

Repeat Class	Repeat Family	Repeat Subfamily	Chromosome	Start	End	Strand	10 dpp	14 dpp	Adult
SINE	Alu	B1_Mus2	chr14	44062955	44063100	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	44293009	44293043	+	0.923247	0.380568	1.84597
Simple_repeat	Simple_repeat	(A)n	chr14	44433069	44433104	+	0	0	0
SINE	Alu	B1_Mus2	chr14	44844922	44845065	+	0.591053	0	0
LTR	MaLR	MTA_Mm-int	chr14	44845066	44845846	+	0.591053	0	0
SINE	B2	B2_Mm1a	chr14	45456446	45456640	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	45456641	45456684	+	0	0	0
LINE	L1	L1Md_T	chr14	45633332	45639965	-	0	0.708814	0
SINE	B2	B2_Mm1a	chr14	46189145	46189333	-	0.332194	0	0
SINE	Alu	B1_Mus2	chr14	46201983	46202128	-	0.591053	0	0
SINE	Alu	B1_Mm	chr10	8402574	8402721	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	8402722	8402744	+	0.591053	0	0
SINE	Alu	B1_Mus2	chr14	47218365	47218512	-	0	0	0
LTR	MaLR	MTA_Mm	chr14	48243261	48243656	-	0	0	0
SINE	B4	B4	chr14	48631557	48631847	+	0	0	0
LTR	MaLR	MTA_Mm	chr14	49155242	49155635	+	0	0	0
SINE	Alu	B1_Mus2	chr14	49180083	49180220	+	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr14	49418185	49418224	+	0	0	0
SINE	B2	B2_Mm1t	chr14	50604083	50604274	+	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr14	51264803	51264846	+	0	0	0
SINE	Alu	B1_Mus2	chr10	59688638	59688784	-	0	0	0
SINE	Alu	B1_Mm	chr14	51625472	51625614	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	51625615	51625647	+	0	0	0
LINE	L1	L1_Mus3	chr14	52087084	52093584	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	52116735	52116775	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	52118910	52118936	+	0	0	0
SINE	Alu	B1_Mus2	chr14	52592394	52592539	-	0	0	0
SINE	Alu	B1_Mus1	chr14	52691766	52691911	+	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	52691912	52691954	+	0	0	0
SINE	Alu	B1_Mus2	chr14	52859736	52859882	-	0	0.708814	0.342623
SINE	Alu	B1_Mus1	chr14	52874067	52874211	+	0	0	0.342623
LTR	MaLR	MTA_Mm	chr14	53044740	53045135	+	0	0	0
LTR	MaLR	MTA_Mm	chr14	53631391	53631786	+	0	0	0
SINE	Alu	B1_Mus2	chr14	53727913	53728061	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	53728062	53728087	+	0	0	0
LTR	MaLR	MTA_Mm	chr14	53942513	53942908	+	0	0	0
LTR	MaLR	MTA_Mm	chr14	54331202	54331598	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	54776580	54776616	+	0	0	0
SINE	Alu	B1_Mus1	chr14	54889908	54890055	-	0.332194	0.761137	0.957947
LTR	MaLR	MTA_Mm	chr14	54956750	54957144	-	0	0	0
Low_complexity	Low_complexity	A-rich	chr14	55112093	55112150	+	0	0	0
LTR	MaLR	MTB	chr14	55178677	55179074	-	0	0	0
SINE	Alu	B1_Mm	chr14	55225561	55225707	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	55225708	55225728	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	55404825	55404852	+	0	0	0
SINE	Alu	B1_Mm	chr14	56008850	56008978	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	56109445	56109491	+	0	0	0.615324
SINE	Alu	B1_Mm	chr14	57021154	57021288	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	57021289	57021314	+	0	0	0
LTR	ERVK	RMER4A	chr14	57280971	57281404	-	0	0.380568	0
SINE	Alu	B1_Mus1	chr14	57281405	57281552	-	0	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr14	57450005	57450042	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	58528342	58528371	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	59671374	59671403	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	59676380	59676407	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	59890064	59890100	+	0	0	0.957947
SINE	Alu	B1_Mm	chr14	60922559	60922700	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	60976143	60976188	+	0	0	0
SINE	Alu	B1_Mus2	chr10	60619009	60619155	+	0	0	0
Low_complexity	Low_complexity	A-rich	chr10	60619156	60619212	+	0	0	0.615324
SINE	B2	B2_Mm1a	chr14	61742582	61742774	-	0	0	0
SINE	B4	B4A	chr14	62120452	62120754	-	0	0	0
SINE	Alu	B1_Mus1	chr14	62542202	62542348	-	0	0	0
SINE	Alu	B1_Mus1	chr14	62818141	62818292	-	0	0	0
LTR	MaLR	MTA_Mm	chr14	62890901	62891296	-	0	0	0
SINE	Alu	B1_Mm	chr14	63248178	63248313	+	0	0.380568	0

Simple_repeat	Simple_repeat	(GAAA)n	chr14	63248314 63248402 +	0.332194	0.380568	0
SINE	Alu	B1_Mm	chr14	64211192 64211337 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	64211338 64211377 +	0	0	0
SINE	Alu	B1_Mus2	chr14	65264333 65264466 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	65264467 65264487 +	0	0	0
SINE	B2	B2_Mm1t	chr14	65722060 65722255 +	0	0	0
SINE	B2	B2_Mm1a	chr14	65725949 65726134 -	0	0	0.342623
SINE	Alu	B1_Mm	chr14	65735920 65736065 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	65736066 65736087 +	0	0	0
LTR	MaLR	MTA_Mm	chr14	65807962 65808357 +	0	0	0
SINE	Alu	B1_Mus1	chr14	66398026 66398172 -	0	0	0
SINE	Alu	B1_Mus2	chr14	66415713 66415857 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	66643777 66643801 +	0.332194	0	0
LTR	ERVK	RLTR45	chr14	66643802 66644272 +	0.332194	0	0
LTR	MaLR	MTA_Mm	chr14	68083178 68083562 -	0	0	0
SINE	Alu	B1_Mus2	chr14	68216752 68216898 -	0	0	0
SINE	Alu	B1_Mus1	chr14	68565890 68566037 -	0	0	0
SINE	Alu	B1_Mm	chr14	68648354 68648499 +	0	0	0.342623
Low_complexity	Low_complexity	A-rich	chr14	68648500 68648542 +	0	0	0
SINE	Alu	B1_Mm	chr14	68678129 68678273 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	69152100 69152134 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	69377617 69377647 +	0	0	0
SINE	Alu	B1_Mm	chr14	69990500 69990641 +	1.18211	0	0.957947
Simple_repeat	Simple_repeat	(A)n	chr14	69990642 69990670 +	0	0	0.957947
SINE	Alu	B1_Mur4	chr14	70766020 70766197 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	71914209 71914247 +	0	0	0
LINE	L1	L1Md_T	chr14	72184355 72191900 -	0	0	0
SINE	Alu	B1_Mur4	chr14	72333049 72333196 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	72333197 72333220 +	0.332194	0	0
SINE	Alu	B1_Mm	chr14	72459596 72459741 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	72549353 72549394 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	72582849 72582901 +	0	0	0
LTR	MaLR	MTA_Mm	chr14	72774262 72774658 -	0	0	0
SINE	B2	B2_Mm1t	chr14	72844259 72844446 -	0	0	0
LTR	MaLR	MTA_Mm	chr14	73291278 73291672 -	0	0	0
SINE	Alu	B1_Mm	chr14	73405670 73405815 -	0	0	0
LTR	MaLR	MTA_Mm	chr14	74240571 74240965 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	74658741 74658775 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	75351422 75351462 +	0	0	0
SINE	Alu	B1_Mur4	chr14	75365503 75365650 -	0	0	0
SINE	Alu	B1_Mm	chr14	76238363 76238510 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr14	76238511 76238557 +	0	0	0
DNA	MER1_type	RCHARR1	chr14	76423579 76424390 -	0	0	0
SINE	Alu	B1_Mm	chr14	76584030 76584176 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	76584177 76584212 +	0.332194	0	0
SINE	Alu	B1_Mus2	chr14	76630104 76630249 +	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chr14	76630250 76630297 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	77093604 77093633 +	0	0	0
SINE	Alu	B1_Mus2	chr14	77339477 77339614 +	0.591053	0	0
SINE	Alu	B1_Mus2	chr14	78415097 78415241 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	78415242 78415295 +	0	0	0
SINE	Alu	B1_Mm	chr14	78789667 78789813 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	78789814 78789839 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	79713665 79713693 +	0	0	0
SINE	Alu	B1_Mus2	chr10	61979292 61979436 -	0.591053	0	0
LINE	L1	L1Md_T	chr14	80228611 80236116 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	80659908 80659979 +	0	0	0
scRNA	scRNA	BC1_Mm	chr14	80984419 80984567 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	81701811 81701858 +	0	0	0
SINE	Alu	B1_Mm	chr10	62068311 62068455 -	0	0	0
SINE	Alu	B1_Mm	chr14	81927179 81927325 -	0	0	0
LINE	L1	L1Md_T	chr14	82190724 82197632 -	0.591053	0	0
SINE	Alu	B1_Mm	chr14	82886887 82887033 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	82887034 82887059 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	83388941 83388995 +	0	0	0
SINE	Alu	B1_Mus1	chr14	83421236 83421368 -	0	0	0
SINE	Alu	B1_Mm	chr14	83573066 83573211 +	0	0.708814	0

Simple_repeat	Simple_repeat	(A)n	chr14	83573212 83573242 +	0	0.708814	0
LINE	L1	L1_Mus1	chr14	84333070 84339502 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	84546868 84546927 +	0.591053	0	0
LINE	L1	L1Md_F2	chr14	84709831 84715027 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	84834692 84834741 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr14	85312470 85312616 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	85469753 85469789 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	85495351 85495387 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	85679649 85679674 +	0	0	0
SINE	Alu	B1_Mm	chr14	85909745 85909886 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	85909887 85909920 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	86545624 86545652 +	0	0	0
Low_complexity	Low_complexity	GA-rich	chr14	86545653 86545829 +	0	0	0
LTR	MaLR	MTB	chr14	87647964 87648361 +	0	0	0
LTR	MaLR	MTA_Mm	chr14	87718375 87718769 +	0	0	0
SINE	Alu	B1_Mm	chr10	62374352 62374490 -	0	0	0
LTR	MaLR	ORR1B1	chr14	88398406 88398728 -	0	0	0
SINE	Alu	B1_Mm	chr14	88622483 88622628 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	89520173 89520204 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	89980350 89980389 +	0	0	0
SINE	Alu	B1_Mus1	chr14	90834208 90834354 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	90834355 90834400 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	91204171 91204196 +	0	0	0
LTR	MaLR	MTA_Mm	chr14	91688330 91688727 +	0	0	0
SINE	Alu	B1_Mus1	chr14	92378666 92378813 +	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chr14	92378815 92378886 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	92405030 92405070 +	0	0	0
LTR	ERVK	RMER6C	chr14	92546049 92546656 +	0	0	0
SINE	Alu	B1_Mm	chr14	92782547 92782692 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	92782693 92782715 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	93016557 93016593 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	93111829 93111907 +	0	0	0
Other	Other	RMER1B	chr14	95329437 95330393 +	0	0	0
SINE	Alu	B1_Mus2	chr14	95348287 95348431 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	95348432 95348488 +	0	0	0.342623
LTR	MaLR	MTA_Mm	chr14	97045928 97046323 -	0	0	0
LTR	MaLR	MTA_Mm	chr14	98107339 98107733 +	0	0	0
SINE	Alu	B1_Mur3	chr10	62884591 62884733 -	0	0	0
SINE	Alu	B1_Mus2	chr14	99381824 99381970 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	99381971 99381994 +	0	0	0
LTR	MaLR	MTA_Mm	chr14	99414897 99415290 -	0	0	0
SINE	Alu	B1_Mus2	chr14	99749395 99749543 -	0	0	0
LTR	MaLR	MLT1D	chr14	10017069 10017110 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr14	10075723 10075731 +	0	0	0
SINE	Alu	B1_Mm	chr14	10095447 10095462 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	10095462 10095465 +	0	0	0
SINE	Alu	B1_Mus1	chr14	10102155 10102167 -	0	0	0
SINE	B2	B2_Mm1a	chr14	10102481 10102500 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr14	10102500 10102503 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr14	10103243 10103256 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	10103256 10103261 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	10133167 10133172 +	0	0	0
LINE	L1	L1_Mus3	chr14	10155160 10155750 -	0	0	0
SINE	Alu	B1_Mm	chr14	10178348 10178363 -	0	0	0
LTR	MaLR	MTB	chr14	10253322 10253359 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	10257111 10257115 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	10443039 10443043 +	0	0	0
SINE	Alu	PB1	chr14	10549469 10549479 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	10557706 10557709 +	0	0.708814	1.48199
LTR	MaLR	MTA_Mm	chr14	10559163 10559202 +	0	0	0
SINE	Alu	B1_Mus2	chr14	10623259 10623274 -	0	0	0
SINE	Alu	B1_Mm	chr14	10623854 10623868 -	0	0	0
LTR	MaLR	MTA_Mm	chr14	10628477 10628515 +	0	0	0
LTR	MaLR	MTA_Mm	chr14	10650143 10650182 +	0	0	0
LINE	L1	L1Md_F2	chr14	10669398 10670009 -	0	0	0
SINE	Alu	B1_Mus1	chr14	10847815 10847830 -	0	0	0
SINE	Alu	B1_Mm	chr14	10918652 10918666 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr14	10918667 10918670 +	0	0	0
SINE	Alu	B1_Mus2	chr14	10922869 10922884 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	10922884 10922887 +	0	0	0
SINE	Alu	B1_Mus1	chr14	10923655 10923669 -	2.10535	2.12644	2.80392
LINE	L1	L1Md_F2	chr14	10962569 10963070 +	0	0	0
SINE	Alu	B1_Mus2	chr14	10999873 10999888 -	0	0	0
LTR	MaLR	MTA_Mm	chr14	11060949 11060988 +	0	0	0
LTR	MaLR	MTA_Mm	chr14	11160574 11160613 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	9011677 9011712 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	11164912 11164916 +	0	0	0
LINE	L1	L1_Mus1	chr14	11206243 11206848 +	0	0	0
SINE	Alu	B1_Mm	chr14	11213639 11213654 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	9020387 9020425 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	11237863 11237867 +	0	0	0
SINE	Alu	B1_Mus1	chr14	11241687 11241701 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	11241701 11241704 +	0	0	0
SINE	Alu	B1_Mm	chr14	11321486 11321501 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	11364173 11364177 +	0	0	0
LTR	MaLR	MTA_Mm	chr14	11378648 11378687 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	11413550 11413554 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	11440201 11440205 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	11470015 11470018 +	0	0	0
LTR	MaLR	MTA_Mm	chr14	11492620 11492660 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	11523098 11523107 +	0	0	0
SINE	Alu	B1_Mur4	chr14	11534407 11534420 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	11534420 11534423 +	0.332194	0	0
SINE	Alu	B1_Mm	chr10	64495753 64495899 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	64495900 64495943 +	0	0	0
LINE	L1	L1Md_T	chr10	64845963 64852617 -	0	0	0
SINE	Alu	B1_Mm	chr14	12035128 12035143 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	12047952 12047955 +	0	0	0
SINE	Alu	B1_Mus2	chr14	12119072 12119086 -	0	0	0
SINE	Alu	B1_Mus1	chr14	12123753 12123766 -	0	0	0
LTR	MaLR	MTA_Mm	chr14	12129336 12129375 +	0	0	0
SINE	Alu	B1_Mus2	chr14	12182666 12182681 -	0	0	0
SINE	Alu	B1_Mm	chr14	12220424 12220438 -	0	0	0
SINE	Alu	B1_Mus2	chr14	12227381 12227396 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	12227396 12227398 +	0	0	0
LTR	MaLR	MTA_Mm	chr14	12248761 12248800 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	12289134 12289140 +	0	0	0.957947
SINE	Alu	B1_Mur3	chr14	12292958 12292971 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr14	12292971 12292975 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr10	65177157 65177211 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	65207753 65207789 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	3807997 3808039 +	0	0	0
LINE	L1	L1Md_F3	chr15	4657870 4663803 -	0	0	0
LTR	MaLR	MTA_Mm	chr15	4827160 4827554 +	0.332194	0	0
LTR	MaLR	MTA_Mm	chr15	4828647 4829019 +	0	0	0
LINE	L1	L1Md_F2	chr15	5677950 5683288 -	0	0	0.615324
LTR	MaLR	MTB	chr15	5790441 5790841 +	0	0	0
LTR	MaLR	MTA_Mm	chr15	5875425 5875819 -	0	0	0
LTR	MaLR	MTA_Mm	chr15	5950127 5950522 -	0	0	0
SINE	Alu	B1_Mus2	chr15	6467087 6467233 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	6467234 6467277 +	0	0	0
LTR	MaLR	MTA_Mm	chr15	7003927 7004319 -	0	0	0
LTR	MaLR	MTA_Mm	chr15	7005413 7005805 -	0	0	0
SINE	Alu	B1_Mm	chr15	8054073 8054219 -	0	0	0
SINE	Alu	B1_Mus1	chr15	8398524 8398667 -	0	0	0
SINE	Alu	B1_Mus2	chr15	8538484 8538629 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	8552489 8552525 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr15	8667671 8667816 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	8667817 8667842 +	0	0	0
LTR	MaLR	MTD	chr15	9858443 9858761 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	9982582 9982638 +	0	0	0
SINE	Alu	B1_Mm	chr15	10044268 10044413 +	0	0	0
SINE	Alu	B1_Mus2	chr15	10375787 10375933 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	10375934 10375959 +	0	0	0

