0
---UGUUUCAUCGUCAGACGGCUCGG------	1	1	0
---UGUUUCAUCGUUAUACGGCUCGGGCG---	1	1	0
---UGUUUCAUCGUUAGACGGCUCGGACG---	1	1	0
---UGUUUCAUCGUUAGACGGCUCGUGCG---	1	1	0
---UGUUUCAUCGUUAGGCGACUCGGG-----	2	1	0
---UGUUUCAUCGUUAGACGGCUCGGGAG---	1	1	0
---UGUUUCAUCGUUAGCCGGCUCGGGCG---	1	1	0
---UGUUUCAUCGUUAGACGGCUCAG------	1	1	0
---UAUUUCAUCGUUAGACGGCUCGGGCG---	1	1	0
--CUGUUUCAUCGUUAGACGGCUCGGGCGG--	0	1	0
---UGUCUCAUCGUUAGACGGCUCGGGCG---	1	1	0
---UGAUUCAUCGUUAGACGGCUCGGGCG---	1	1	0
---UGUUUCAUCGUUAGACGGCUCGGGCAA--	2	1	0
---UAUUUCAUCGUUAGACGGCUCGGC-----	2	1	0
---UGUUUCAUCGUUAGACGGCCCGGG-----	1	1	0
---UGUUUCAUCGUUAGACCGCUCGGGCG---	1	1	0
---UGAUUCAUCGUUAGACGGCUCGGG-----	1	1	0
---UGUUUCAUCGUUAGACGGCCCGGGCG---	1	1	0
---UGUUUCAUCGUUAGGCGGCUCGG------	1	1	0
---UGUUUCAUCGUUAGGCGGCUCGGGCG---	1	1	0
---UGUUUCAUCGUUAGACGGCUUGGG-----	1	1	0
---UGUUUCAUCGUUAGACGGCGCGGG-----	1	1	0
	Total	**327**	**0**

AT-chX-1 (Aub unique)

Alignment	Mismatches	AGO3	Aub
---UGUUUCAUCGUUAGACGGCUGGGGC----	1	0	35
---UGUUUCAUCGUUAGACGGCUAGGGC----	1	0	33
---UGUUUCAUCUUUAGACGGCUCGGG-----	1	0	21
---UGUUUCAUCGUUAGACGGCUUGGGC----	1	0	19
--AUGUUUCAUCGUUAGACGGCUCGGGCC---	2	0	19
---UGUUUCAUCGUUAGACGUCUCGGG-----	1	0	19
---UGUUUCAUCGUUAGAUGGCUCGGGC----	1	0	18
-----UCUCAUCGUUAGACGGCUCGGG-----	1	0	15
---UCUUUCAUCGUUAGACGGCUCGGGC----	1	0	15
---UGUUUCAUCGUUAGACUGCUCGGGC----	1	0	14
-UCUGUUUCAUCGUUAGACGGCUCGG------	0	0	13
---UGUUUCAUCGUUAGACGGCUCGGGCU---	1	0	12
----GUUUCAUCGUUAGACGGCUCGGG-----	0	0	12
---UGUUUCAUCGUUAGACGGUUCGGGC----	1	0	12
----UUUUCAUCGUUAGACGGCUCGGG-----	1	0	11
---UGUUUCAUCGUUAGACGGCUCCGGC----	1	0	10
---UGUUUCAUCGUUAGACUGCUCGGG-----	1	0	10
---UUUCUCAUCGUUAGACGGCUCGGGC----	2	0	9
-----UCUCAUCGUUAGACGGCUCGGGC----	1	0	9
---UGGUUCAUCGUUAGACGGCUCGGG-----	1	0	8
-----UUUCAUCGUUAGACGGCUCGGG-----	0	0	7
---UGUUUUAUCGUUAGACGGCUCGGGC----	1	0	6