LTR	MaLR	MTA_Mm	chr15	10839967 10840361 +	0	0	0
LTR	MaLR	MTA_Mm	chr15	11497143 11497536 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	11960795 11960822 +	0	0	0
SINE	Alu	B1_Mus2	chr15	11993895 11994041 -	0	0	0
SINE	Alu	B1_Mus1	chr15	12132740 12132884 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	12132896 12132932 +	0	0	0
LTR	ERV1	RLTR23	chr10	66240038 66240386 +	0	0	0
LTR	MaLR	MTC	chr15	12611227 12611700 +	0	0	0
LTR	MaLR	MTA_Mm	chr15	12624811 12625205 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	14393360 14393405 +	0	0	0
LTR	MaLR	MTA_Mm	chr15	14439838 14440233 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	14484912 14484938 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr15	14532786 14532831 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	14694583 14694613 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	14708750 14708775 +	0	0	0
SINE	Alu	B1_Mus1	chr10	66398035 66398181 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	66398182 66398246 +	0	0	0
SINE	Alu	B1_Mm	chr10	66441809 66441961 +	0	0	0
LINE	L1	L1Md_T	chr15	15832021 15838443 -	0	0.380568	0
LTR	MaLR	MTA_Mm	chr15	15859483 15859876 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	16019378 16019398 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr15	16076084 16076223 -	0	0	0.342623
LTR	MaLR	MTA_Mm	chr15	16223362 16223756 -	0	0	0
SINE	Alu	B1_Mus2	chr15	17360675 17360820 +	0	0	0
LTR	MaLR	MTA_Mm	chr15	17408821 17409205 +	0	0	0
SINE	Alu	B1_Mus1	chr15	17572178 17572325 -	0	0	0.342623
LTR	MaLR	MTB	chr15	17782107 17782506 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	17880567 17880673 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	17976628 17976703 +	0	0	0
LINE	L1	L1Md_T	chr15	18594167 18600499 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	19194615 19194645 +	0	0	0
LINE	L1	L1Md_F2	chr15	19226885 19232918 -	0	0	0
LINE	L1	L1Md_F2	chr15	19488473 19493557 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	19863428 19863458 +	0	0	0
SINE	Alu	B1_Mus1	chr15	20052884 20053023 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	20053024 20053094 +	0	0	0
LTR	MaLR	MTB	chr15	20719902 20720299 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	20723717 20723784 +	0.591053	2.83525	0
LINE	L1	L1Md_F2	chr15	21839003 21844238 +	0	0	0
LINE	L1	L1Md_F2	chr15	22118695 22125281 -	0	0	0
SINE	Alu	B1_Mm	chr15	22322082 22322229 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	22322230 22322254 +	0	0	0
SINE	Alu	B1_Mus1	chr15	22692075 22692202 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	22762211 22762250 +	3.23032	2.43746	2.80392
SINE	Alu	B1_Mus2	chr15	22771370 22771515 -	0.591053	0	0
SINE	Alu	B1_Mus1	chr15	22929537 22929683 +	1.64268	1.85052	1.57327
Simple_repeat	Simple_repeat	(A)n	chr15	22929684 22929765 +	0.591053	1.85052	1.91589
Simple_repeat	Simple_repeat	(A)n	chr15	23708069 23708092 +	0	0	0
SINE	Alu	B1_Mus1	chr15	24976069 24976222 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	24976223 24976264 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	67111207 67111243 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	26218792 26218830 +	0	0	0
LTR	MaLR	MTA_Mm	chr15	26948751 26949146 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	26954149 26954183 +	0	0	0
LTR	MaLR	MTA_Mm	chr15	27141469 27141865 -	0	0	0
LTR	MaLR	MTA_Mm	chr15	27302761 27303155 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	67300918 67300955 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	28846888 28846973 +	0	0	0
SINE	Alu	B1_Mus2	chr15	28860535 28860683 -	0	0	0
SINE	Alu	B1_Mus2	chr15	31860530 31860676 -	0	0	0
LTR	MaLR	MTA_Mm	chr15	32005147 32005543 +	0	0	0
SINE	Alu	B1_Mus2	chr15	34367090 34367193 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	34367194 34367224 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	34609414 34609448 +	1.77316	0.708814	0.685246
Simple_repeat	Simple_repeat	(A)n	chr15	34690107 34690178 +	0	0	0
LINE	L1	L1_Mus2	chr15	35899137 35904590 -	0.591053	0	0
SINE	Alu	B1_Mus2	chr15	36567894 36568045 -	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr15	36658916 36658938 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	36764070 36764130 +	0	0	0
LTR	MaLR	MTA_Mm	chr15	36789268 36789664 -	0	0	0
LTR	MaLR	MTA_Mm	chr15	36790368 36790764 -	0	0	0
SINE	Alu	B1_Mm	chr15	37057208 37057290 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	38105225 38105252 +	0	0	0
SINE	Alu	B1_Mus2	chr15	38124395 38124540 -	0	0	0
LTR	MaLR	MTA_Mm	chr15	38276168 38276561 +	0	0	0
SINE	Alu	B1_Mur4	chr15	38720895 38721041 +	0.591053	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr15	38721042 38721063 +	0.591053	1.08938	0
SINE	Alu	B1_Mus1	chr15	38723785 38723925 +	0	1.08938	0
Simple_repeat	Simple_repeat	(CAAA)n	chr15	38723926 38723977 +	0	1.08938	0.615324
SINE	Alu	B1_Mus1	chr15	39596807 39596952 -	0	0	0
scRNA	scRNA	4.5SRNA	chr15	39596953 39597044 -	0	0	0
SINE	B2	B2_Mm2	chr10	68503600 68503788 -	0	0	0
LTR	MaLR	MTA_Mm	chr15	40086067 40086460 -	0	0	0
LTR	MaLR	MTA_Mm	chr15	40114188 40114572 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	40360695 40360745 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	68561307 68561347 +	0	0	0
SINE	Alu	B1_Mus1	chr15	40460915 40461055 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	40461056 40461088 +	0	0	0
LTR	MaLR	MTA_Mm	chr15	40974871 40975266 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	68648895 68648930 +	0	0	0
LTR	MaLR	MTA_Mm	chr15	43209044 43209437 -	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	68786713 68786748 +	0	0	0
SINE	Alu	B1_Mus1	chr15	43703792 43703937 -	0	0	0
LTR	MaLR	MTA_Mm	chr15	43769725 43770119 -	0	0	0
LTR	MaLR	MTA_Mm	chr15	43873844 43874239 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	68866747 68866786 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	45222530 45222586 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	45232684 45232716 +	0	0	0
LINE	L1	L1Md_T	chr15	45415068 45422145 -	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chr15	45456999 45457160 +	0	0	0
SINE	Alu	B1_Mm	chr15	46154516 46154662 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	46338493 46338529 +	0	0	0
SINE	Alu	B1_Mus2	chr15	46548015 46548157 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	46917830 46917865 +	0	0	0
SINE	Alu	B1_Mus2	chr15	48883032 48883176 -	0	0	0
SINE	Alu	B1_Mus1	chr15	48889679 48889825 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	49483091 49483123 +	0	0	0
LTR	MaLR	MTA_Mm	chr15	49752109 49752505 -	0	0	0
LTR	MaLR	MTA_Mm	chr15	50109869 50110265 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	50425071 50425100 +	0	0	0
LTR	MaLR	MTA_Mm	chr15	52040744 52041138 -	0	0	0
SINE	Alu	B1_Mus1	chr15	52041206 52041335 +	0	0	0
LINE	L1	L1Md_T	chr15	52562376 52568805 -	0.591053	0	0
LINE	L1	L1Md_T	chr15	54036273 54042835 -	0	0	0
SINE	Alu	B1_Mus1	chr15	54234439 54234570 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	55121571 55121650 +	0	0	0.342623
SINE	Alu	B1_Mm	chr10	69685115 69685256 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	69685257 69685279 +	0	0	0
SINE	B2	B2_Mm1t	chr15	56545699 56545891 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	56545892 56545916 +	0.664388	0	0
SINE	Alu	B1_Mus1	chr15	56946073 56946219 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	57520881 57520932 +	0	0	0
SINE	Alu	B1_Mus1	chr15	57803702 57803847 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	69901714 69901749 +	0	0	0
SINE	Alu	B1_Mus1	chr15	58933180 58933329 +	0	0	0
Simple_repeat	Simple_repeat	(CAAA)n	chr15	58933331 58933384 +	0	0	0
SINE	Alu	B1_Mus1	chr15	59434692 59434839 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	59492871 59492895 +	0	0	0
SINE	Alu	B1_Mus1	chr15	60073931 60074077 -	0	0	0
SINE	Alu	B1_Mus1	chr15	60390706 60390841 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	60390842 60390868 +	0	0	0
LTR	MaLR	MTA_Mm	chr15	60490096 60490491 +	0	0	0
LINE	L1	L1Md_F2	chr15	60918898 60924740 -	0	0	0.866667
SINE	Alu	B1_Mm	chr15	61070114 61070245 +	0	0	0

SINE	Alu	B1_Mm	chr15	61474873 61475019 -	0	0	0
SINE	Alu	B1_Mus1	chr15	61482345 61482491 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	61482492 61482527 +	0	0	0
SINE	Alu	B1_Mm	chr15	61678052 61678183 -	0	0	0
SINE	B2	B2_Mm2	chr15	62084004 62084194 -	0	0	0
LINE	L1	L1Md_T	chr15	62518380 62525329 -	0	0	0
LINE	L1	L1Md_T	chr15	62595637 62601999 -	0	0	0
SINE	Alu	B1_Mus1	chr15	63224714 63224860 -	0	0	0
LTR	MaLR	MTB	chr15	63899778 63900148 -	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	64272356 64272396 +	0	0.380568	0
LINE	L1	L1_Mus1	chr15	64434059 64440297 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	66807917 66807950 +	0	0	0
SINE	Alu	B1_Mm	chr15	67791574 67791720 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	67791721 67791755 +	0	0	0
LTR	MaLR	MTA_Mm	chr15	67832761 67833162 -	0	0	0
LTR	MaLR	MTA_Mm	chr15	68535637 68536031 +	0	0	0
SINE	Alu	B1_Mus2	chr15	69508447 69508581 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	69508582 69508610 +	0	0	0
LTR	MaLR	ORR1D2	chr15	70085287 70085634 -	0	0	0
SINE	Alu	B1_Mus2	chr15	70210401 70210547 -	0	0	0
LTR	MaLR	MTA_Mm	chr15	70279789 70280183 -	0	0	0
LTR	MaLR	MTA_Mm	chr15	70720534 70720928 +	0	0	0
LINE	L1	L1_Mus3	chr15	70902819 70908431 +	0	0	0
SINE	Alu	B1_Mus2	chr15	71901244 71901390 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	72320428 72320463 +	0	0	0
SINE	Alu	B1_Mus1	chr15	73775312 73775458 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	73775459 73775483 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	74009693 74009741 +	0	0	0
LTR	MaLR	MTA_Mm	chr10	71331831 71332227 -	0	0	0
SINE	Alu	B1F	chr15	74704995 74705120 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	74705121 74705145 +	0.591053	0	0
LINE	L1	L1Md_F	chr15	74930032 74934585 -	0	0	0
LTR	MaLR	MTA_Mm	chr15	75085896 75086297 +	0	0	0
SINE	B2	B2_Mm1a	chr10	71447435 71447625 -	0	0	0
SINE	B2	B2_Mm1a	chr15	75704296 75704483 -	0.332194	0	0
SINE	B2	B2_Mm1t	chr15	75705155 75705287 -	0	0	0
SINE	Alu	B1_Mus1	chr15	75705307 75705447 -	0	0	0
SINE	Alu	B1_Mm	chr15	76172913 76173047 -	0	0	0
SINE	Alu	B1_Mm	chr15	76361352 76361498 +	0	0.708814	0.342623
Simple_repeat	Simple_repeat	(A)n	chr15	76361499 76361540 +	0	0.708814	0.342623
SINE	Alu	B1_Mm	chr15	76660238 76660351 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	76660352 76660395 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	77153214 77153260 +	0	0	0
SINE	B2	B2_Mm2	chr15	77165556 77165743 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	77165744 77165770 +	0	0	0
SINE	Alu	B1_Mm	chr15	77507257 77507402 -	0	0	0
SINE	Alu	B1_Mus2	chr15	77693991 77694137 +	0	0	0
SINE	Alu	B1_Mus1	chr15	77697941 77698087 -	0	0	0
LTR	MaLR	MTA_Mm	chr15	77700938 77701304 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	78646632 78646661 +	0	0	0
SINE	Alu	B1_Mus2	chr15	78751588 78751734 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	78893721 78893753 +	0	0	0
SINE	B4	B4A	chr15	79159325 79159535 -	0	0	0
SINE	Alu	B1_Mus2	chr15	79246418 79246564 -	0	0	0
SINE	Alu	B1_Mus2	chr15	79286518 79286663 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	79286664 79286689 +	0	0	0.342623
SINE	Alu	B1_Mm	chr15	79385452 79385597 -	0	0	0
SINE	Alu	B1_Mus2	chr15	79386946 79387087 -	0	0	0
SINE	Alu	B1_Mus1	chr10	71932560 71932702 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	71932703 71932797 +	0	0	0
LTR	MaLR	MTA_Mm	chr10	72017834 72018228 -	0	0	0
SINE	Alu	B1_Mus2	chr15	80439658 80439794 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	80446059 80446085 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr10	72084495 72084564 +	0	0.708814	0
SINE	Alu	B1_Mus2	chr10	72137753 72137899 +	6.86594	7.60204	14.3299
Simple_repeat	Simple_repeat	(A)n	chr10	72137900 72137922 +	1.77316	2.12644	2.18859
LTR	MaLR	MTB	chr10	72188701 72189085 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr10	72244293 72244325 +	0	0	0
SINE	Alu	B1_Mus2	chr15	81658572 81658718 +	0	2.17876	0.685246
Simple_repeat	Simple_repeat	(A)n	chr15	81794624 81794649 +	0	0	0
SINE	B2	B2_Mm1a	chr15	81973809 81974002 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	82392497 82392525 +	0	0	0
SINE	Alu	B1_Mm	chr15	82466231 82466377 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	82466378 82466408 +	0	0	0
SINE	Alu	B1_Mus1	chr15	83014802 83014939 -	0	0	0
SINE	Alu	B1_Mus2	chr15	83428734 83428880 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	83428881 83428911 +	0	0	0
SINE	Alu	B1_Mus1	chr10	72740036 72740182 +	1.5143	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	72740183 72740208 +	1.5143	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	72744935 72744970 +	1.05163	2.7378	0.615324
Simple_repeat	Simple_repeat	(A)n	chr15	85346790 85346819 +	0	0	0
SINE	B4	B4	chr15	85435549 85435802 -	0.591053	0	0.342623
SINE	Alu	B1_Mm	chr15	86022722 86022868 -	0	0	0
LTR	MaLR	MTA_Mm	chr15	86605478 86605872 -	0	0	0
SINE	Alu	B1_Mus1	chr15	87358153 87358299 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	87358300 87358330 +	0	0	0
SINE	Alu	B1_Mur2	chr15	88454134 88454281 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	88454282 88454307 +	0	0	0
SINE	Alu	B1_Mur3	chr15	89041325 89041465 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	89041466 89041488 +	0	0	0
SINE	Alu	B1_Mus2	chr15	89214462 89214608 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	89295381 89295458 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr15	89500371 89500437 +	0	0	0
SINE	Alu	B1_Mus2	chr15	89505794 89505940 -	0	0	0
LTR	MaLR	MTA_Mm	chr15	89751116 89751512 -	0	0	0
LTR	MaLR	MTB	chr15	90569600 90569992 +	0	0	0
LTR	MaLR	MTA_Mm	chr15	90589570 90589962 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	91487022 91487069 +	0	0.380568	0.615324
Simple_repeat	Simple_repeat	(A)n	chr15	95447177 95447211 +	0	0	0
SINE	Alu	B1_Mus2	chr15	96238596 96238742 -	0	0	0
SINE	Alu	B1_Mur4	chr15	96505799 96505947 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	96505948 96505970 +	0	0	0
Simple_repeat	Simple_repeat	(CAA)n	chr15	96505969 96505998 +	0	0	0
SINE	Alu	B1_Mus2	chr15	96549722 96549868 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	96549869 96549921 +	0	0	0
SINE	Alu	B1_Mus2	chr15	96912031 96912176 -	0	0	0
LTR	MaLR	MTB	chr15	97440885 97441268 -	0	0	0
SINE	B2	B2_Mm1t	chr15	98014059 98014243 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	98089719 98089753 +	0	0	0
SINE	Alu	B1_Mm	chr15	98135001 98135146 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	98135147 98135213 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	98161455 98161498 +	0	0	0
SINE	Alu	PB1D7	chr15	98286021 98286148 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	98475680 98475719 +	0	0	0
SINE	Alu	B1_Mus2	chr15	98855711 98855855 -	0	0	0
SINE	Alu	B1_Mus2	chr15	98876443 98876579 -	0	0	0
SINE	Alu	B1_Mus2	chr15	98896462 98896608 -	0	0	0
SINE	Alu	B1_Mm	chr15	98898166 98898315 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	98898316 98898338 +	0	0	0
SINE	B2	B2_Mm2	chr15	99100231 99100418 -	0	0	0
LTR	MaLR	MTA_Mm	chr15	99387593 99387989 +	0	0	0
SINE	Alu	B1_Mus1	chr15	99782723 99782870 -	0	0	0
SINE	Alu	B1_Mus1	chr15	99862846 99862992 -	0	0	0
SINE	Alu	B1_Mm	chr15	10026898 10026912 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	10026912 10026915 +	0	0	0
SINE	Alu	B1_Mm	chr15	10056549 10056564 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr15	10056564 10056568 +	0	0	0
SINE	Alu	B1_Mus1	chr10	74672793 74672939 -	0	0	0
SINE	Alu	B1_Mur4	chr15	10100171 10100183 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr15	10100183 10100189 +	0	0	0
SINE	Alu	B1_Mm	chr15	10212170 10212185 +	0	0	0
Simple_repeat	Simple_repeat	(CAAAA)n	chr15	10212186 10212190 +	0	0	0
SINE	B2	B2_Mm1a	chr15	10222182 10222201 +	60.8852	34.5905	36.4764
SINE	Alu	B1_Mm	chr15	10247784 10247799 -	0	0	0