---UGUUACAUCGUUAGACGGCUCGGG-----	1	0	6
---UGUUUCAUCCUUAGACGGCUCGGGC----	1	0	6
---UGUUUCAUCGUUAGACGGCUCGA------	1	0	6
---UGUUUCAUAGUUAGACGGCUCGGGC----	1	0	6
---UGUUUCAUCGUCAGACGGCUCGGG-----	1	0	6
---UGUUUCAUCGUUAGACGGCUCGGCCCG--	2	0	5
---UGUUUCAUCGUUAGACGGCUGGC------	2	0	5
---UGUUUCACCGUUAGACGGCUCGGG-----	1	0	5
---CGUUUCAUCGUUAGACGGCUCGGCC----	2	0	4
---UGUUUCAUCGUUAUACGGCUCGGC-----	2	0	4
---UCUUUCAUCGUUAGACGGCUCGGG-----	1	0	4
---UGUUUCAUCGUUUGACGGCUCGGGC----	1	0	4
---UGUUUAAUCGUUAGACGGCUCGGG-----	1	0	4
---AUUUUCAUCGUUAGACGGCUCGGGC----	2	0	4
---UGUUUCAUCGUUAGACAGCUCGGG-----	1	0	4
-----UUUCAUCGUUAGACGGCUCGGGC----	0	0	4
---UGUUUAAUCGUUAGACGGCUCGGGC----	1	0	4
---UGUUUCAUCGUUAGACGACUCGGGCU---	2	0	4
---UGUUUCAUCAUUAGACGGCUCGGGC----	1	0	4
---UGUUUCAUCGUUAGACGGCACGGG-----	1	0	4
---UGUUUCAUCGUUAGACGGCUCGGGCCG--	1	0	4
---UGUUUCAUCGUUAGACGCCUCGGG-----	1	0	3
---CGUUUCAUCGUCAGACGGCUCGGGC----	2	0	3
-----UGUCAUCGUUAGACGGCUCGGGC----	1	0	3
---UGCUUCAUCGUUAGACGGCUCGCGC----	2	0	3
---UGUUUCAUCGUUAGACGACUCUGGC----	2	0	3
--GUGUUUCAUCGUUAGACGGCUCGGGU----	2	0	3
---UGAUUCAUCGUUAGACGGCUCGGGC----	1	0	3
---UGUUUCAUCGUUAGACGGCUAGGG-----	1	0	2
---UGUUUCAUCGUUAGAUGGCUCGGG-----	1	0	2
---UGUUUCAUCGUUAGACGGCUCGGCCC---	2	0	2
---UGUUUCAUCGUUAGACGGCUCGAGC----	1	0	2
-AAUGUUUCAUCGUUAGACGGCUCGGGC----	2	0	2
---UGUUUCAUCGUUAGACGGCUCCGCC----	2	0	2
---UUUUUCAUCGUUAGACGGCUCGGG-----	1	0	2
---UGUUUCAUCGUUAGACGGCUCGGGUA---	2	0	2
---UGUUUCAUCGUUACACGGCUCGGG-----	1	0	2
---UGUCUCAUCGUUAUACGGCUCGGGC----	2	0	2
--AUGUUUCAUCGUUAUACGGCUCGGGC----	2	0	2
---UGUUUCAUCGUUAGACGGCUCGGUCC---	2	0	2
---UGUUUCAUAGUUAGACGGCUCGGCC----	2	0	2
---UGUUUCAUCGUUAGACGGCUCGAA-----	2	0	2
---UGUUUCAUAGUUAGACGGCUCGGG-----	1	0	1
---UGUUUCAUCGUUAGACUGCUCGGCC----	2	0	1
---UGUUUCAUCGUUAGACGGCUCGUG-----	1	0	1
---UGUUUCAUCGUUAUACUGCUCGGGC----	2	0	1

---UGUUUCAUCGUUAGACGGCUCGGCCG---	1	0	1
---UGUUUCAUCGUUAGACGGCUCGGGCCUA-	2	0	1
---UGUUUCAUCGUAAGACGGCUCGGGC----	1	0	1
--CUGUUUCAUCGUUAGACGGCUCCGGCC---	2	0	1
---UGUUUCAUCGUUAAACGGCUCGGGA----	2	0	1
---UGUCUCAUCGUUAGACGACUCGGGC----	2	0	1
---UGUGUCAUCGUUAGACGGCUCGGG-----	1	0	1
---UGUUUCAUCGUUAGACGGCUCGGGCUC--	2	0	1
-----UCUCAUCGUUAGACGGCUCGG------	1	0	1
---UGUUUCAUCGUUAGACUGCUCGGC-----	2	0	1
---UGUUUCAUCGUUAGACGGCUCGGCCGAAA	2	0	1
---UGUUUCAUCGCUAGACGGCUCGG------	1	0	1
---UGUUUCAUCGUUAGACGUCUCGUGC----	2	0	1
-AAUGUUUCAUCGUUAGACGGCUCGGG-----	2	0	1
--AUGUUUCAUCGUUAGACGGCUCGGGCU---	2	0	1
---CGUUUCAUCGUUAGACUGCUCGGG-----	2	0	1
---UGUUUCAUCGUUAGACGGCUCGCC-----	2	0	1
---UGUUUCAACGUUAGACGGCUCGGGC----	1	0	1
---UGUUUCAUCGUUAUACGGCUCGG------	1	0	1
---UGUUUCAUCGUUAGACGGCUCCAG-----	2	0	1
--GUGUUUCAUCGUUAGACGGCACGGGC----	2	0	1
---UGUUUCAUCUUUAGACGGCUCGGCC----	2	0	1
---UGUUUCAUCGUUAGACGGCUCAGG-----	1	0	1
-----UCUCAUCGUUAUACGGCUCGGGC----	2	0	1
---AGUUUCAUCGUUAGACGGCUCGGGC----	1	0	1
---UGUUUCAUCGCUAGACGGCUCGGG-----	1	0	1
---UGUUUCAUCGUUAGAAGGCUCGGGC----	1	0	1
-UCUGUUUCAUCGUUAUACGGCUCGGGC----	1	0	1
---UGUGUCAUCGUUAGACGGCUCGGGC----	1	0	1
UACUGUUUCAUCGUUAGACGGCUCGGG-----	2	0	1
--GUGUUUCAUCGUUAGACGGCUCGGGCU---	2	0	1
---UGUUUCAUCGUUAGGCGGCUCGGGCCG--	2	0	1
--CUGUUUCAUCGUUAGGCGGCUCGGG-----	1	0	1
--AUGGUUCAUCGUUAGACGGCUCGGGC----	2	0	1
---UGUUUCAUCGUUAGACUUCUCGGGC----	2	0	1
--CUGUUUCAUCGUUAGACGGCUCCGG-----	1	0	1
---UGUUUCAUCGUUAGUCGGCUCGGGC----	1	0	1
---UGUUUCAUCGUUAGACGGCCCCGGC----	2	0	1
---UGUUUCGUCUUUAGACGGCUCGGGC----	2	0	1
---UUUUUCAUCGUUAGACGGCUCUGGC----	2	0	1
---UGUCUCAUCGUUAGACGGCUCGG------	1	0	1
---UGUUUCAUCGUUAGACGGCUCGGGGCG--	2	0	1
---UGUUUCAUUGUUAGACGGCUCGGGU----	2	0	1
---UGUUUCAUCGGUAGACGGCUCGGGC----	1	0	1
--AUGUUUCAUCGUUAGACGACUCGGGC----	2	0	1
---UUUUUCAUCGUUAGACGACUCGGGC----	2	0	1
-UCUGCUUCAUCGUUAGACGGCUCGGG-----	1	0	1