SINE	Alu	B1_Mm	chr15	10265363	10265375	-	0	0	0
SINE	Alu	B1_Mm	chr10	74968654	74968800	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	74968801	74968836	+	0	0	0
SINE	Alu	B1_Mus1	chr16	4014802	4014951	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	4235194	4235217	+	0	0	0
SINE	Alu	B1_Mus1	chr16	4448644	4448780	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	4560096	4560130	+	0	0	0.342623
SINE	Alu	B1_Mm	chr16	4568679	4568825	-	0	0	0
SINE	Alu	B1_Mus2	chr16	4581159	4581304	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	4581305	4581356	+	0	0	0
SINE	B2	B3	chr16	4870505	4870705	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	8470347	8470381	+	0	0	0
SINE	Alu	B1_Mus1	chr16	8779667	8779814	-	0	0	0
SINE	Alu	B1_Mus2	chr16	8794616	8794761	-	0	0	0
SINE	B2	B2_Mm1a	chr16	8924187	8924394	+	0	0	0
SINE	Alu	PB1D9	chr10	75494233	75494352	+	0	0	0
SINE	Alu	B1_Mus1	chr16	10054680	10054826	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	10816042	10816069	+	0	0	0
SINE	Alu	B1_Mus1	chr16	10903764	10903910	-	0	0	0
LTR	MaLR	ORR1A1	chr16	10949002	10949325	+	0	0	0
SINE	Alu	B1_Mus2	chr16	11955939	11956084	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	11956085	11956107	+	0	0	0
LTR	MaLR	MTA_Mm	chr16	12388043	12388439	+	0	0	0
LTR	MaLR	MTA_Mm	chr16	12969444	12969838	-	0	0	0
SINE	Alu	B1_Mus1	chr16	13043111	13043257	-	0	0	0
SINE	B2	B2_Mm1a	chr16	13056655	13056840	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	13435821	13435857	+	0	0	0
SINE	Alu	B1_Mm	chr16	13509001	13509146	-	0.332194	0	0
LTR	MaLR	MTA_Mm	chr16	14333143	14333536	+	0	0	0
LTR	MaLR	MTA_Mm	chr16	14534685	14535080	+	0	0	0
LTR	MaLR	MTA_Mm	chr16	15024876	15025255	-	0	0	0
SINE	Alu	B1_Mus2	chr16	15474966	15475111	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	15475112	15475140	+	0	0	0
LTR	MaLR	MTA_Mm	chr10	76125170	76125563	+	0	0	0
SINE	Alu	B1_Mus1	chr16	15606704	15606849	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	15606850	15606875	+	0	0	0
SINE	Alu	B1_Mus2	chr16	16856149	16856278	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	16856279	16856303	+	0	0	0
LTR	ERVK	RMER13A	chr16	16856304	16857114	+	0	0	0
SINE	B2	B2_Mm1a	chr16	17231872	17232064	-	0	0	0
SINE	Alu	PB1D9	chr16	17266473	17266590	-	0	0	0
scRNA	scRNA	BC1_Mm	chr16	17724447	17724579	+	0	0	0
SINE	Alu	B1_Mus2	chr10	76450934	76451075	-	0	0	0
SINE	Alu	B1_Mus2	chr16	18675318	18675463	-	0	0	0.615324
SINE	Alu	B1_Mus1	chr16	18728464	18728595	-	0	0	0
SINE	Alu	B1_Mus2	chr16	18732273	18732417	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	19127181	19127272	+	0	0.380568	0
SINE	Alu	B1_Mus1	chr10	76632881	76633027	-	0	0	0
LTR	MaLR	MTA_Mm	chr16	19875620	19876019	-	0	0	0
SINE	Alu	B1_Mus1	chr16	20068788	20068935	-	0	0	0
SINE	Alu	B1_Mus1	chr16	20138532	20138679	+	0	0	0
SINE	Alu	B1_Mus1	chr16	20437459	20437573	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	20437574	20437596	+	0.332194	0	0
SINE	Alu	B1_Mus2	chr16	20495584	20495734	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	20495735	20495769	+	0	0	0
SINE	Alu	B1_Mus1	chr16	20669264	20669411	-	0	0.380568	0
SINE	Alu	B1_Mus1	chr16	20671094	20671225	-	0	0	0
SINE	Alu	B1_Mm	chr16	21178963	21179109	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	21369020	21369057	+	0	0	0
SINE	Alu	B1_Mus2	chr16	21697498	21697644	-	0	0	0
LTR	MaLR	MTA_Mm	chr16	21725778	21726165	+	0	0	0
SINE	Alu	B1_Mus2	chr10	76885020	76885164	+	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr10	76885165	76885214	+	0	0	0
SINE	Alu	B1_Mm	chr16	22199752	22199903	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	22199904	22199926	+	0	0	0
Simple_repeat	Simple_repeat	(CAA)n	chr16	22199925	22199948	+	0	0	0
SINE	Alu	B1_Mus2	chr16	22215337	22215482	-	0	0	0

SINE	Alu	B1_Mus2	chr16	22238762 22238908 -	0	0	0
SINE	Alu	B1_Mm	chr16	22350954 22351102 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	22351103 22351126 +	0	0	0
SINE	Alu	B1_Mm	chr16	22354254 22354400 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	76989359 76989403 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr16	22758589 22758636 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	77038004 77038059 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	23200059 23200098 +	0	0	0
LTR	MaLR	MTA_Mm	chr16	23694555 23694952 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	24136768 24136797 +	0	0	0
LTR	MaLR	MTA_Mm	chr10	77157696 77158094 -	0	0	0
SINE	Alu	B1_Mus2	chr10	77172550 77172696 -	0	0	0
SINE	Alu	B1_Mm	chr16	25018679 25018825 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr16	25018826 25018857 +	0	0	0.342623
SINE	Alu	B1_Mus1	chr16	25022133 25022279 -	0	0	0
SINE	Alu	B1_Mm	chr10	77209614 77209766 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	77209767 77209793 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	26814956 26814988 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	27099973 27100001 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	27498761 27498789 +	0	0	0
LINE	L1	L1Md_F2	chr16	27691664 27696693 +	0	0.708814	0
SINE	Alu	B1_Mm	chr10	77474328 77474474 +	0	0	0.342623
Simple_repeat	Simple_repeat	(GAAAA)n	chr10	77474475 77474528 +	0.591053	0	0.342623
LTR	MaLR	MTA_Mm	chr16	27999488 27999881 +	0	0	0
LTR	MaLR	MTA_Mm	chr16	28147218 28147601 +	0	0	0
LTR	MaLR	MTA_Mm	chr16	28781778 28782173 +	0	0	0
LTR	MaLR	MTA_Mm	chr16	28783268 28783663 +	0.332194	0	0
SINE	Alu	B1_Mus2	chr16	29574703 29574862 -	0	0	0
SINE	B2	B2_Mm1a	chr16	29765099 29765291 -	0	1.01984	0
SINE	Alu	B1_Mm	chr16	29949683 29949822 +	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr16	29949823 29949846 +	0	0.708814	0
SINE	Alu	B1_Mm	chr16	30117953 30118104 -	0	0	0
SINE	Alu	B1_Mus2	chr16	30167029 30167175 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	30185999 30186033 +	0	0	0
LTR	MaLR	MTA_Mm	chr16	30199502 30199897 +	0	0	0
LTR	MaLR	MTA_Mm	chr16	30685123 30685494 +	0	0	0
SINE	Alu	B1_Mus2	chr16	30767442 30767588 -	0	0	0
SINE	Alu	B1_Mus2	chr16	30874959 30875102 -	0	0	0
SINE	Alu	B1_Mm	chr10	77825532 77825670 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	77825671 77825690 +	0.332194	0	0
SINE	Alu	B1_Mus2	chr16	31524608 31524741 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	31524742 31524768 +	0	0	0
LTR	MaLR	MTC	chr16	31525303 31525651 -	0	0	0
SINE	Alu	B1_Mus2	chr16	31525652 31525798 -	0	0	0
SINE	Alu	B1_Mm	chr16	31625368 31625514 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	77856501 77856534 +	36.1014	24.1903	20.2875
SINE	B2	B2_Mm1a	chr16	32207136 32207324 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	32207325 32207348 +	0.591053	1.08938	0.615324
SINE	Alu	B1_Mus2	chr16	32260522 32260667 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	32260676 32260728 +	0	0	0
LTR	MaLR	MTA_Mm	chr16	32596055 32596450 +	0	0	0
LTR	MaLR	MTA_Mm	chr16	32597542 32597937 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	32687873 32687904 +	0	0	0
SINE	B2	B3	chr16	32687905 32688086 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	77994813 77994833 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	33142032 33142096 +	0	0	0
LTR	MaLR	MTA_Mm	chr16	33156072 33156463 -	0	0	0
SINE	Alu	B1_Mm	chr16	33166622 33166766 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	33166767 33166793 +	0	0	0
SINE	Alu	B1_Mus2	chr16	33348726 33348872 -	0	0.708814	0
LINE	L1	L1Md_F2	chr10	78099785 78106008 -	0	0	0
SINE	Alu	B1_Mm	chr16	33825027 33825173 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	33825174 33825196 +	0	0	0
LTR	MaLR	MTA_Mm	chr16	34626440 34626836 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	34641967 34642025 +	0	0	0
Simple_repeat	Simple_repeat	(T)n	chr16	36096166 36096185 +	0	0	0
SINE	Alu	B1_Mm	chr16	36114423 36114569 -	0	0	0

SINE	Alu	B1_Mus1	chr16	36192594 36192740 -	0	0	0
LINE	L1	L1VL4	chr16	36574461 36577650 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	37635667 37635706 +	0	0	0
SINE	Alu	B1_Mus2	chr16	38296354 38296498 +	0	0	0
SINE	Alu	B1_Mus2	chr16	38436732 38436880 -	0	0	0
SINE	Alu	B1_Mm	chr16	38557824 38557945 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	38557946 38557969 +	0	0	0
LINE	L1	L1VL4	chr16	38734842 38739732 -	0	0	0
SINE	Alu	B1_Mus1	chr16	39033036 39033179 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	39882022 39882085 +	0	0	0
SINE	Alu	B1_Mm	chr16	40461025 40461171 +	7.75265	6.0891	5.88752
Simple_repeat	Simple_repeat	(A)n	chr16	40461172 40461209 +	7.32516	7.17848	5.68474
LINE	L1	L1Md_T	chr16	40660159 40666852 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	40765710 40765750 +	2.34225	0.380568	2.87384
SINE	Alu	B1_Mus1	chr16	40779703 40779849 +	1.64268	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr16	40779850 40779898 +	1.97487	0.380568	0
SINE	Alu	B1_Mm	chr16	40788581 40788731 +	1.84649	0.380568	0.615324
Simple_repeat	Simple_repeat	(A)n	chr16	40788732 40788757 +	1.5143	1.08938	0.957947
LTR	MaLR	MTA_Mm	chr16	40950225 40950623 +	1.18211	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr16	40980920 40980958 +	5.85339	0	1.57327
Simple_repeat	Simple_repeat	(A)n	chr16	41136291 41136331 +	3.98493	3.52684	0.615324
Simple_repeat	Simple_repeat	(A)n	chr16	41253870 41253902 +	4.20861	2.12644	7.18586
Simple_repeat	Simple_repeat	(A)n	chr16	42235820 42235848 +	0	0	0
SINE	Alu	B1_Mm	chr16	44345287 44345433 -	0	0	0
LTR	MaLR	MTA_Mm	chr16	45006088 45006469 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	45046752 45046781 +	0	0	0
SINE	Alu	PB1D9	chr16	45710508 45710627 -	1.18211	0.708814	1.23065
SINE	Alu	B1_Mus2	chr10	10831361 10831505 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	46280706 46280754 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	46514674 46514720 +	0	0	0
LTR	MaLR	MTA_Mm	chr16	47099149 47099544 +	0	0	0
LTR	MaLR	MTA_Mm	chr16	47100637 47101032 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	47102003 47102031 +	0	0	0
LTR	MaLR	MTA_Mm	chr16	47270600 47270995 +	0	0	0
SINE	Alu	B1_Mus2	chr16	47782367 47782502 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	47782503 47782525 +	0	0	0
SINE	Alu	B1_Mus2	chr16	48058707 48058855 -	0	0	0
SINE	Alu	B1_Mus1	chr16	48181940 48182093 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	48304219 48304254 +	0	0	0
LTR	MaLR	MTA_Mm	chr16	48391230 48391625 -	0	0	0
SINE	Alu	B1_Mus1	chr16	48665677 48665824 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	48665825 48665868 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	48816786 48816821 +	0	0	0
SINE	Alu	B1_Mus2	chr16	48822778 48822924 -	0.923247	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr16	48862503 48862547 +	0	0	0
SINE	Alu	B1_Mm	chr16	48900963 48901109 +	0	0	0
SINE	Alu	B1_Mus2	chr10	79626268 79626417 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	79626418 79626446 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	49227237 49227259 +	0	0	0
LTR	MaLR	MTA_Mm	chr16	49323930 49324308 -	0	0	0
SINE	B2	B2_Mm1a	chr10	79674245 79674436 +	0	0	0
SINE	Alu	B1_Mus1	chr10	79783710 79783851 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	79783852 79783874 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	51207955 51207997 +	0.332194	0	0
LINE	L1	L1Md_F2	chr16	51235920 51241841 -	0	0	0
SINE	Alu	B1_Mm	chr16	51588031 51588172 -	0	0	0
LTR	MaLR	MTA_Mm	chr16	51834667 51835062 +	0.332194	0	0
SINE	Alu	B1_Mm	chr10	79906684 79906830 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	53153637 53153665 +	6.22949	8.29006	14.1657
SINE	B2	B2_Mm1t	chr16	53271536 53271727 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr16	53271728 53271810 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	53601351 53601439 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	53858837 53858878 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr16	53913303 53913329 +	0	0	0
SINE	Alu	B1_Mm	chr16	53929746 53929898 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	54108342 54108383 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	54724747 54724782 +	0	0	0

LTR	MaLR	MTA_Mm	chr16	54935952 54936267 +	0	0	0
LTR	MaLR	MTA_Mm	chr16	55217169 55217569 +	0	0	0
SINE	Alu	B1_Mus2	chr10	80045747 80045883 +	0	0	0
SINE	Alu	B1_Mus1	chr16	55757854 55758013 -	0	0	0
SINE	Alu	B1_Mus2	chr16	55899587 55899738 -	0	0	0
SINE	Alu	PB1D9	chr16	56057240 56057312 +	0	0	0
SINE	B2	B2_Mm1a	chr10	80124687 80124879 +	0	0	0
LINE	L1	L1_Mus2	chr16	56959159 56965602 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	57890992 57891019 +	0	0	0
SINE	Alu	B1_Mus1	chr16	59092679 59092830 +	0	0	0
SINE	Alu	B1_Mus1	chr10	80387272 80387402 -	0	0	0
SINE	Alu	B1_Mm	chr16	60965637 60965783 -	0	0	0
SINE	Alu	B1_Mus2	chr16	61059989 61060137 -	0	0	0
LINE	L1	L1Md_F2	chr16	61304654 61310539 -	0	0	0
SINE	Alu	B1_Mus1	chr16	61398582 61398726 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	61487811 61487861 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	61967774 61967802 +	0	0	0
LTR	MaLR	MTA_Mm	chr16	62930284 62930677 -	0	0	0
LTR	MaLR	MTB_Mm	chr16	63202862 63203284 +	0	0	0
LINE	L1	L1Md_T	chr16	63257508 63263145 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	63903888 63903938 +	0	0	0
SINE	Alu	B1_Mus1	chr10	80638365 80638506 -	0	0	0
LTR	MaLR	MTA_Mm	chr16	64283878 64284273 +	0	0	0
SINE	Alu	B1_Mus2	chr16	64846771 64846917 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	64846918 64846942 +	0	0	0
SINE	Alu	B1F2	chr10	80719791 80719921 -	0	0	0
LINE	L1	L1Md_T	chr16	65745404 65751850 -	0	0	0
SINE	Alu	B1_Mus2	chr16	66045842 66045992 -	0	0	0
SINE	Alu	B1_Mur2	chr16	66046004 66046142 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	66229406 66229445 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	66280129 66280154 +	0	0	0
LINE	L1	L1Md_T	chr16	66518330 66523846 -	0.332194	1.08938	0.342623
SINE	B2	B2_Mm1a	chr16	66523847 66524033 -	0.332194	0.380568	0.342623
Simple_repeat	Simple_repeat	(A)n	chr16	67629545 67629581 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	67652636 67652698 +	0	0	0
SINE	Alu	B1_Mus1	chr16	68173995 68174140 -	0	0	0
SINE	Alu	B1_Mus2	chr16	68374729 68374874 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	68510223 68510267 +	0	0	0
LINE	L1	L1Md_F3	chr16	68669512 68675713 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	68777967 68778000 +	0	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr16	68957962 68958009 +	0	0	0
LTR	MaLR	MTA_Mm	chr16	69315600 69315995 -	0.591053	0.708814	0
SINE	Alu	B1_Mus1	chr16	69360343 69360461 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	69360462 69360520 +	0	0.380568	0.615324
Simple_repeat	Simple_repeat	(A)n	chr16	69548646 69548673 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	69600051 69600090 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	70019154 70019208 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	70165944 70165973 +	0	0	0
SINE	Alu	B1_Mus1	chr10	81153037 81153184 -	0	0	0
LINE	L1	L1Md_A	chr16	71147905 71154513 -	0	0	0
LTR	MaLR	MTA_Mm	chr10	81187355 81187750 -	0	0.380568	0
SINE	Alu	B1_Mm	chr10	81241637 81241786 +	0	0	0
LINE	L1	L1Md_F2	chr16	72012597 72018594 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	81241787 81241822 +	0	0	0
SINE	Alu	B1_Mm	chr16	72114204 72114349 -	0	0	0
SINE	Alu	B1_Mus1	chr10	81312132 81312279 -	0	0	0
SINE	Alu	B1_Mus2	chr16	73255135 73255283 -	0	0	0
LTR	MaLR	MTA_Mm	chr10	81346447 81346842 -	0	0.380568	0
SINE	Alu	B1_Mm	chr16	73701525 73701670 +	0	0	0
SINE	Alu	B1_Mm	chr10	81400978 81401127 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	81401128 81401169 +	0	0	0
SINE	Alu	B1_Mm	chr10	81464491 81464636 -	0	0	0
LTR	MaLR	MTB	chr16	75000744 75001127 +	0	0	0
LTR	MaLR	MTA_Mm	chr10	81479976 81480370 -	0	0	0.615324
SINE	Alu	B1_Mus2	chr16	75223637 75223782 -	1.84649	1.46995	1.23065
Simple_repeat	Simple_repeat	(A)n	chr16	75272037 75272075 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	76054821 76054857 +	0	0	0

SINE	Alu	B1_Mus2	chr16	76278117 76278263 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	76278264 76278303 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	76692842 76692871 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr16	78197632 78197674 +	0	0	0
SINE	Alu	B1_Mus2	chr10	81743628 81743774 -	0	0	0
LTR	MaLR	MTA_Mm	chr16	78517945 78518342 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	78651066 78651103 +	0	0	0
SINE	Alu	B1_Mus1	chr16	78749964 78750111 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	79494828 79494875 +	0	0	0
LINE	L1	L1Md_F2	chr16	79510837 79516583 -	0	0	0
LTR	MaLR	MTA_Mm	chr16	79529312 79529716 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	79741577 79741627 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	80895679 80895704 +	0	0	0
LINE	L1	L1Md_T	chr16	81029627 81036581 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	81180253 81180294 +	0	0	0
SINE	Alu	B1_Mm	chr16	82101837 82101979 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	82101980 82102010 +	0	0	0
SINE	Alu	B1_Mm	chr16	83424062 83424208 -	0	0	0
LTR	MaLR	MTA_Mm	chr16	83493148 83493542 +	0	0	0
SINE	Alu	B1_Mus1	chr16	83657045 83657197 -	0	0	0
SINE	Alu	B1_Mm	chr16	83661650 83661796 -	0	0	0
LTR	MaLR	MTA_Mm	chr16	84085084 84085479 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	84529775 84529801 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr10	82293547 82293582 +	0.591053	0	0
LTR	MaLR	MTA_Mm	chr16	84764878 84765271 +	0	0	0
SINE	B2	B2_Mm1a	chr16	84882440 84882631 -	0	0	0.342623
SINE	Alu	B1_Mus1	chr16	85214137 85214284 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr16	85214297 85214404 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	85804206 85804233 +	0	0	0
LTR	MaLR	MTA_Mm	chr16	85829364 85829758 -	0	0	0
SINE	Alu	B1_Mus2	chr16	86157329 86157475 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	86157476 86157498 +	0	0	0
LINE	L1	L1_Mus4	chr10	11237726 11243407 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	87213890 87213919 +	0	0	0
SINE	Alu	B1_Mm	chr16	87674561 87674707 -	0	0	0
SINE	Alu	B1_Mus2	chr16	87687401 87687545 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	87687546 87687572 +	0	0	0
LINE	L1	L1Md_F2	chr16	88658617 88665017 -	0	0	0
LTR	MaLR	MTA_Mm	chr16	89232428 89232824 -	0	0	0
LTR	MaLR	MTA_Mm	chr16	89453924 89454318 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	89542494 89542538 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	89675110 89675136 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	90504652 90504685 +	0.591053	0	0
LTR	MaLR	MTA_Mm	chr16	90668054 90668460 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr16	91097405 91097440 +	0	0	0
SINE	Alu	B1_Mus2	chr10	82801210 82801356 -	0	0	0
SINE	B2	B3	chr16	91311376 91311559 -	0	0	0
SINE	Alu	B1_Mus2	chr10	82818299 82818444 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	82818445 82818481 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	91978355 91978391 +	0	0	0
SINE	Alu	B1_Mus1	chr16	92572276 92572405 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	11330238 11330281 +	1.58764	0.380568	1.84597
LINE	L1	L1Md_F3	chr16	93617191 93622227 +	0.923247	1.41763	0
LTR	MaLR	MTA_Mm	chr16	93673360 93673755 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	94512622 94512656 +	0	0	0
LTR	MaLR	MTA_Mm	chr16	94573110 94573504 +	0	0	0
SINE	Alu	B1_Mm	chr16	95830797 95830937 -	0	0	0
SINE	Alu	B1_Mus1	chr16	96031795 96031942 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	96031943 96031977 +	0	0	0
SINE	Alu	B1_Mus2	chr16	96172914 96173058 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	96173059 96173084 +	0	0	0
SINE	Alu	B1_Mus2	chr16	96324471 96324617 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	97422049 97422089 +	0	0	0
SINE	Alu	B1_Mus1	chr16	97484485 97484632 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	97484633 97484660 +	0	0	0.342623
SINE	Alu	B1_Mus1	chr16	98274037 98274177 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr16	98274178 98274199 +	0	0	0