---UGUUUCAUCGUUAGACGGCUGGG------	1	0	1
---UGUUUCAUCGUUAGACGGCUCGAG-----	1	0	1
--AUGUUUCAUCGUUAGACGGCUCGGAC----	2	0	1
--AAGUUUCAUCGUUAGACGGCUCGGGC----	2	0	1
---UGUUUCAUCGUUAGACGGCUCCGGG----	2	0	1
---UGUUUCAUCGUUAGACGGCACGGGCG---	1	0	1
---UGUUUCAUCGUUAGACGGCUCGGCCGG--	1	0	1
---UGUUUCAUCGUUAUACGGCUCGGA-----	2	0	1
---CGUUUCAUCGUUAGACGGCUCGGGA----	2	0	1
---UGUUUCAUCUUUAGACGGCUCGGA-----	2	0	1
---UGUUUCAACGUUAGACGGCUCGGG-----	1	0	1
---UGUUUCAUCGUUAGAAGGCUCGGG-----	1	0	1
---UGUUUCAUCGUUACACGGCUCGG------	1	0	1
---UUUUUCAUCGUUAGACGGCUCGGGCG---	1	0	1
---UGUUUCAUCGUUAGACGGCUCGGGCCU--	2	0	1
--AUGUUUCAUCGUUAGACGGCUCGGGA----	2	0	1
-UCUGUUUCAUCGUUAGACGACUCGGG-----	1	0	1
----UUUUCAUCGUUAGACGGCUCGUGC----	2	0	1
---UGUUUCAUCGUUAAACGGCUCGGG-----	1	0	1
-GAUGUUUCAUCGUUAGACGGCUCGG------	2	0	1
----GUUUCAUCGUUAGACGGCUCGGGUC---	2	0	1
-AGUGUUUCAUCGUUAGACGGCUCGGGC----	2	0	1
---UGUUUCAUCGAUAGACGGCUCGGGC----	1	0	1

---UGUUUCAUCGUUAGACGGAUCGGGC----	1	0	1
---AUUUUCAUCGUUAGACGGCUCGGG-----	2	0	1
--CUGUUUCAUCGUUAGACGGCUAGGG-----	1	0	1
---UGUUUCAUCGUUAUACGGCUCGGGCA---	2	0	1
---UGUUUCAGCGUUAGACGGCUCGGGC----	1	0	1
--AUGUUUCAUCAUUAGACGGCUCGGGC----	2	0	1
---CGUUUCAUCGUUAGACGGCUCGUGC----	2	0	1
---UGUUUCAUCGUUAGACGUCUCGGC-----	2	0	1
---UGUUUCAUCGUUGGACGGCUCGG------	1	0	1
----UUUUCAUCUUUAGACGGCUCGGGC----	2	0	1
--AUGUUUCAUCGUUAGACGGCCCGGGC----	2	0	1
---UGUUUCAUCUUUACACGGCUCGGG-----	2	0	1
	Total	**0**	**559**

qRT-PCR primer

Gene	Forward primer	Reverse primer
RP49	5'-CCGCTTCAAGGGACAGTATCTG-3'	5'-ATCTCGCCGCAGTAAACGC-3'
Aub	5'-GGGATGACCAGCAAGAAAGG-3'	5'-AAGACCGTCGCAGTCGTGTA-3'
AGO3	5'-ACGCCCACTTATCTTGTTGG-3'	5'-CTGCATTTGTGGCCTCCATA-3'
Stellate	5'-GTCAGAGCGTGGCGTGATTGC-3'	5'-GGCCCGAACATCGCTCCGTCC-3'
1731	5'-AGCAAACGTCTGTTGGAAGG-3'	5'-CGACAGCAAAACAACACTGC-3'
412	5'-CACCGGTTTGGTCGAAAG-3'	5'-GGACATGCCTGGTATTTTGG-3'