SINE	Alu	B1_Mus2	chr17	3089203	3089349	-	0	0	0
LTR	MaLR	MTA_Mm	chr10	83520503	83520897	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	4106198	4106234	+	0	0	0
SINE	Alu	B1_Mus2	chr17	4709991	4710136	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	4710137	4710186	+	0	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr17	4962042	4962078	+	0	0	0
SINE	Alu	B1_Mm	chr17	4983240	4983380	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	4983381	4983421	+	0	0	0
SINE	Alu	B1_Mus2	chr17	6354163	6354310	-	0.664388	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr17	6461288	6461326	+	0	0	0
SINE	Alu	B1_Mm	chr17	6494237	6494383	-	0	0	0
SINE	Alu	B1_Mus1	chr17	6539963	6540111	-	0	0	0
SINE	Alu	B1_Mus1	chr17	7049637	7049784	-	0	0	0
SINE	Alu	B1_Mus2	chr17	7640719	7640864	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	7640865	7640887	+	0	0	0
SINE	Alu	B1_Mus2	chr17	7721554	7721699	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	7721700	7721740	+	0	0	0
SINE	Alu	B1_Mus1	chr17	7884446	7884592	+	0	0	0
SINE	Alu	B1_Mus1	chr17	8351392	8351538	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	8351539	8351569	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	8944414	8944450	+	0	0	0
SINE	Alu	B1_Mm	chr17	10552820	10552966	-	0	0	0
SINE	Alu	B1_Mus2	chr17	12826580	12826708	-	0	0	0
LTR	MaLR	MTA_Mm	chr17	13504469	13504860	-	0	0	0
SINE	Alu	B1_Mus1	chr17	13750221	13750339	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	14172212	14172241	+	0	0	0
SINE	Alu	B1_Mus2	chr17	15244781	15244927	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	15244928	15244951	+	0	0	0
SINE	Alu	B1_Mus2	chr17	15277236	15277380	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	15277381	15277420	+	0	0	0.342623
LTR	MaLR	MTA_Mm	chr17	16124234	16124628	+	0	0	0
LTR	MaLR	MTA_Mm	chr17	16246961	16247357	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	16380048	16380082	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	16754679	16754706	+	0	0	0
SINE	Alu	B1_Mus2	chr17	17086791	17086937	-	0	0	0
SINE	Alu	B1_Mus2	chr17	17483557	17483700	+	0	0	0.615324
LTR	MaLR	MTB	chr17	17493623	17494023	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	17504220	17504244	+	1.18211	0	0
SINE	B2	B2_Mm2	chr17	17616062	17616250	-	0	0	0
SINE	Alu	B1_Mm	chr17	18037257	18037401	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	18037402	18037431	+	0	0	0
LTR	MaLR	MTA_Mm	chr17	18334696	18335084	-	0	0	0
SINE	Alu	B1_Mus1	chr17	18468903	18469052	+	0	0	0
Low_complexity	Low_complexity	A-rich	chr17	18469053	18469158	+	0	0	0.342623
LINE	L1	L1Md_F2	chr17	18482792	18488033	+	0	0	0
SINE	Alu	B1_Mm	chr17	18505707	18505853	-	0	0	0
SINE	Alu	B1_Mus1	chr17	18535278	18535425	-	0	0	0
LTR	MaLR	MTA_Mm	chr17	18780391	18780769	-	0	0	0
Low_complexity	Low_complexity	A-rich	chr17	19669523	19669690	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	20054632	20054687	+	0	0	0
LINE	L1	L1Md_F2	chr17	20424258	20430476	-	0	0	0
SINE	Alu	B1_Mus2	chr17	20574865	20575010	-	0	0	0
SINE	Alu	B1_Mm	chr17	20627771	20627917	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	20627918	20627942	+	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	20782608	20782652	+	0	0	0
LINE	L1	L1_Mus2	chr17	21016581	21022547	+	0	0	0
LTR	MaLR	MTA_Mm	chr17	21037184	21037577	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	21108397	21108446	+	0.923247	1.08938	0.685246
LTR	MaLR	MTA_Mm	chr17	21223768	21224159	-	0	0	0
LTR	MaLR	MTA_Mm	chr17	21374049	21374442	+	0	0	0
SINE	Alu	B1_Mm	chr17	21725912	21726022	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	21726023	21726047	+	0	0	0
SINE	Alu	B1_Mus1	chr17	21812801	21812945	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	21983619	21983667	+	0	0	0
SINE	Alu	B1_Mm	chr17	22308708	22308854	+	0	0	0
SINE	Alu	B1_Mus1	chr17	22793320	22793464	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	22793465	22793511	+	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr17	23668583 23668617 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	23782854 23782897 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	23837684 23837713 +	0.591053	0	1.23065
SINE	Alu	B1_Mm	chr17	24395732 24395875 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	24395876 24395902 +	0	0	0
SINE	Alu	B1_Mus2	chr10	85394623 85394769 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	85394770 85394797 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	25848550 25848590 +	0	0	0
SINE	Alu	B1_Mus1	chr17	26108157 26108304 -	0	0	0
SINE	Alu	B1_Mus2	chr17	26270494 26270632 -	0	0	0
LTR	MaLR	MTA_Mm	chr17	26385266 26385660 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	26697303 26697341 +	0	0	0
LTR	MaLR	MTA_Mm	chr17	26943872 26944267 +	0	0	0
SINE	Alu	B1_Mus1	chr17	27040084 27040224 -	0	0	0
SINE	Alu	B1_Mus1	chr17	27114557 27114705 +	1.77316	0	0.615324
SINE	Alu	B1_Mus1	chr17	27121671 27121818 +	2.82479	1.08938	0.615324
Low_complexity	Low_complexity	A-rich	chr17	27121819 27121875 +	1.18211	1.08938	0
Simple_repeat	Simple_repeat	(TTTA)n	chr17	27974637 27974660 +	0	0	0
SINE	B2	B2_Mm1t	chr17	28527787 28527978 -	1.16015	0	0
SINE	Alu	B1_Mus1	chr17	28528020 28528167 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	28648978 28649014 +	0	0	0
SINE	Alu	B1_Mm	chr17	28678465 28678613 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	28678614 28678643 +	0.664388	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	28700253 28700289 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	28726294 28726380 +	0	0	0
SINE	Alu	B1_Mus1	chr17	29055143 29055288 +	0	0	0
LTR	ERV1	MURVY-LTR	chr17	29055289 29055720 +	0	0	0
SINE	Alu	B1_Mm	chr17	29261151 29261296 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	29261297 29261332 +	0	0	0
SINE	Alu	B1_Mus2	chr17	29402688 29402836 -	0	0	0
SINE	Alu	B1_Mus1	chr17	29404263 29404410 -	0	0	0
SINE	Alu	B1_Mus1	chr17	29454208 29454353 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	29454354 29454385 +	0	0	0
SINE	Alu	B1_Mm	chr17	29591028 29591173 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	29591174 29591196 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	29622061 29622089 +	0	0	0
SINE	Alu	B1_Mus2	chr17	29643593 29643739 -	0	0	0
SINE	Alu	B1_Mus1	chr17	30124209 30124354 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	30124355 30124377 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr17	30137774 30137919 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	30137920 30137956 +	0	0	0
SINE	Alu	B1_Mur1	chr17	30138187 30138324 -	0	0	0
LTR	ERVK	RMER19B	chr17	31149283 31150195 +	0	0	0
SINE	Alu	B1_Mm	chr10	11730916 11731053 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	31866499 31866539 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	11731054 11731110 +	0	0	0
SINE	Alu	B1_Mm	chr17	32544965 32545111 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	32546465 32546502 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	32670469 32670496 +	0	0	0
SINE	Alu	B1_Mus2	chr17	33174369 33174515 +	0	0	0
LTR	MaLR	MTA_Mm	chr17	33394123 33394503 +	0	0	0
LTR	MaLR	MTA_Mm	chr17	33424071 33424451 -	0	0	0
SINE	Alu	B1_Mus1	chr17	34635113 34635257 +	0	0	0
Simple_repeat	Simple_repeat	(CAAAA)n	chr17	34635260 34635325 +	0	0	0
SINE	Alu	B1_Mus2	chr17	34916715 34916861 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	34916862 34916886 +	0	0	0
SINE	Alu	B1_Mus1	chr17	35081062 35081207 -	0	0	0
SINE	Alu	B1_Mus2	chr17	35199156 35199302 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	3918711 3918751 +	0	0	0
SINE	Alu	B1_Mus1	chr17	35342731 35342879 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	35514972 35515005 +	0	0	0
SINE	Alu	B1_Mus2	chr17	35635555 35635699 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	35635700 35635725 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	86749472 86749502 +	0	0	0
SINE	Alu	B1_Mus1	chr17	35750417 35750555 -	0	0	0
LTR	MaLR	MTA_Mm	chr17	35852525 35852914 +	0	0.761137	0
SINE	Alu	B1_Mus1	chr17	35951332 35951466 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr17	35951467 35951512 +	0	0	0
LINE	L1	L1Md_F2	chr17	36880706 36886565 +	0	0	0
SINE	Alu	B1_Mm	chr17	37108817 37108962 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	37108963 37109018 +	0	0	0
SINE	Alu	B1_Mm	chr17	37163046 37163192 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	37163193 37163219 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	37300596 37300623 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	37503935 37503982 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr17	37554022 37554056 +	0	0	0
SINE	Alu	B1_Mus1	chr17	37556619 37556766 +	0	0	0
LTR	MaLR	MTA_Mm	chr17	38123301 38123684 -	0	0	0
SINE	Alu	B1_Mus2	chr17	38135577 38135721 -	0	0	0
SINE	Alu	B1_Mm	chr17	38643454 38643596 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	11826371 11826393 +	0	0	0
SINE	Alu	B1_Mus1	chr10	3921946 3922090 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	3922091 3922120 +	0	0	0
LINE	L1	L1_Mus1	chr17	40150250 40156284 -	0	0.380568	0
SINE	Alu	B1_Mus1	chr10	87249970 87250083 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	87250084 87250125 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	40610338 40610415 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	41387614 41387659 +	0	0	0.615324
LTR	MaLR	MTA_Mm	chr17	41511802 41512203 +	0	0	0
LINE	L1	L1Md_F2	chr17	41521576 41527076 -	0	0	0
LINE	L1	L1Md_F3	chr17	41546557 41551765 +	0	0	0
SINE	Alu	B1_Mus1	chr17	41696276 41696422 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	42035332 42035362 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	42277593 42277623 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	42289867 42289895 +	0	0	0
LTR	MaLR	MTA_Mm	chr17	42397957 42398350 -	0	0	0
SINE	Alu	B1_Mm	chr17	42920252 42920398 -	0	0	0
SINE	Alu	B1_Mus2	chr17	43044631 43044776 -	0	0	0
LTR	MaLR	MTA_Mm	chr17	45343679 45344060 -	0	0	0
SINE	Alu	B1_Mus1	chr17	45623493 45623604 +	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chr17	45623605 45623746 +	0	0	0
SINE	Alu	B1_Mus1	chr17	45713187 45713315 -	0	0	0
SINE	Alu	B1_Mus2	chr17	46498660 46498808 -	0	0	0
SINE	Alu	B1_Mus2	chr17	46999538 46999683 -	2.10535	1.08938	0
SINE	Alu	B1_Mus1	chr17	47243828 47243975 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	87976211 87976238 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	88071859 88071897 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	48243890 48243923 +	0	0	0
SINE	Alu	B1_Mus1	chr17	48475636 48475781 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	48475782 48475808 +	0	0	0
SINE	Alu	B1_Mm	chr10	88123731 88123877 -	0	0	0
SINE	Alu	B1_Mm	chr10	88128170 88128315 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	88128316 88128335 +	0	0	0
SINE	Alu	B1_Mus2	chr17	49383362 49383508 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	49423217 49423258 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	49423706 49423742 +	0	0	0
SINE	Alu	B1_Mus1	chr17	51380185 51380330 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	51477181 51477208 +	0	0	0
LTR	MaLR	MTA_Mm	chr17	51778371 51778766 -	0	0	0
LTR	MaLR	MTA_Mm	chr17	51779865 51780260 -	0	0	0
LTR	MaLR	MTA_Mm	chr17	51990508 51990903 +	0.332194	2.12644	0.615324
SINE	Alu	B1_Mus2	chr17	52003306 52003452 +	3.10194	3.97696	3.21646
Simple_repeat	Simple_repeat	(A)n	chr17	52183347 52183379 +	0	0	0
SINE	Alu	B1_Mm	chr17	52188898 52189044 -	0	0	0
SINE	Alu	B1_Mus2	chr17	53693077 53693223 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	53693224 53693247 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	53700207 53700276 +	0.332194	0	0
SINE	Alu	B1_Mm	chr17	54171011 54171157 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	54171158 54171183 +	0	0	0
LINE	L1	L1_Mus4	chr17	54901791 54906250 -	0.332194	0	0
LTR	MaLR	MTA_Mm	chr17	54950684 54951077 +	0	0	0
LINE	L1	Lx	chr17	55722172 55727669 -	0	0	0
SINE	Alu	B1_Mus2	chr17	56364767 56364913 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	56364914 56364949 +	0	0	0