HMS-Beagle	5'-AAGGGCAGCAATCCGAACTA-3'	5'-AGGTGACCTTGAGGGTCGAA-3'
Stalker-2	5'-GGGTTAAAACCGCTCCACAA-3'	5'-TTGAATGGTCAAGCGAATGC-3'
Stalker-4	5'-GGAGCCACAATTCTCGAAGG-3'	5'-TTCTGCTGCCTGGCTCTGTA-3'
Diver-2	5'-CTTCAGCCAGCAAGGAAAAC-3'	5'-CTGGCAGTCGGGTGTAATTT-3'
gtwin	5'-TTCGCACAAGCGATGATAAG-3'	5'-GATTGTTGTACGGCGACCTT-3'
Idefix	5'-AACAAAATCGTGGCAGGAAG-3'	5'-TCCATTTTTCGCGTTTACTG-3'
mdg1	5'-CACATGTTCTCATTCCCAACC-3'	5'-TTCGCTTTTTATATTTGCGCTAC-3'
roo	5'-CGTCTGCAATGTACTGGCTCT-3'	5'-CGGCACTCCACTAACTTCTCC-3'
Doc	5'-GGGTGACTATAACGCCAAGC-3'	5'-GCAAAATCGATCAGGTCTGG-3'
F-element	5'-GCTGGTAGATACCGCTGAGG-3'	5'-GTAGTCGTCCTCCGTTTTCG-3'
Het-A	5'-CGCGCGGAACCCATCTTCAGA-3'	5'-CGCCGCAGTCGTTTGGTGAGT-3'
Helena	5'-TGCGAGTACTCCAACGCTGT-3'	5'-AGCAGTAGGGCGCAGGTAGA-3'
I-element	5'-TGAAATACGGCATACTGCCCCCA-3'	5'-GCTGATAGGGAGTCGGAGCAGATA-3'
R1	5'-CGTTGTTTCCACTGCCGTTA-3'	5'-CCTAGGCTGCGGAAACTGAT-3'
TART	5'-AGAGAGGGAAAGAAGGGAAAGGGA-3'	5'-ATTTCCTGCCTGGTTAGATCGCCA
hopper2	5'-GAACTACCGAATCCGCCTTG-3'	5'-TGGAGAATCGTGGGTGGAAT-3'

DNA oligo (for Northern blot)

	piRNA	probe
Su(Ste)-4 antisense	5'-TCTCATCGTCGTAGAACAAGCCCGA-3'	5'-TCGGGCTTGTTCTACGACGATGAGA-3'
Su(Ste)-4 sense	5'-GACGATGAGAAAAAGTGGTACGGCA-3'	5'-TGCCGTACCACTTTTTCTCATCGTC-3'
AT-chX-1 antisense	5'-TGTTTCATCGTTAGACGGCTCGGGC-3'	5'-GCCCGAGCCGTCTAACGATGAAACA-3'
AT-chX-1 sense	5'-CGATGAAACAGAGATCTTCAGCAGT-3'	5'-ACTGCTGAAGATCTCTGTTTCATCG-3'
Su(Ste)-6 antisense	5'-TAGATCTTCACCGTTGACTAGCCCCA-3'	5'-TGGGGCTAGTCAACGGTGAAGATCTA-3'
Su(Ste)-6 sense	5'-TGAAGATCTACTGCCCACGCTGTAAA-3'	5'-TTTACAGCGTGGGCAGTAGATCTTCA-3'
Su(Ste) pair4-antisense	5'-TAGAAGAACATTGAAGCGCTTGAC-3'	5'-GTCAAGCGCTTCAATGTTCTTCTA-3'
AT-chX 71 antisense	5'-TTAATGCCGGAAGCGATCCCACTGC-3'	5'-GCAGTGGGATCGCTTCCGGCATTAA-3'
AT-chX 71 sense	5'-CCGGCATTAATTTCTTTAAATATGA-3'	5'-TCATATTTAAAGAAATTAATGCCGG-3'
U6 snRNA		5'-GGGCCATGCTAATCTTCTCTGTA-3'