SINE	Alu	B1_Mus2	chr17	56391296 56391441 -	0	0	0
SINE	Alu	B1_Mm	chr17	56463499 56463646 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	56463647 56463674 +	0	0	0
SINE	Alu	B1_Mus1	chr17	56662694 56662840 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr17	56662841 56662896 +	0	0	0
SINE	B2	B2_Mm2	chr17	56689079 56689272 -	0	0	0
SINE	Alu	B1_Mm	chr17	56865259 56865405 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	56865406 56865428 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	56973859 56973891 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	57141771 57141792 +	0	0	0
LINE	L1	L1Md_F2	chr10	12050073 12055123 +	0	0	0
SINE	Alu	B1_Mus1	chr17	57269747 57269894 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	57269895 57269929 +	0	0	0
SINE	Alu	B1_Mus1	chr10	89000301 89000443 -	0	0	0
LTR	MaLR	MTA_Mm	chr17	58933115 58933510 +	0.591053	0	0.615324
LTR	MaLR	MTA_Mm	chr10	89167052 89167447 +	0	0	0
SINE	Alu	B1_Mus2	chr17	60523656 60523804 -	0	0	0
SINE	Alu	B1_Mm	chr17	60548977 60549122 +	0	0	0
SINE	Alu	B1_Mm	chr10	12098767 12098911 -	0	0	0
Low_complexity	Low_complexity	A-rich	chr17	61723210 61723271 +	0	0	0
SINE	Alu	B1_Mm	chr17	61801670 61801808 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	61801809 61801831 +	0	0	0
SINE	Alu	B1_Mus2	chr17	62357571 62357718 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	62863329 62863368 +	0	0	0
SINE	Alu	B1_Mus2	chr17	63896669 63896815 -	0	0	0
SINE	Alu	B1_Mm	chr17	63997038 63997173 -	0	0	0
SINE	Alu	B1_Mm	chr17	64083145 64083291 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	64083292 64083315 +	0	0	0
SINE	Alu	B1_Mus2	chr17	64615435 64615581 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	64615582 64615607 +	0	0	0
SINE	Alu	B1_Mus1	chr17	64946205 64946352 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	64946353 64946377 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	65226967 65227005 +	0	0	0
SINE	B2	B2_Mm1a	chr17	66518990 66519175 +	0	0	0
SINE	Alu	B1_Mm	chr17	66809630 66809777 -	0	0	0
SINE	Alu	B1_Mus2	chr17	66831332 66831478 -	0	0	0
SINE	Alu	B1_Mm	chr17	67735328 67735473 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	68526394 68526432 +	0	0	0
LTR	MaLR	MTA_Mm	chr17	69318725 69319121 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	69919837 69919863 +	3.46721	2.98884	0
SINE	Alu	B1_Mm	chr17	70279777 70279923 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	70506604 70506668 +	1.58764	1.41763	1.30057
SINE	Alu	B1_Mus2	chr17	71190591 71190737 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	71190738 71190761 +	0	0	0
SINE	Alu	B1_Mur2	chr17	71191754 71191861 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	71191862 71191887 +	0	0	0
SINE	Alu	B1_Mus2	chr17	71220491 71220627 -	0	0.380568	0
LTR	MaLR	MTA_Mm	chr17	74084609 74085004 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr10	90448338 90448358 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	74484343 74484381 +	0	0	0
SINE	Alu	B1_Mus2	chr17	74724108 74724254 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	74724255 74724293 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	74734769 74734797 +	0	0	0
Low_complexity	Low_complexity	GA-rich	chr17	74885304 74885388 +	0	0	0
SINE	Alu	B1_Mus2	chr17	74906920 74907065 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	74907066 74907107 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr17	75104467 75104511 +	1.5143	0.708814	2.06302
SINE	Alu	B1_Mus2	chr17	75264868 75264998 -	0	0	0
SINE	Alu	B1_Mm	chr17	76206011 76206151 -	0	0	0
LTR	MaLR	MTA_Mm	chr17	76759659 76760050 +	0	0	0
SINE	B4	B4	chr17	77334637 77334923 +	0	0	0
LINE	L1	L1Md_F3	chr17	78097627 78102811 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	78257725 78257765 +	0	0	0
SINE	Alu	B1_Mus1	chr10	90837604 90837739 +	0	0	0
SINE	Alu	B1_Mus2	chr17	80314386 80314529 +	0	0	0
SINE	Alu	B1_Mur3	chr17	80512062 80512181 -	0	0	0
SINE	Alu	B1_Mus2	chr17	81133328 81133457 -	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr17	81247064 81247098 +	0	0	0
SINE	Alu	B1_Mm	chr17	81693260 81693406 +	0	0	0
LTR	MaLR	MTA_Mm	chr17	82788378 82788774 -	0	0	0
LTR	MaLR	MTA_Mm	chr17	82804240 82804633 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	82870098 82870141 +	0	0	0
SINE	Alu	B1_Mus1	chr17	83632960 83633080 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	83633081 83633131 +	0	0	0
SINE	Alu	B1_Mm	chr17	83934920 83935068 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	84081195 84081232 +	0	0	0
LTR	MaLR	MTB	chr17	84084115 84084505 +	0	0	0
SINE	Alu	B1_Mm	chr17	84100570 84100715 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	84100716 84100764 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	91316842 91316871 +	0	0	0.615324
LTR	MaLR	MTA_Mm	chr17	84567201 84567597 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	84582280 84582311 +	0	0	0
SINE	Alu	B1_Mus1	chr10	91427990 91428134 -	0	0	0
SINE	Alu	B1_Mus1	chr17	85099860 85100008 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	91539514 91539540 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	85929415 85929441 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	87328727 87328752 +	0	0	0
SINE	Alu	B1_Mus1	chr17	88015348 88015494 -	0	0	0
SINE	Alu	B1_Mus2	chr17	88061572 88061717 -	0	0	0.342623
SINE	Alu	B1_Mus2	chr17	88358722 88358868 -	0	0	0
SINE	Alu	B1_Mus1	chr17	88363793 88363925 -	0	0	0
SINE	Alu	B1_Mus2	chr17	88478736 88478885 -	0	0	0
LTR	MaLR	MTA_Mm	chr17	88501436 88501831 +	0	0	0
SINE	Alu	B1_Mus1	chr17	88524210 88524343 -	0	0	0
SINE	Alu	B1_Mus2	chr17	88648903 88649049 -	0	0	0
SINE	Alu	B1_Mus1	chr17	88838885 88839033 -	0	0	0
LINE	L1	Lx3B	chr17	89631799 89636902 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	91987262 91987306 +	0.332194	0	0
SINE	B2	B2_Mm1a	chr17	92711419 92711608 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	92726095 92726141 +	0	0	0
LINE	L1	L1Md_T	chr17	93095180 93101626 -	0	0	0
SINE	Alu	B1_Mus2	chr10	92389892 92390008 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	93923399 93923427 +	0	0	0
LINE	L1	L1Md_T	chr17	94022864 94029305 -	0	0	0
LTR	MaLR	MTA_Mm	chr17	94115791 94116179 -	0	0	0
LTR	MaLR	MTB	chr17	94144431 94144815 -	0	0	0
LTR	MaLR	MTA_Mm	chr17	94319055 94319448 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr17	94484465 94484501 +	0	0	0
SINE	Alu	B1_Mus1	chr17	95268817 95268962 -	0	0	0
SINE	Alu	B1_Mm	chr18	3089757 3089876 -	0	0	0
LTR	MaLR	MTA_Mm	chr18	3568208 3568602 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	3569700 3570094 +	0.591053	0	0
SINE	Alu	B1_Mus2	chr18	4477735 4477881 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	4500541 4500571 +	0	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr18	4981768 4981812 +	0	0	0
SINE	Alu	B1_Mm	chr18	5426557 5426702 -	0	0	0
LTR	MaLR	MTA_Mm	chr18	5804371 5804764 +	0	0	0
SINE	Alu	B1_Mus2	chr18	5822532 5822678 -	0	0	0
LTR	ERV1	MuRRS-int	chr18	6546344 6551000 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	7770449 7770841 -	0	0	0
SINE	Alu	B1_Mus2	chr18	7843342 7843488 +	0	0.380568	0
SINE	Alu	B1_Mm	chr18	7856267 7856410 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	7936239 7936262 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	8381589 8381983 -	0	0	0
SINE	Alu	B1_Mm	chr18	8608302 8608454 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	8608455 8608477 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	8697113 8697158 +	0.591053	1.41763	0.615324
LINE	L1	L1VL2	chr18	8798966 8803655 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	9003743 9003787 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	9020935 9021330 -	0	0	0
LTR	MaLR	MTB	chr18	9048732 9049127 +	0	0	0
SINE	Alu	B1_Mus2	chr18	9502658 9502804 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	9502805 9502859 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	9573108 9573156 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr18	10052261 10052284 +	0	0	0
SINE	Alu	B1_Mus2	chr18	10263444 10263586 -	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chr18	10615678 10615736 +	0	0	0
SINE	Alu	B1_Mm	chr18	10861769 10861914 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	10861915 10861938 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	10898002 10898397 -	0	0	0
LINE	L1	L1Md_F2	chr18	11337701 11344102 -	0	0	0
LTR	MaLR	MTA_Mm	chr18	11650337 11650730 -	1.05163	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	11776783 11776804 +	0	0	0
SINE	B2	B2_Mm2	chr10	93283926 93284105 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	12481354 12481748 +	0	0	0
SINE	Alu	B1_Mus2	chr18	13728873 13728994 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	14482562 14482598 +	0	0	0
SINE	Alu	B1_Mus2	chr18	15479206 15479351 -	0	0	0
LTR	MaLR	MTA_Mm	chr18	15494204 15494598 -	0	0	0
SINE	B4	B4	chr18	16514649 16514897 -	0	0	0
Simple_repeat	Simple_repeat	(T)n	chr18	16606319 16606343 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	16674718 16675112 +	0	0	0
LINE	L1	L1Md_T	chr18	17979385 17986064 -	0	0	0
SINE	Alu	B1_Mm	chr18	18388935 18389081 -	0	0	0
SINE	Alu	B1_Mm	chr18	18723522 18723667 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	18997476 18997501 +	0	0	0
SINE	Alu	B1_Mus2	chr18	19552050 19552196 -	0	0	0
SINE	Alu	B1_Mm	chr18	19613017 19613163 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	19983848 19983885 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	20655836 20655873 +	0	0	0
LINE	L1	MusHAL1	chr18	20662336 20664536 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	21081785 21081823 +	0	0	0
SINE	Alu	B1_Mm	chr18	21091495 21091618 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	21091619 21091646 +	0	0	0
SINE	Alu	B1_Mus2	chr18	21203685 21203831 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	21203832 21203869 +	0	0	0
SINE	Alu	B1_Mus2	chr18	21274250 21274396 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	21274397 21274430 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	21643063 21643094 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	21907660 21907688 +	0	0	0
LINE	L1	L1_Mus3	chr18	22337358 22342617 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	22455250 22455297 +	0	0	0
SINE	Alu	B1_Mur4	chr10	94079442 94079572 +	0	0	0
Low_complexity	Low_complexity	GA-rich	chr10	94079573 94079657 +	0	0	0
SINE	B2	B2_Mm1a	chr10	94105709 94105901 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	24074915 24074990 +	0.332194	0	0
SINE	Alu	B1_Mus1	chr18	24281169 24281301 -	0.591053	0	0
SINE	Alu	B1_Mus2	chr18	24354239 24354385 -	0	0	0
SINE	Alu	B1_Mur2	chr18	24462428 24462575 +	0	1.08938	0
Simple_repeat	Simple_repeat	(A)n	chr18	24462576 24462620 +	0	1.08938	0
Simple_repeat	Simple_repeat	(A)n	chr18	24592020 24592083 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	24597607 24597646 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	25972017 25972411 -	0	0	0
LTR	MaLR	MTA_Mm	chr18	25973506 25973900 -	0	0	0
LTR	MaLR	MTA_Mm	chr18	26707258 26707652 -	0	0	0
LTR	MaLR	MTA_Mm	chr18	26844096 26844478 +	0.591053	1.01984	0
LTR	MaLR	MTA_Mm	chr18	27274716 27275095 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	27607534 27607565 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	27807682 27807719 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	29241465 29241514 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	29271975 29272368 -	0	0	0
SINE	Alu	B1_Mus1	chr18	29322149 29322296 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	29461790 29461835 +	0	0	0
SINE	Alu	B1_Mus2	chr18	29777561 29777707 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	29777708 29777745 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	29948978 29949373 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	30653631 30653665 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	30766618 30766652 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	31065538 31065932 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	31107065 31107090 +	0	0	0
SINE	Alu	B1_Mus2	chr18	32457135 32457279 -	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr18	32742638 32742680 +	0	0	0
SINE	Alu	B1_Mus1	chr18	33052291 33052437 -	0	0	0
LTR	MaLR	MTA_Mm	chr18	33548373 33548740 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	33715092 33715132 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	33738065 33738459 -	0	0	0
SINE	Alu	B1_Mus1	chr10	94919419 94919565 -	0	0	0
LTR	MaLR	MTA_Mm	chr18	34487848 34488243 -	0	0	0
SINE	Alu	B1_Mus2	chr18	34716415 34716561 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	34716562 34716585 +	0	0	0
LTR	MaLR	MTA_Mm	chr10	12689712 12690083 +	0	0	0
SINE	Alu	B1_Mus2	chr18	34848356 34848484 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	34848485 34848510 +	0.332194	0	0
SINE	Alu	B1_Mur1	chr18	34935847 34935982 -	0	0	0
SINE	Alu	B1_Mm	chr18	35119173 35119318 +	0	0	0
SINE	Alu	B1_Mus2	chr18	35158065 35158210 +	0	0	0
SINE	B2	B2_Mm1a	chr18	35916925 35917105 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	35917106 35917132 +	0	0	0.342623
SINE	Alu	B1_Mm	chr10	95085374 95085520 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	95085521 95085560 +	0	0	0
SINE	Alu	B1_Mm	chr18	36465370 36465512 +	1.18211	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	36465513 36465548 +	0.591053	0	0
SINE	Alu	B1_Mus2	chr18	36553446 36553591 -	0	0	0
LTR	MaLR	MTA_Mm	chr18	36653378 36653772 -	0	0	0
SINE	Alu	B1_Mus2	chr18	36888987 36889133 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	36889134 36889159 +	0	0	0
LTR	MaLR	MTA_Mm	chr10	95255082 95255476 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	38184591 38184620 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	38744664 38744695 +	0	0	0
SINE	Alu	PB1D10	chr18	38988271 38988373 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	38988374 38988398 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	39728964 39728992 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr18	39733971 39734006 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	41244228 41244264 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	41353615 41353650 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	41654423 41654473 +	0	0	0
SINE	Alu	B1_Mm	chr18	41825925 41826058 -	0.591053	0	0
LTR	MaLR	MTA_Mm	chr18	42161228 42161621 -	0	0	0
SINE	Alu	B1_Mm	chr18	42358387 42358533 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	95697541 95697579 +	0	0	0
SINE	Alu	B1_Mus2	chr18	42584979 42585115 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	42585116 42585148 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr18	42618184 42618215 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	95727222 95727260 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	42753173 42753211 +	0	0	0
LINE	L1	L1Md_T	chr18	43870217 43876428 -	0	0	0
SINE	Alu	B1_Mus1	chr18	43979917 43980064 -	0	0	0
SINE	Alu	B1_Mus1	chr18	44091297 44091438 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	44091439 44091484 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	96008661 96008693 +	0	0	0
SINE	Alu	B1_Mm	chr18	46780107 46780252 -	0	0	0
LTR	MaLR	MTA_Mm	chr18	47277938 47278333 +	0	0	0
SINE	Alu	B1_Mm	chr18	47665935 47666087 +	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr18	47666088 47666120 +	0	0	0
LINE	L1	L1Md_T	chr18	48370273 48377565 -	0	0.761137	0
Simple_repeat	Simple_repeat	(A)n	chr18	48466716 48466787 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	48611184 48611578 +	0	0	0
LINE	L1	L1_Mus2	chr18	48746984 48752642 +	0	0	0
SINE	Alu	B1_Mus1	chr18	49243111 49243257 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	96357538 96357577 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr10	96363882 96363918 +	0	0	0
SINE	Alu	B1F2	chr18	50384246 50384368 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	50384369 50384415 +	0	0	0
LINE	L1	L1Md_T	chr18	50421619 50428042 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	51059636 51059686 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	96486012 96486060 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	52114575 52114614 +	0	0	0
SINE	Alu	B1_Mm	chr18	52735047 52735172 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr18	52735173 52735196 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	52769966 52770362 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	52847341 52847733 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	52917911 52917955 +	0	0	0
SINE	Alu	B1_Mus1	chr10	96653357 96653475 -	0	0	0
SINE	Alu	B1_Mur4	chr18	53321787 53321937 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	53321938 53321991 +	0	0	0
SINE	Alu	B1_Mus2	chr18	53393377 53393524 -	0	0	0.615324
SINE	Alu	B1_Mus1	chr18	54268311 54268435 -	0	0	0
SINE	Alu	B1_Mm	chr18	54500993 54501138 +	0	0	0
LTR	MaLR	ORR1B2	chr18	54501139 54501487 +	0.332194	0	0
SINE	Alu	B1_Mus1	chr10	96812518 96812665 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr10	96812668 96812776 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	54899107 54899143 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	54958509 54958566 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	54977832 54977857 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	96895878 96895923 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	96923244 96923304 +	0	0	0
SINE	Alu	B1_Mm	chr18	55925669 55925822 +	0	0	0
SINE	Alu	B1_Mm	chr18	55991243 55991389 -	0	0	0
Other	Other	RMER1A	chr18	56353455 56354337 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	56441985 56442018 +	0	0	0
SINE	Alu	B1_Mm	chr18	56775449 56775594 -	0	0	0
SINE	Alu	B1_Mus2	chr18	56836666 56836813 -	0	0	0
SINE	Alu	B1_Mus2	chr18	57088719 57088865 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	57616671 57617066 -	0	0	0
SINE	Alu	B1_Mm	chr18	58123858 58124005 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	58124006 58124076 +	0	0	0.342623
LTR	MaLR	MTA_Mm	chr18	58666906 58667305 +	0	0	0
SINE	Alu	B1_Mus1	chr18	58917896 58918042 -	0	0	0
SINE	Alu	B1_Mus1	chr18	59297080 59297213 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	59579556 59579596 +	2.36421	1.41763	0.615324
Simple_repeat	Simple_repeat	(A)n	chr18	59741725 59741771 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	59897335 59897361 +	0	0	0
LINE	L1	L1Md_T	chr18	60107452 60114739 -	0	0	0
LINE	L1	L1_Mus1	chr18	60321314 60327628 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	61449375 61449405 +	0	0.708814	0
SINE	Alu	B1_Mus1	chr18	62102137 62102280 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	62264007 62264043 +	0	0	0
LTR	MaLR	MTB	chr18	62437989 62438387 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	97693772 97693807 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	62952345 62952381 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	64494502 64494540 +	0	0	0
SINE	Alu	B1_Mus1	chr18	65708481 65708626 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	65708627 65708658 +	0	0	0
SINE	Alu	B1_Mm	chr18	65849648 65849794 -	0	0	0
LTR	MaLR	MTA_Mm	chr18	65947192 65947572 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	66475516 66475549 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	66884398 66884790 -	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	67409855 67409895 +	0	0	0.615324
SINE	Alu	B1_Mus2	chr18	67433700 67433827 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	67433828 67433859 +	0	0	0
SINE	Alu	B1_Mus2	chr18	67821100 67821240 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	67821241 67821288 +	0.332194	0	0
SINE	Alu	B1_Mm	chr18	67925778 67925925 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	68073372 68073767 -	0	0	0.342623
LTR	MaLR	MTA_Mm	chr18	68074865 68075260 -	0	0	0
LTR	MaLR	MTB	chr18	69431918 69432301 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	69919004 69919397 -	0	0	0
LTR	MaLR	MTA_Mm	chr18	69920497 69920891 -	0	0	0
SINE	Alu	B1_Mus1	chr18	69958118 69958266 -	0	0	0
LTR	MaLR	MTB	chr18	70135641 70136040 -	0	0	0
SINE	Alu	B1_Mus1	chr18	70696821 70696968 -	0	0	0
LINE	L1	L1Md_T	chr18	70934968 70941530 -	0	0	0
SINE	Alu	B1_Mm	chr18	71041419 71041565 -	0	0	0
LTR	MaLR	MTB	chr18	72762758 72763155 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	73201707 73201744 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr18	73254331 73254363 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	73265269 73265301 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	73531376 73531768 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	73756797 73756838 +	0	0	0
SINE	Alu	B1_Mus2	chr18	74268085 74268225 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	74268226 74268252 +	0	0	0
SINE	Alu	B1_Mus1	chr18	74396249 74396396 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	74396397 74396434 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	75448185 75448228 +	0	0	0.615324
LTR	MaLR	MTB	chr18	75578559 75578959 +	0	0	0
LTR	MaLR	MTA_Mm	chr10	99165030 99165410 -	0	0	0
Low_complexity	Low_complexity	A-rich	chr18	77978973 77979078 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	78806194 78806563 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	78816285 78816311 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	79716502 79716541 +	0	0.380568	0
SINE	Alu	B1_Mm	chr18	79815913 79816059 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	80016999 80017031 +	0	0	0
LTR	MaLR	MTB	chr18	80107558 80107955 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	80576457 80576853 +	0	0	0
LTR	MaLR	MTB	chr10	99574978 99575374 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	81553335 81553374 +	0	0	0.615324
LTR	MaLR	MTA_Mm	chr18	81698536 81698930 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	82319965 82319994 +	0	0	0
SINE	Alu	B1_Mus1	chr18	83052231 83052372 -	0	0	0
SINE	Alu	B1_Mm	chr18	83259215 83259361 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	83371820 83371855 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	83551244 83551275 +	0	0	0
LTR	MaLR	MTA_Mm	chr18	84766836 84767226 -	0	0	0
SINE	Alu	B1_Mus2	chr18	84856809 84856955 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	84856956 84856979 +	0	0	0
SINE	Alu	B1_Mus2	chr18	84864102 84864248 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	84864249 84864271 +	0	0	0
SINE	Alu	B1_Mus2	chr18	85239437 85239583 -	0	0	0
LINE	L1	L1Md_T	chr18	85243799 85249965 -	0	0	0
LTR	MaLR	MTB	chr18	85762455 85762854 -	0.591053	0	0
LINE	L1	L1Md_F2	chr18	86052965 86058206 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	86126430 86126464 +	0	0	0
LINE	L1	L1Md_T	chr10	10013795 10014421 -	0	0	0
SINE	Alu	B1_Mm	chr18	86553986 86554135 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	86893504 86893535 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	87120587 87120634 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	87402035 87402077 +	0	0	0
LINE	L1	L1Md_T	chr18	87946452 87952725 -	0	0	0
LINE	L1	L1Md_F3	chr18	88095741 88100732 +	0	0	0
SINE	Alu	B1_Mus1	chr18	88184852 88184999 +	0	0	0
LTR	ERVK	RLTR40	chr18	88302633 88303354 -	0	0	0
SINE	Alu	B1_Mus1	chr18	88449432 88449582 -	0	0	0
SINE	Alu	B1_Mus1	chr18	88723407 88723553 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr18	89939591 89939636 +	0.332194	0.708814	0.615324
LTR	MaLR	MTA_Mm	chr10	10053254 10053294 -	0.591053	0	0
LTR	MaLR	MTA_Mm	chr10	10053400 10053439 -	0	0	0
SINE	Alu	B1_Mus1	chr19	3705813 3705953 -	0	0	0
SINE	Alu	B1_Mm	chr19	3827422 3827568 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	3827569 3827608 +	0	0	0
SINE	Alu	B1_Mm	chr19	3828136 3828274 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	3828275 3828297 +	0	0	0
LTR	MaLR	MTA_Mm	chr19	4074319 4074712 -	0	0	0
SINE	Alu	B1_Mm	chr19	4090151 4090297 -	0	0	0
SINE	Alu	B1_Mus1	chr19	4211142 4211289 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	4211290 4211309 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	10078193 10078196 +	0	0	0
SINE	Alu	B1_Mus1	chr19	4845064 4845208 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	4845209 4845240 +	0	0	0
SINE	Alu	B1_Mus2	chr19	5017178 5017324 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	5017325 5017347 +	0	0	0
SINE	Alu	B1_Mus1	chr19	5077728 5077874 -	0	0	0.615324
SINE	Alu	B1_Mm	chr19	5124031 5124178 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr19	5124179	5124226	+	0	0	0
SINE	Alu	B1_Mus1	chr19	5427612	5427730	+	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr19	5427731	5427751	+	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr19	6096511	6096538	+	0	0	0
SINE	Alu	B1_Mm	chr19	6584348	6584468	-	0	0	0
SINE	B2	B2_Mm1a	chr19	6679833	6680024	-	0	0	0
SINE	B2	B2_Mm1a	chr19	7293964	7294154	-	0	0	0
SINE	B2	B2_Mm1a	chr19	7327208	7327400	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	7327401	7327436	+	0	0	0
SINE	Alu	PB1D9	chr19	8760132	8760229	-	0	0	0
SINE	Alu	B1_Mus2	chr19	8939330	8939476	-	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr19	8944989	8945020	+	0	0	0
LTR	MaLR	MTA_Mm	chr19	10435750	10436143	-	0	0	0
SINE	Alu	B1_Mus1	chr19	10706805	10706950	+	0	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr19	10706951	10706999	+	0	0	0
SINE	Alu	B1_Mm	chr19	11802292	11802438	+	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chr19	11802439	11802562	+	0	0	0
SINE	Alu	B1_Mus1	chr19	11957739	11957860	-	0	0	0
SINE	Alu	B1_Mus1	chr19	12231754	12231896	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	12231897	12231916	+	0	0	0
SINE	Alu	B1_Mm	chr19	12796790	12796936	-	0	0	0
SINE	Alu	B1_Mm	chr19	12829338	12829483	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	12829484	12829534	+	0	0.380568	0
LTR	MaLR	MTA_Mm	chr19	13314864	13315258	+	0	0	0
SINE	Alu	B1_Mus1	chr19	13334458	13334606	-	0	0	0
LTR	MaLR	MTA_Mm	chr19	14174456	14174851	+	0	0	0
SINE	Alu	B1_Mus2	chr19	14717569	14717715	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	14717716	14717745	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	14890298	14890347	+	0	0	0
SINE	Alu	B1_Mus1	chr10	10194394	10194409	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	10194409	10194412	+	0	0	0
SINE	Alu	B1_Mus2	chr19	15003945	15004091	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	15004099	15004127	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	15393298	15393324	+	0	0	0
SINE	Alu	B1_Mus2	chr19	15475002	15475153	-	0	0	0
SINE	Alu	B1_Mm	chr19	15605740	15605890	-	0	0	0
SINE	B2	B2_Mm1a	chr19	15959439	15959633	-	0	0	0
SINE	Alu	B1_Mm	chr19	16001482	16001628	+	0	0	0
LTR	MaLR	MTA_Mm	chr19	16087981	16088370	+	0	0	0
LTR	MaLR	MTA_Mm	chr19	18213872	18214265	+	0	0	0
LTR	MaLR	MTA_Mm	chr10	10230185	10230224	+	0	0	0
SINE	Alu	B1_Mus2	chr19	18647040	18647195	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	18647196	18647234	+	0	0	0
SINE	Alu	B1_Mus1	chr19	18648926	18649073	-	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	10233774	10233783	+	0	0	0
SINE	Alu	B1_Mm	chr19	18743477	18743621	-	0	0	0
SINE	Alu	B1_Mus1	chr19	18794272	18794418	-	0	0	0
LTR	MaLR	MTA_Mm	chr19	19593980	19594375	-	0	0	0
LTR	MaLR	MTA_Mm	chr19	20034010	20034405	+	0	0	0
LTR	MaLR	MTA_Mm	chr19	20122207	20122602	+	0	0	0
LTR	MaLR	MTA_Mm	chr19	20261434	20261829	-	0	0	0
SINE	Alu	B1_Mus1	chr19	20304059	20304177	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	20304178	20304211	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	10250894	10250896	+	0.332194	0	0
LINE	L1	L1Md_T	chr19	20641820	20648285	-	0	0	0
LTR	MaLR	MTA_Mm	chr19	22039581	22039975	-	0.332194	0	0
LTR	MaLR	MTA_Mm	chr19	22041075	22041469	-	0	0	0
SINE	Alu	B1_Mm	chr19	22164720	22164867	+	0.591053	1.7982	1.23065
Simple_repeat	Simple_repeat	(A)n	chr19	22164868	22164888	+	0.591053	1.08938	1.23065
LTR	MaLR	MTA_Mm	chr19	22204308	22204702	-	0	0	0
LTR	MaLR	MTA_Mm	chr19	23104524	23104920	-	0	0	0
SINE	Alu	B1_Mm	chr19	23133864	23134010	-	0	0	0.342623
SINE	Alu	B1_Mus2	chr19	23134048	23134194	-	0	0	0
Simple_repeat	Simple_repeat	(GAA)n	chr19	23149084	23149140	+	0	0	0
LTR	MaLR	MTA_Mm	chr19	23175485	23175880	+	0	0	0
SINE	Alu	B1_Mus1	chr19	23177712	23177859	-	0	0	0
SINE	Alu	B1_Mus2	chr10	10291654	10291669	+	0	0	0

LTR	MaLR	MTA_Mm	chr10	10298541 10298581 -	0	0	0
LTR	MaLR	MTA_Mm	chr10	10298690 10298729 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	10306608 10306612 +	0	0	0
LTR	MaLR	MTA_Mm	chr19	25907230 25907629 +	0	0	0
LTR	MaLR	MTEa	chr19	26133047 26133394 -	0	0	0
SINE	Alu	B1_Mus1	chr19	26361369 26361516 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	26361517 26361549 +	0.332194	0	0
LTR	MaLR	MTA_Mm	chr19	26610794 26611184 -	0	0	0
LTR	MaLR	MTB	chr19	26617862 26618241 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr19	26998710 26998747 +	0	0	0
SINE	Alu	B1_Mus1	chr19	27102135 27102283 -	0	0	0.342623
SINE	Alu	B1_Mus2	chr19	27417240 27417386 +	0	0	0
LTR	MaLR	MTA_Mm	chr19	27513852 27514248 -	0	0	0
SINE	Alu	B1_Mm	chr19	27572527 27572674 -	0	0	0
LTR	MaLR	MTB	chr19	28258806 28259201 +	0	0	0
SINE	Alu	B1_Mus2	chr19	28889841 28889987 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	28889988 28890008 +	0	0	0
SINE	Alu	B1_Mm	chr19	28908738 28908884 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	28908885 28908910 +	0	0	0
LTR	MaLR	MTB	chr10	10338822 10338860 +	0	0	0
SINE	Alu	B1_Mus2	chr19	29909683 29909829 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	29909830 29909876 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	10354169 10354176 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	10356117 10356123 +	0	0	0
SINE	B2	B2_Mm1t	chr19	32192069 32192256 +	0	0	0
LTR	MaLR	MTA_Mm	chr19	33053204 33053602 -	0	0	0
scRNA	scRNA	4.5SRNA	chr19	33191619 33191711 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	34525384 34525431 +	0.332194	0	0
SINE	Alu	B1_Mus2	chr19	34870294 34870439 -	0	0	0.342623
SINE	Alu	B1_Mm	chr19	34984337 34984485 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	34984486 34984524 +	0	0	0
SINE	Alu	B1_Mur2	chr19	35098062 35098207 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	35779707 35779758 +	0	0	0.615324
LTR	MaLR	MTA_Mm	chr19	35794164 35794557 -	0	0	0
SINE	Alu	B1_Mus1	chr19	35923113 35923258 +	0	0	0
SINE	Alu	B1_Mus2	chr19	36893551 36893697 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	37449267 37449309 +	0	0.380568	0.615324
SINE	Alu	B1_Mm	chr19	37943755 37943897 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	37943898 37943918 +	0	0	0
SINE	Alu	B1_Mus2	chr19	37946337 37946482 +	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chr19	37946485 37946550 +	0	0	0
SINE	Alu	B1_Mm	chr19	38402108 38402254 -	0.591053	0	0
SINE	Alu	B1_Mm	chr19	38409012 38409158 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	38409159 38409182 +	0	0	0
SINE	Alu	B1_Mus2	chr19	38496837 38496984 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	10447293 10447297 +	0	0	0
LINE	L1	Lx2	chr19	40189487 40194397 +	0	0	0
LTR	MaLR	MTA_Mm	chr19	40363599 40363996 +	0	0	0
SINE	Alu	B1_Mus2	chr19	41608543 41608655 +	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr19	41608656 41608682 +	0	0.708814	0
SINE	Alu	B1_Mus2	chr19	41651768 41651900 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr19	41651901 41651926 +	0	0.380568	0.342623
SINE	Alu	B1_Mur4	chr19	41897400 41897552 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	41897553 41897592 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	41904105 41904152 +	0	0	0
SINE	Alu	B1_Mur3	chr19	41985776 41985909 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	41985910 41985934 +	0	0	0
SINE	Alu	B1_Mus1	chr19	42231957 42232103 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	42232104 42232131 +	0.332194	0.380568	0
SINE	Alu	B1_Mm	chr19	43718488 43718631 -	0	0	0
LTR	MaLR	MTA_Mm	chr19	44914848 44915243 -	1.16015	0	0
LTR	MaLR	MTA_Mm	chr19	44916601 44916995 -	0	0	0
SINE	Alu	B1_Mm	chr19	45384107 45384243 -	0	0	0
LTR	MaLR	MTA_Mm	chr19	45788033 45788427 -	0	0	0
SINE	Alu	B1_Mus2	chr19	46124381 46124516 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	46124517 46124543 +	0	0	0
SINE	Alu	B1_Mm	chr19	46199986 46200132 +	0	0	0



Simple_repeat	Simple_repeat	(A)n	chr19	46200133	46200170 +	0	0	0
SINE	Alu	B1_Mus2	chr19	46422425	46422570 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	46422571	46422596 +	0	0	0
SINE	Alu	B1_Mus2	chr19	46430761	46430906 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	48883820	48883866 +	0	0	0
LTR	MaLR	MTA_Mm	chr19	49081293	49081684 -	0	0	0
LTR	ERVK	RMER6A	chr19	49400680	49401465 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	49570268	49570304 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	49783242	49783281 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	51439220	51439287 +	0	0	0
LINE	L1	L1Md_T	chr19	51618350	51624488 -	0	0	0
LTR	MaLR	MTA_Mm	chr19	52967003	52967384 -	0	0	0
SINE	Alu	B1_Mm	chr19	53655435	53655581 -	0	0	0
SINE	Alu	B1_Mus2	chr19	53730937	53731080 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	53731081	53731104 +	0	0	0
LTR	ERVK	RMER19C	chr19	54134835	54135556 -	0	0	0
SINE	Alu	B1_Mus2	chr19	54152806	54152955 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	54253620	54253653 +	0	0	0
SINE	Alu	B1_Mus1	chr19	54581927	54582072 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	54582073	54582096 +	0	0	0
SINE	B4	B4	chr19	54582609	54582942 -	0	0	0
SINE	Alu	B1_Mm	chr19	56592810	56592956 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	56592957	56593003 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	57317526	57317556 +	0	0	0
SINE	Alu	B1_Mm	chr19	57484311	57484459 -	0	0	0
SINE	Alu	B1_Mus1	chr19	58668834	58668981 +	0	0	0
SINE	Alu	B1_Mus2	chr19	58672924	58673067 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	58673068	58673099 +	0	0	0
SINE	Alu	B1_Mm	chr19	58730706	58730852 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	58730853	58730879 +	0	0	0
SINE	Alu	B1_Mm	chr19	59254812	59254952 +	0	0	0
SINE	Alu	B1_Mus1	chr19	60448860	60449004 -	0	0	0
SINE	Alu	B1_Mm	chr19	60876199	60876335 -	0	0	0
SINE	Alu	B1_Mus2	chr19	60958032	60958180 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr19	60958181	60958204 +	0	0	0
LTR	MaLR	MLT1A	chr1	3158390	3158755 +	0	0	0
SINE	Alu	B1_Mus1	chr1	3895094	3895241 +	0	0	0
Low_complexity	Low_complexity	GA-rich	chr1	3895243	3895307 +	0.591053	0	0
LTR	MaLR	MTA_Mm	chr1	4459694	4460087 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	5363027	5363399 +	0	0	0
SINE	Alu	B1_Mm	chr1	5974350	5974496 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	6118540	6118930 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	6166783	6166808 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	6316887	6317281 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	7186902	7186942 +	0	0	0
LINE	L1	L1Md_T	chr1	7318763	7325821 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	7379898	7379924 +	0	0	0
SINE	Alu	B1_Mus1	chr1	7383886	7384024 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	7384121	7384170 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	10758043	10758047 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	7736046	7736092 +	0	0	0
SINE	Alu	B1_Mus2	chr1	8225645	8225791 -	0	0	0
LTR	MaLR	MTA_Mm	chr10	13926283	13926680 +	0	0	1.10507
Simple_repeat	Simple_repeat	(A)n	chr1	10280912	10281032 +	0	0	0
SINE	Alu	B1_Mus2	chr1	10743193	10743339 -	0	0	0
SINE	Alu	B1_Mm	chr1	10869851	10869944 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr1	10869945	10869990 +	0	0	0.342623
SINE	Alu	B1_Mus1	chr10	10786471	10786485 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr10	10786485	10786489 +	0	0	0.342623
SINE	Alu	B1_Mus1	chr1	12076729	12076875 -	0	0	0.615324
LTR	MaLR	MTA_Mm	chr1	12137289	12137684 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	12213548	12213577 +	0	0	0
LTR	MaLR	MTB	chr1	12214505	12214876 +	0	0	0
LINE	L1	L1_Mus2	chr1	12350343	12355914 +	0	0	0
LINE	L1	L1Md_F2	chr1	12447775	12453014 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	12507060	12507455 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	13024878	13025261 -	0	0	0

SINE	Alu	B1_Mm	chr1	13438957 13439103 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	13442647 13443045 -	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	13579979 13580019 +	0	0.380568	0
SINE	Alu	B1F	chr1	13582238 13582381 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	13701247 13701641 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	13712190 13712583 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	13713683 13714076 +	0	0	0
SINE	Alu	B1_Mm	chr1	14460313 14460459 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	14808372 14808426 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	14909142 14909180 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	14911900 14912289 -	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr1	15950048 15950108 +	0	0	0
SINE	Alu	B1_Mus1	chr1	16087059 16087203 +	0	0	0
SINE	Alu	B1_Mm	chr1	16712179 16712324 -	0	0	0
SINE	Alu	B1_Mus1	chr1	16842186 16842333 -	0	0	0
LINE	L1	L1_Mus2	chr1	17128125 17133630 -	0	0	0
SINE	B2	B2_Mm1t	chr1	17224679 17224871 -	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	17688764 17688804 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	17701866 17702245 -	0	0	0
LINE	L1	L1Md_T	chr1	17906212 17912668 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	17996654 17997047 -	0	0	0.342623
SINE	Alu	B1_Mus1	chr1	18470778 18470924 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	18631436 18631484 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	18771358 18771724 +	0	0	0
LINE	L1	L1_Mus1	chr1	18904036 18909923 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	19014247 19014641 +	0	0	0
LINE	L1	L1Md_F	chr1	19261551 19266776 +	0	0	0
SINE	Alu	B1_Mus1	chr10	10871125 10871140 +	0	0	0
Low_complexity	Low_complexity	GA-rich	chr10	10871140 10871149 +	0	0	0
SINE	Alu	B1_Mm	chr1	19532328 19532459 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	22905052 22905445 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	23046296 23046329 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	23216073 23216100 +	0	0	0
SINE	Alu	B1_Mus1	chr1	23227188 23227334 +	0	0	0
Simple_repeat	Simple_repeat	(CAAAA)n	chr1	23227335 23227443 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	23628441 23628835 +	0	0	0
LTR	MaLR	MTB	chr1	23724107 23724501 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	24892423 24892490 +	0	0	0
LTR	ERVK	RLTR25A	chr1	24904866 24905869 -	0	0	0
SINE	Alu	B1_Mm	chr10	14089893 14090035 -	0	0	0
SINE	Alu	B1_Mm	chr1	25067710 25067856 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	25068897 25068927 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr1	25909043 25909065 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	14094117 14094187 +	0	0	0
LINE	L1	L1_Mus3	chr1	26013440 26019720 -	0	0	0
SINE	Alu	B1_Mm	chr1	26092146 26092292 +	0	0	0
SINE	Alu	B1_Mus1	chr1	26457791 26457938 -	0	0	0
SINE	Alu	B1_Mus2	chr1	26801855 26802000 +	0	0	0.615324
LINE	L1	L1Md_F2	chr1	27047027 27052077 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	27246077 27246111 +	0.332194	0	0
LTR	MaLR	MTA_Mm	chr10	10955646 10955685 -	7.93678	4.15543	4.86694
LTR	MaLR	MTA_Mm	chr1	27744272 27744665 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	28557389 28557422 +	0	0	0
LINE	L1	L1Md_F	chr1	28571431 28575921 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	28685159 28685282 +	0	0	0
SINE	Alu	B1_Mus2	chr1	28861585 28861731 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	29279945 29279975 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	10977416 10977419 +	0	0	0
SINE	B2	B3A	chr1	29833803 29834006 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	29971928 29971964 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	30158936 30159330 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	30528675 30528710 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	30532249 30532289 +	0	0	0
LINE	L1	L1Md_T	chr1	30717415 30723900 -	0	0	0
SINE	Alu	B1_Mus2	chr1	30963441 30963587 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	30963588 30963611 +	0	0	0
SINE	Alu	B1_Mus2	chr1	31669624 31669770 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr1	31669771 31669823 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	31827931 31828326 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	31829418 31829813 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	32036162 32036198 +	0	0	0
LTR	MaLR	MTA_Mm	chr10	11010427 11010466 -	0	0	0
LTR	MaLR	MTA_Mm	chr10	11010576 11010616 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	32804228 32804251 +	0	0	0
SINE	Alu	B1_Mus2	chr10	11015227 11015241 -	0	0	0
SINE	Alu	B1_Mus2	chr1	33261708 33261847 -	0	0	0
LINE	L1	L1_Mus2	chr1	33399530 33405715 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	33801796 33801828 +	0	0	0
SINE	Alu	PB1D9	chr1	34458705 34458815 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	34797641 34797678 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	35232788 35233182 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	35236546 35236942 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	35319340 35319729 +	0	0	0
SINE	Alu	B1_Mus1	chr1	35394174 35394317 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	35394318 35394346 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	35756051 35756082 +	0	0	0
SINE	Alu	B1_Mm	chr1	36079100 36079250 +	0	0	0
SINE	B2	B2_Mm2	chr1	36101237 36101424 +	0	0	0
LTR	MaLR	MTB	chr10	11049098 11049138 +	0	0	0
SINE	Alu	B1_Mur3	chr1	37140556 37140708 -	0	0	0
SINE	Alu	B1_Mus2	chr10	11054555 11054570 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	11054570 11054576 +	0	0	0
SINE	Alu	B1_Mus1	chr1	37468941 37469087 -	0	0	0
SINE	Alu	B1_Mm	chr10	11057331 11057346 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	11057346 11057350 +	0	0	0
SINE	Alu	B1_Mm	chr10	11059640 11059655 -	0	0	0
LINE	L1	Lx2B	chr1	37993708 38000000 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	38037315 38037708 -	0	0	0
SINE	Alu	B1_Mus2	chr1	38731063 38731208 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	39672535 39672570 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	40103785 40104179 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	40105275 40105669 -	0	0	0
SINE	Alu	B1_Mm	chr10	11079942 11079956 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	40428697 40429073 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	40609711 40610109 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	41332932 41333006 +	0	0	0
LINE	L1	L1Md_F2	chr1	41376983 41381722 -	0	0	0
LINE	L1	L1Md_F2	chr10	14294518 14300080 -	0	0	0.342623
SINE	Alu	B1_Mm	chr10	14300081 14300227 -	0	0	0.342623
SINE	Alu	B1_Mm	chr10	11088394 11088408 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	41867243 41867277 +	0	0	0
SINE	Alu	B1_Mus1	chr1	41919013 41919159 -	0	0	0
SINE	Alu	B1_Mus1	chr1	42343155 42343295 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	42768187 42768226 +	0	1.08938	0.342623
Simple_repeat	Simple_repeat	(A)n	chr1	42827815 42827843 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	43001033 43001427 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	11100530 11100533 +	0	0	0
SINE	Alu	B1_Mus1	chr1	44545194 44545341 +	0	0	0
SINE	B2	B2_Mm1a	chr1	44586160 44586352 -	0	0	0
LINE	L1	L1Md_F2	chr1	45136138 45142296 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	45884874 45884910 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	45894464 45894513 +	0	0	0
LINE	L1	L1Md_F2	chr1	46552425 46558574 -	0	0	0
SINE	Alu	B1_Mm	chr1	46649209 46649354 -	0	0	0
SINE	B2	B2_Mm1t	chr1	46842767 46842959 +	0	0.380568	0
SINE	Alu	B1_Mus2	chr1	46970938 46971084 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	46971085 46971125 +	0	0	0
SINE	Alu	B1_Mus1	chr1	47229595 47229743 -	0	0	0.342623
SINE	B2	B2_Mm1a	chr1	47543669 47543857 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	47543858 47543878 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	47702515 47702562 +	0	0	0
SINE	Alu	B1_Mus1	chr1	48498396 48498541 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	48871131 48871184 +	0	0	0
SINE	Alu	B1_Mus1	chr1	49248345 49248492 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr1	49248493 49248522 +	0	0	0
SINE	Alu	B1_Mus2	chr10	11148585 11148599 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	11148599 11148602 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	49316357 49316751 +	0	0	0
SINE	Alu	B1_Mm	chr1	49708529 49708674 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	50028698 50029092 -	0.923247	0.380568	0.342623
LTR	MaLR	MTA_Mm	chr1	50030190 50030584 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	50344401 50344433 +	0	0	0
LINE	L1	L1Md_F2	chr1	50493795 50499762 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	50684891 50684929 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	51253632 51253666 +	0	0.708814	0.342623
SINE	Alu	B1_Mus1	chr10	11167123 11167138 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	52429869 52430264 -	0	0	0
SINE	Alu	B1_Mus1	chr1	52617115 52617262 -	0	0	0
SINE	Alu	B1_Mm	chr1	52895859 52896006 -	0	0	0
LTR	MaLR	MTA_Mm	chr10	11191084 11191124 -	0	0	0
SINE	Alu	B1_Mus2	chr1	53789066 53789211 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	53789212 53789240 +	0	0	0
SINE	Alu	B1_Mus2	chr1	55176705 55176851 +	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr1	55176852 55176905 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	56087064 56087458 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	56088554 56088948 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	57038499 57038545 +	0	0	0.342623
LTR	MaLR	MTA_Mm	chr1	57165838 57166232 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	57690903 57690938 +	0.332194	0	0.342623
SINE	Alu	B1_Mus1	chr10	11232431 11232446 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	11232447 11232450 +	0	0	0
SINE	Alu	PB1D9	chr1	58437743 58437857 -	0	0	0
SINE	Alu	B1_Mus1	chr1	58688443 58688589 -	0	0	0
SINE	Alu	B1_Mus2	chr1	58695278 58695422 -	0	0	0
SINE	Alu	B1_Mus1	chr1	59778547 59778688 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	59785820 59785864 +	0	0	0
SINE	Alu	B1_Mm	chr1	60103686 60103828 +	0	0	0
SINE	Alu	B1_Mus2	chr1	60119095 60119241 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	60119242 60119266 +	0	0	0
SINE	Alu	B1_Mus1	chr1	60210878 60211019 -	0	0	0
SINE	Alu	B1_Mus1	chr1	60234634 60234780 +	0	0.708814	0.342623
Simple_repeat	Simple_repeat	(A)n	chr1	60234781 60234833 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr1	61573288 61573325 +	0	0.380568	0
SINE	Alu	B1_Mus1	chr1	63379270 63379417 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	63379422 63379442 +	0	0	0
SINE	Alu	B1_Mm	chr1	64463532 64463678 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	64574900 64574949 +	0	0	0
SINE	Alu	B1_Mus2	chr1	64676923 64677068 +	0	0.380568	0
SINE	Alu	B1_Mm	chr1	65004753 65004921 +	0	0	0
SINE	Alu	B1_Mm	chr1	65005722 65005867 -	0	0	0
SINE	Alu	B1_Mm	chr1	65018166 65018313 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	11309956 11309961 +	0	0	0
SINE	Alu	B1_Mus2	chr10	11312522 11312537 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	11312537 11312541 +	0	0	0.342623
LTR	MaLR	MTA_Mm	chr1	65989316 65989697 -	0	0	0
SINE	Alu	B1_Mus1	chr1	67479764 67479910 -	0	0	0
LINE	L1	Lx	chr1	68034051 68039844 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	11343793 11343797 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	11358143 11358147 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr1	71582665 71582702 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	71586102 71586135 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	71774368 71774425 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	72473394 72473422 +	0	0	0
SINE	B2	B2_Mm2	chr1	72750873 72751064 +	0	0	0
SINE	Alu	B1_Mus2	chr1	72847535 72847681 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	72847682 72847716 +	0	0	0
SINE	Alu	B1_Mus1	chr1	74231092 74231205 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	74231206 74231229 +	0	0	0
SINE	Alu	B1_Mus2	chr1	74870171 74870315 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	74870322 74870350 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	75268033 75268429 +	0	0.708814	0.342623

SINE	Alu	B1_Mus2	chr1	75647256 75647392 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	75834615 75835010 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	75888599 75888984 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	14665430 14665456 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	75920600 75920637 +	0	0.380568	0
LINE	L1	L1Md_F3	chr1	76375361 76381361 -	0	0	0
LINE	L1	L1VL4	chr1	76513567 76516547 -	0	0	0
SINE	Alu	B1_Mus2	chr1	76838671 76838803 -	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr1	77040601 77040639 +	0	0	0
LTR	MaLR	MTB	chr1	77092600 77092996 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	11441380 11441383 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	77842130 77842535 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	77999505 77999543 +	0	0	0
SINE	Alu	B1_Mm	chr1	78245116 78245261 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	78245262 78245287 +	0	0	0
LTR	MaLR	MTB	chr1	78537264 78537637 +	0	0	0
SINE	Alu	B1_Mm	chr1	78548664 78548810 +	0	0	0
SINE	Alu	B1_Mus1	chr1	78604597 78604724 -	0	0	0
SINE	Alu	B1_Mm	chr10	11451049 11451063 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	11451064 11451068 +	0	0	0
SINE	Alu	B1_Mm	chr1	78630435 78630581 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	79544362 79544746 +	0.332194	0	0
SINE	Alu	B1_Mus2	chr1	79695462 79695597 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	79695598 79695626 +	0	0	0
SINE	Alu	B1_Mus1	chr1	80368926 80369071 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	80371992 80372031 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	80759732 80759770 +	0	0	0
SINE	Alu	B1_Mur4	chr1	80885014 80885160 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	81022049 81022082 +	0	0	0
SINE	Alu	B1_Mus2	chr1	81037805 81037949 +	0	0	0
SINE	Alu	B1_Mm	chr1	81802938 81803085 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	81845800 81845829 +	0	0	0
SINE	Alu	B1_Mus1	chr1	82771555 82771702 -	0	0	0
SINE	Alu	B1_Mus2	chr1	82815080 82815225 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	82815226 82815247 +	0	0	0
LINE	L1	L1Md_T	chr1	83507613 83513485 -	0.332194	0	0.342623
SINE	Alu	B1_Mus2	chr10	11488503 11488518 -	0	0	0
SINE	Alu	B1_Mus2	chr1	84286579 84286725 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	84286726 84286754 +	0	0	0
SINE	Alu	B1_Mus2	chr1	85164942 85165087 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	85329392 85329429 +	1.25544	1.41763	0.342623
SINE	Alu	B1_Mus2	chr1	87392031 87392176 -	0	0	0
SINE	Alu	B1_Mus1	chr1	88328710 88328856 -	0	0	0
SINE	Alu	B1_Mus1	chr1	88342228 88342376 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	88342377 88342403 +	0.591053	0	0
SINE	Alu	B1_Mus1	chr1	88406560 88406707 +	0	0	0.615324
Low_complexity	Low_complexity	GA-rich	chr1	88406708 88406784 +	0	0	0.615324
SINE	Alu	PB1D9	chr1	88417721 88417826 -	0	0	0
SINE	Alu	B1_Mus2	chr1	89217411 89217557 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	89513727 89514121 +	0	0	0.615324
LTR	MaLR	MTA_Mm	chr1	89515215 89515609 +	0	0	0.685246
LTR	MaLR	MTA_Mm	chr1	89948657 89949051 +	0	0	0
SINE	Alu	B1_Mm	chr1	90958757 90958903 -	0	0	0
SINE	Alu	B1_Mus2	chr1	92316637 92316783 -	0	0	0
SINE	Alu	B1_Mm	chr1	93144220 93144378 -	0	0	0.342623
SINE	Alu	B1_Mm	chr1	93357799 93357945 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	94092247 94092284 +	0	0	0
SINE	Alu	B1_Mus1	chr1	94329757 94329903 -	0	0	0
SINE	Alu	B1_Mus1	chr1	94330038 94330185 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	94330186 94330212 +	0	0	0
SINE	Alu	B1_Mur4	chr1	94869199 94869348 -	0	0	0
SINE	Alu	B1_Mus1	chr1	95551990 95552137 -	0	0	0
SINE	Alu	PB1D7	chr1	95576725 95576831 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	95787570 95787604 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	96496164 96496219 +	0	0	0
SINE	Alu	B1_Mus2	chr1	96700005 96700150 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	96700151 96700191 +	0	0	0

LINE	L1	L1Md_F	chr1	97242919 97247255 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr1	97360974 97361013 +	0	0	0
LINE	L1	L1Md_T	chr1	97407974 97414683 -	0	0	0
SINE	Alu	B1_Mus1	chr1	97978262 97978402 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	99476851 99476885 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	10120414 10120420 +	0	0	0
SINE	Alu	B1_Mus2	chr1	10121164 10121179 -	0	0	0
LINE	L1	L1Md_F	chr1	10126110 10126658 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	10163186 10163192 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	10253237 10253241 +	0	0	0.342623
SINE	Alu	B1_Mm	chr1	10280856 10280871 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	10280871 10280873 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	10291367 10291376 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	10292913 10292918 +	0	0	0
SINE	Alu	B1_Mm	chr1	10315447 10315462 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	10315462 10315464 +	0	0	0
LINE	L1	L1Md_F2	chr1	10393977 10394614 -	0	0	0
SINE	Alu	B1_Mus1	chr1	10396751 10396766 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	10396766 10396772 +	0	0	0
LINE	L1	L1Md_F3	chr1	10397622 10398130 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr1	10478708 10478717 +	0	0	0
SINE	Alu	B1_Mus1	chr1	10481710 10481724 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	10481724 10481730 +	0	0	0
LTR	MaLR	MTA_Mm	chr10	11662319 11662358 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	10509100 10509105 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	10523012 10523015 +	0	0	0
SINE	Alu	B1_Mus1	chr10	11664588 11664601 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	11664601 11664604 +	0	0	0
SINE	Alu	B1_Mus1	chr1	10564359 10564374 -	0	0	0
SINE	B2	B2_Mm1t	chr10	11675181 11675198 -	0	0	0
SINE	Alu	B1_Mus2	chr10	11675201 11675215 -	0	0	0
SINE	Alu	B1_Mus1	chr1	10708863 10708876 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	10714107 10714111 +	0	0	0
SINE	Alu	B1_Mus2	chr10	11688358 11688372 -	0	0	0
scRNA	scRNA	4.5SRNA	chr1	10890147 10890156 -	0	1.01984	0
SINE	Alu	B1_Mus2	chr1	10909839 10909854 -	0	0	0
LINE	L1	L1Md_T	chr1	10909929 10910613 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	10930568 10930572 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	10973940 10973944 +	0	0	0
LTR	MaLR	MTB	chr10	11694408 11694447 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	10995290 10995293 +	0	0	0
LINE	L1	L1Md_F2	chr1	11029331 11029905 -	0	0	0
SINE	Alu	B1_Mm	chr1	11060952 11060965 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	11085281 11085283 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	11109373 11109377 +	0	0	0
SINE	Alu	B1_Mus2	chr1	11132066 11132081 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	11132081 11132083 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	11143826 11143866 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	11144024 11144063 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	11205179 11205186 +	0	0	0
SINE	Alu	B1_Mus1	chr1	11223710 11223724 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	11238147 11238150 +	0	0	0
SINE	Alu	B1_Mus2	chr1	11295658 11295673 -	0	0	0
SINE	Alu	B1_Mm	chr1	11297351 11297365 -	0	0	0
SINE	Alu	B1_Mus1	chr1	11304918 11304933 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	11322190 11322197 +	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chr1	11323865 11323873 +	0	0	0
LINE	L1	L1_Mus1	chr1	11325896 11326511 -	0	0	0
SINE	Alu	B1_Mus2	chr10	11714975 11714989 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	11334748 11334753 +	0	0	0
LINE	L1	L1Md_A	chr1	11335729 11336255 +	0	0	0
SINE	Alu	B1_Mus2	chr10	11715763 11715777 -	0	0	0
SINE	Alu	B1_Mm	chr10	11715923 11715938 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	11715938 11715940 +	0	0	0
SINE	Alu	B1_Mus2	chr1	11391113 11391127 -	0	0	0
SINE	Alu	B1_Mm	chr1	11396307 11396321 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	11485948 11485954 +	0	0	0

LINE	L1	L1_Mus2	chr1	11488659 11489231 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	11539049 11539089 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	11576142 11576145 +	0	0	0
LINE	L1	L1Md_F3	chr1	11628967 11629584 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	11637803 11637809 +	0	0	0
LTR	ERV_L	RMER15	chr1	11641217 11641273 +	0	0	0
SINE	Alu	B1_Mus2	chr1	11654815 11654829 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	11654829 11654833 +	0.332194	0	0
SINE	Alu	B1_Mus1	chr1	11671479 11671493 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	11671493 11671496 +	0	0	0
SINE	Alu	B1_Mus2	chr1	11679012 11679026 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	11679026 11679029 +	0	0	0
SINE	Alu	B1_Mus1	chr1	11700889 11700904 -	0	0	0
LINE	L1	Lx	chr1	11745692 11746279 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr1	11755369 11755372 +	0	0	0
LTR	MaLR	MTB	chr10	11748531 11748571 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	11848686 11848725 +	0	0	0
SINE	Alu	B1_Mus2	chr1	11864350 11864365 +	0.591053	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr1	11864365 11864367 +	0.591053	0.380568	0
LTR	MaLR	MTA_Mm	chr1	11870243 11870283 +	0	0	0
SINE	Alu	B1_Mus2	chr1	11872334 11872349 +	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr1	11884649 11884652 +	0	0	0
LINE	L1	L1_Mus1	chr1	11979934 11980445 +	0	0	0
SINE	Alu	B1_Mm	chr1	12051677 12051690 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	12130500 12130503 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	12234005 12234009 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	12332727 12332729 +	0.591053	0	0
LINE	L1	L1Md_F2	chr1	12362083 12362613 +	0	0	0
SINE	Alu	B1_Mm	chr1	12385738 12385751 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr1	12385751 12385757 +	0	0	0
SINE	Alu	B1_Mm	chr1	12394364 12394379 -	0	0	0
SINE	Alu	B1_Mus1	chr1	12423698 12423709 +	0	0	0
Simple_repeat	Simple_repeat	(GAAAA)n	chr1	12423709 12423718 +	0	0	0
SINE	Alu	B1_Mm	chr1	12440453 12440468 -	0	0	0
SINE	Alu	B1_Mus2	chr10	11827377 11827391 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	12449961 12449966 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	12450786 12450825 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr1	12595714 12595717 +	0	0	0
SINE	Alu	B1_Mus1	chr1	12680749 12680763 +	0	0	0
SINE	Alu	B1_Mus2	chr1	12739253 12739268 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	12739268 12739272 +	0	0	0.342623
LTR	MaLR	MTB_Mm	chr10	11868676 11868717 +	0	0	0
LTR	MaLR	MTB	chr10	11873989 11874028 -	0	0	0
LTR	MaLR	MTA_Mm	chr10	4328933 4329328 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	11877148 11877152 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	13013734 13013773 -	0	0	0
SINE	Alu	B1_Mm	chr1	13032274 13032289 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	13108106 13108145 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	13108255 13108294 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	13232685 13232724 +	0	0	0
SINE	Alu	B1_Mm	chr1	13384750 13384764 -	0	0	0
SINE	Alu	B1_Mus1	chr1	13388301 13388316 -	0	0	0
SINE	Alu	B1_Mus2	chr1	13397016 13397030 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	13397030 13397038 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	13404887 13404891 +	0	0	0
SINE	B2	B3	chr1	13538516 13538533 -	0	0	0.342623
SINE	Alu	B1_Mus2	chr1	13556225 13556239 +	0	0	0
LTR	MaLR	ORR1C2	chr1	13556241 13556273 +	0	0	0
SINE	Alu	B1_Mus2	chr1	13594823 13594838 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	13594838 13594842 +	0	0	0
SINE	B2	B3A	chr1	13605945 13605962 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	13606228 13606267 -	0	0	0
SINE	Alu	B1_Mm	chr1	13614407 13614421 -	0	0	0
SINE	Alu	B1_Mus2	chr1	13642916 13642931 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	13642931 13642933 +	0	0	0
SINE	Alu	B1_Mus2	chr10	11973053 11973067 +	0	0	0
SINE	Alu	B1_Mus2	chr1	13708823 13708838 -	0	0	0

SINE	Alu	B1_Mm	chr1	13718184	13718198	-	0	0	0
SINE	Alu	B1_Mus2	chr10	11992208	11992222	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	11992222	11992229	+	0	0	0
SINE	Alu	B1_Mm	chr10	11995171	11995185	-	0	0	0
SINE	Alu	B1_Mm	chr1	13850821	13850836	-	0	0	0
SINE	Alu	B1_Mus2	chr1	13854444	13854459	-	0	0	0
SINE	Alu	B1_Mus1	chr1	13859194	13859208	-	0	0	0
LTR	MaLR	MTA_Mm	chr1	13864657	13864695	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	13864884	13864888	+	0	0	0
SINE	Alu	B1_Mus1	chr10	11998611	11998626	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	11998626	11998629	+	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	12000377	12000380	+	0	0	0
LINE	L1	L1Md_T	chr1	13923543	13924208	-	0	0	0
SINE	Alu	B1_Mus2	chr10	12004878	12004892	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	13951629	13951632	+	0	0	0
SINE	Alu	B1_Mus2	chr1	13983178	13983192	+	1.5143	0.708814	0.615324
Simple_repeat	Simple_repeat	(A)n	chr1	13983192	13983194	+	1.25544	0.708814	1.23065
LINE	L1	L1Md_T	chr1	14020199	14020893	-	0	0	0
SINE	Alu	B1_Mus2	chr1	14079966	14079981	-	0	0	0
LTR	MaLR	MTA_Mm	chr10	12017468	12017507	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	12019818	12019823	+	0	0	0
LTR	MaLR	MTA_Mm	chr1	14146380	14146421	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	14181060	14181064	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	14181968	14181973	+	0	0	0
SINE	Alu	B1_Mur4	chr1	14296440	14296453	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	14296453	14296456	+	0	0	0
SINE	Alu	B1_Mus2	chr1	14328887	14328901	-	0	0	0
LTR	MaLR	MTA_Mm	chr1	14364799	14364839	-	0	0	0
LTR	MaLR	MTA_Mm	chr1	14364948	14364988	-	0	0	0
SINE	Alu	B1_Mus2	chr10	12042584	12042599	+	0	0	0
Low_complexity	Low_complexity	A-rich	chr10	12042599	12042603	+	0	0	0
SINE	Alu	B1_Mus1	chr1	14395316	14395331	-	0	0	0
LINE	L1	L1MA4	chr1	14424562	14425024	-	0	0	0
SINE	Alu	B1_Mm	chr1	14472084	14472099	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	14496056	14496059	+	0	0	0
LINE	L1	L1Md_F2	chr1	14513118	14513640	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	14540217	14540221	+	0	0	0
SINE	Alu	B1_Mm	chr1	14607830	14607845	-	0	0	0
SINE	Alu	B1_Mm	chr1	14633201	14633215	-	0	0	0
LTR	MaLR	MTA_Mm	chr1	14655686	14655725	-	0	0	0
LTR	ERVK	MYSERV16_l-int	chr1	14663571	14663778	-	0	0	0
LTR	MaLR	MTA_Mm	chr1	14669456	14669495	+	0	0	0
LTR	MaLR	MTA_Mm	chr10	15516148	15516547	-	0	0	0
SINE	Alu	B1_Mm	chr1	14738447	14738459	-	0	0	0
LTR	MaLR	MTA_Mm	chr1	14743159	14743198	+	0	0	0.342623
LTR	MaLR	MTA_Mm	chr1	14825875	14825915	-	0	0	0.615324
LTR	MaLR	MTA_Mm	chr1	14883835	14883873	-	0	0	0
LTR	MaLR	MTA_Mm	chr10	12084294	12084333	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	14973345	14973348	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	14989865	14989868	+	0	0	0
LTR	MaLR	MTA_Mm	chr1	15015908	15015948	-	0	0	0
SINE	Alu	B1_Mus2	chr1	15115082	15115097	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	15115097	15115106	+	0	0	0
LTR	MaLR	MTA_Mm	chr1	15115192	15115230	-	0	0	0
SINE	Alu	B1_Mus2	chr1	15120184	15120199	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	15120199	15120202	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	15155955	15155959	+	0	0	0
SINE	Alu	B1_Mm	chr10	12097639	12097654	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	12097654	12097657	+	0	0	0
LINE	L1	L1Md_F3	chr1	15184503	15184952	+	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	15186650	15186654	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	15208674	15208679	+	0	0	0
SINE	Alu	B1_Mus2	chr1	15314533	15314547	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	15314547	15314550	+	0	0	0
SINE	Alu	B1_Mus2	chr1	15338097	15338111	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	15338111	15338115	+	0	0	0
SINE	Alu	B1_Mus1	chr1	15423440	15423454	+	0	0	0



Simple_repeat	Simple_repeat	(A)n	chr1	15423455	15423457 +	0	0	0
SINE	Alu	B1_Mus2	chr1	15477272	15477287 -	0	0	0
SINE	Alu	B1_Mm	chr1	15518605	15518620 +	0	0	0
SINE	Alu	B1_Mus1	chr1	15521264	15521278 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	15521279	15521283 +	0	0	0
SINE	Alu	B1_Mus1	chr1	15523166	15523179 -	0	0	0
SINE	Alu	B1_Mm	chr1	15534755	15534770 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	15534770	15534772 +	0	0	0
SINE	Alu	B1_Mm	chr1	15540089	15540103 -	0	0	0
SINE	Alu	B1_Mus1	chr1	15549467	15549482 -	0	0	0
SINE	Alu	B1_Mm	chr1	15581492	15581507 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	15581507	15581509 +	0	0	0
LINE	L1	L1Md_T	chr10	15610249	15616584 -	0	0	0
SINE	Alu	B1_Mus1	chr1	15709574	15709588 -	0	0	0
SINE	Alu	B1_Mus1	chr1	15710966	15710980 -	0	0.708814	0
SINE	Alu	B1_Mm	chr1	15736029	15736044 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	15736044	15736047 +	0.332194	0	0
SINE	B2	B3	chr1	15736047	15736064 +	0.332194	0	0
SINE	Alu	B1_Mm	chr1	15737106	15737121 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	15782503	15782543 -	0	0	0
SINE	Alu	B1_Mus1	chr1	15783568	15783582 -	0	0	0
SINE	Alu	B1_Mus2	chr1	15818059	15818073 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	15818074	15818076 +	0	0	0
SINE	B2	B2_Mm1a	chr1	15845290	15845309 -	0	0	0
SINE	Alu	B1_Mus2	chr1	15848166	15848180 -	0	0	0
SINE	Alu	B1_Mm	chr1	15873126	15873141 -	0	0	0
SINE	B2	B2_Mm1a	chr1	15887296	15887315 -	0	0	0
SINE	Alu	B1_Mm	chr1	15887628	15887642 +	0	0	0
SINE	Alu	B1_Mus2	chr1	16320040	16320054 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	16320054	16320056 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	16348587	16348626 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	16359025	16359061 +	0.332194	0	0
SINE	B2	B2_Mm1t	chr1	16381962	16381981 -	0	0	0
SINE	Alu	B1_Mus1	chr1	16384561	16384575 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	16391583	16391587 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	16603060	16603099 -	0	0	0
SINE	Alu	B1_Mus2	chr1	16644782	16644796 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	16644797	16644799 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	16650788	16650792 +	0	0.380568	0.342623
SINE	Alu	B1_Mm	chr1	16719420	16719437 -	0	0	0
SINE	Alu	B1_Mm	chr1	16826711	16826726 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	16826726	16826728 +	0	0	0
SINE	Alu	B1_Mus1	chr10	12264241	12264255 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	12264256	12264258 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	16878946	16878949 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	16908306	16908308 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	16941142	16941183 +	0	0	0
SINE	Alu	B1_Mus1	chr1	17049117	17049131 +	0	0	0
SINE	Alu	B1_Mus2	chr1	17067491	17067506 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	17098593	17098632 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	17125178	17125217 +	0	0	0
SINE	Alu	B1_Mus1	chr1	17140339	17140353 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	17224313	17224317 +	0	0	0
SINE	Alu	B1_Mm	chr1	17283154	17283168 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	17283169	17283171 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	17293364	17293367 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	17318020	17318024 +	0	0	0.615324
SINE	Alu	B1_Mus2	chr1	17432604	17432618 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	12341748	12341752 +	0	0	0
SINE	Alu	B1_Mur4	chr10	12344922	12344936 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	12344936	12344941 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	17607958	17607962 +	0	0	0
SINE	Alu	B1_Mus2	chr1	17697111	17697126 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	17697126	17697129 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	17874213	17874217 +	0	0	0.342623
SINE	Alu	B1_Mm	chr1	17926703	17926717 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	17926717	17926720 +	0	0	0

SINE	Alu	B1_Mm	chr1	17926954	17926969 -	0	0	0
SINE	Alu	B1_Mus2	chr10	12383812	12383827 -	0	0	0
LINE	L1	L1Md_T	chr1	17985947	17986592 -	0	0	0
SINE	Alu	B1_Mur3	chr1	18008099	18008114 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	18008114	18008118 +	0	0	0
SINE	B2	B2_Mm2	chr1	18018847	18018866 +	0	0.708814	0
SINE	Alu	B1_Mm	chr1	18022641	18022657 -	0	0	0
SINE	Alu	B1_Mus1	chr1	18027269	18027283 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	18027283	18027286 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	18248458	18248464 +	0	0	0
SINE	Alu	B1_Mus2	chr1	18249198	18249212 -	0	0	0
SINE	Alu	B1_Mm	chr1	18256524	18256538 -	0	0	0
Simple_repeat	Simple_repeat	(CCAA)n	chr1	18320542	18320545 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	18320545	18320548 +	0	0	0
LTR	MaLR	MTB	chr1	18320558	18320596 +	0	0	0
LTR	MaLR	MTC	chr1	18325040	18325083 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	18327431	18327433 +	0	0	0
SINE	Alu	B1_Mus1	chr10	12429515	12429529 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	12429529	12429532 +	0	0	0
SINE	Alu	B1_Mus2	chr1	18356546	18356561 -	0	0	0
SINE	Alu	B1_Mus2	chr1	18369061	18369076 -	0	0	0
SINE	Alu	B1_Mus2	chr1	18400639	18400654 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	18400654	18400656 +	0	0	0
SINE	Alu	B1_Mm	chr1	18407683	18407699 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	18407699	18407701 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	18410166	18410170 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	18417173	18417207 +	0	0	0
SINE	Alu	B1_Mus1	chr1	18421971	18421986 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	18427452	18427490 +	0	0	0
SINE	B2	B2_Mm2	chr1	18448274	18448292 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	18460665	18460704 -	0	0	0
LTR	MaLR	MTB	chr1	18468697	18468735 +	0	0	0
LINE	L1	L1Md_A	chr1	18563364	18563884 +	0	0	0.615324
SINE	Alu	B1_Mm	chr1	18564024	18564039 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	18564039	18564041 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	18579494	18579533 -	0	0	0
SINE	Alu	B1_Mm	chr1	18582573	18582588 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	18591635	18591672 -	0	0	0
scRNA	scRNA	4.5SRNA	chr1	18628971	18628980 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	18643865	18643869 +	0	0	0
SINE	Alu	B1_Mus2	chr1	18649547	18649561 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	18649562	18649568 +	0	0	0
SINE	Alu	B1_Mus2	chr1	18651108	18651123 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	12460273	12460277 +	0	0	0
SINE	Alu	B1F1	chr1	18687420	18687435 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	18687435	18687437 +	0.591053	0	0
LTR	MaLR	MTA_Mm	chr1	18766752	18766791 +	0	0	0
SINE	B2	B2_Mm1t	chr1	18862658	18862676 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	18930818	18930858 -	0	0	0
LTR	MaLR	MTA_Mm	chr10	12502629	12502668 +	0	0	0
LTR	MaLR	MTA_Mm	chr1	19078965	19079003 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	19092659	19092662 +	0	0	0
SINE	Alu	B1_Mus2	chr1	19182386	19182400 +	0	0	0
SINE	Alu	B1_Mus1	chr1	19191874	19191889 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	19245327	19245366 -	0	0	0
LTR	MaLR	MTA_Mm	chr1	19252164	19252203 +	0	0	0
LTR	MaLR	ORR1D2-int	chr1	19296447	19296586 -	0	0	0
SINE	Alu	B1_Mus2	chr1	19348887	19348901 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	19348901	19348904 +	0	0	0
SINE	Alu	B1_Mus2	chr1	19511831	19511845 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	19511845	19511848 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	19542514	19542519 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	19591308	19591311 +	0	0	0
SINE	B2	B2_Mm1t	chr1	19636896	19636915 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr1	19636915	19636919 +	0	0.380568	0
SINE	Alu	B1_Mus2	chr2	3174406	3174524 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	3174525	3174560 +	0	0	0

SINE	Alu	B1_Mus1	chr2	4910347	4910485	-	0	0	0
SINE	Alu	B1_Mus2	chr2	4941248	4941393	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	4941394	4941420	+	0	0	0
SINE	Alu	B1_Mus2	chr2	5188456	5188602	-	0	0.761137	0
SINE	ID	ID4_	chr2	5639734	5639813	-	0	0	0
SINE	Alu	B1_Mm	chr10	12605613	12605628	-	0	0	0
LTR	MaLR	MTB	chr10	12613028	12613067	-	0	0	0
LINE	L1	L1Md_F	chr2	7668777	7674169	+	0	0	0
LTR	MaLR	MTA_Mm	chr2	8019490	8019896	+	0	0	0.342623
LTR	MaLR	MTA_Mm	chr2	8094306	8094695	-	0	0	0
LINE	L1	L1Md_T	chr2	8104924	8112296	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	8137806	8137832	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	9344965	9345004	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	9346563	9346599	+	0	0	0
SINE	Alu	B1_Mus2	chr2	9535307	9535432	+	0	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr2	9535433	9535457	+	0	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr2	9633930	9633967	+	0	0	0
SINE	Alu	B1_Mus1	chr2	9974454	9974601	-	0	0	0.342623
SINE	Alu	B1_Mm	chr2	10069667	10069812	-	0	0	0
SINE	Alu	B1_Mus2	chr2	10184888	10185030	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	10185031	10185063	+	0	0	0
SINE	Alu	B1_Mm	chr2	10996014	10996166	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	11042593	11042632	+	0	0	0
SINE	Alu	B1_Mm	chr2	11536529	11536665	-	0	0	0
LTR	MaLR	MTB	chr2	11658208	11658603	-	0.332194	0	1.10507
LTR	MaLR	MTA_Mm	chr2	12411968	12412365	-	0	0	0.342623
LTR	MaLR	MTA_Mm	chr2	12704276	12704671	-	0	0	0
LTR	MaLR	MTA_Mm	chr2	12792305	12792684	-	0	0	0
LTR	MaLR	MTA_Mm	chr2	12793772	12794151	-	0	0	0
LINE	L1	L1_Mus3	chr2	13838048	13843246	-	0	0	0
SINE	Alu	B1_Mus1	chr2	15183472	15183605	-	0	0	0
LINE	L1	L1_Mus1	chr2	15571270	15577248	-	0	0	0
SINE	Alu	B1_Mus1	chr2	15891210	15891323	-	0	0	0
LINE	L1	L1Md_F3	chr2	15987699	15993003	-	0	0	0
SINE	B2	B2_Mm1a	chr2	15993004	15993194	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	12682722	12682725	+	0	0	0
LINE	L1	L1Md_T	chr2	17045803	17053054	-	0	0.708814	0.615324
SINE	Alu	B1_Mus2	chr2	17810394	17810540	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	17810541	17810568	+	0	0	0
SINE	Alu	B1_Mus1	chr2	17835780	17835927	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	17859018	17859044	+	0	0	0
SINE	B2	B3	chr2	17942916	17943132	+	0	0	0
Low_complexity	Low_complexity	A-rich	chr2	17943133	17943148	+	0	0	0
SINE	Alu	B1_Mus1	chr2	18489599	18489751	-	0	0	0
SINE	Alu	PB1	chr2	18570164	18570294	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	18570295	18570330	+	0	0	0
LTR	MaLR	MTA_Mm	chr2	19716613	19717007	+	0	0	0.615324
LTR	MaLR	MTA_Mm	chr2	19874272	19874666	+	0	0	0.615324
SINE	Alu	B1_Mus2	chr2	20738694	20738840	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	20761569	20761616	+	0.332194	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr2	22609311	22609357	+	0	0	0.342623
SINE	Alu	B1_Mm	chr2	24222450	24222582	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	24421154	24421184	+	0	0	0
SINE	B2	B2_Mm1a	chr2	25112232	25112423	-	4.2745	4.08066	1.55647
SINE	Alu	B1_Mus2	chr2	25234837	25234983	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	25308113	25308140	+	0	0	0
SINE	Alu	B1_Mus1	chr2	25408962	25409108	-	0	0	0
SINE	Alu	B1_Mm	chr2	25846423	25846540	-	0	0	0
SINE	Alu	B1_Mus2	chr2	26312691	26312836	-	0	0	0
SINE	Alu	B1_Mm	chr2	26480540	26480686	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	26642995	26643040	+	0	0	0
SINE	Alu	B1_Mus2	chr2	26886568	26886721	-	0	0	0
SINE	Alu	B1_Mm	chr2	27366863	27367008	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	27367009	27367035	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	27432622	27432652	+	0	0	0.342623
Low_complexity	Low_complexity	A-rich	chr2	27471180	27471254	+	0	0	0
LTR	ERV_L	MT2B	chr2	27646356	27646921	-	0	0	0

SINE	Alu	B1_Mm	chr2	27646922 27647070 -	0	0	0
SINE	B2	B2_Mm2	chr2	27648007 27648211 -	0	0	0
SINE	Alu	B1_Mm	chr2	27648212 27648358 -	0	0	0
SINE	Alu	B1_Mm	chr2	28298026 28298182 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	28298183 28298208 +	0	0	0
SINE	Alu	B1_Mur4	chr2	28333744 28333875 -	0	0	0
SINE	B2	B2_Mm2	chr2	28792407 28792561 +	0	0	0
SINE	Alu	B1_Mus1	chr10	12765388 12765403 -	0	0	0
SINE	Alu	B1_Mus2	chr2	28965950 28966095 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	28966096 28966126 +	0	0	0
SINE	Alu	B1_Mus2	chr2	28970577 28970722 -	0	0	0
SINE	Alu	B1_Mus2	chr2	29045436 29045582 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	29049719 29049761 +	0	0	0
SINE	Alu	B1_Mus2	chr2	29987658 29987804 -	0	0	0
SINE	Alu	B1_Mus2	chr2	30036357 30036502 -	0	0	0
SINE	Alu	B1_Mus2	chr2	30072376 30072508 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	30072509 30072536 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	30219269 30219304 +	0	0	0
SINE	Alu	B1_Mus2	chr2	30584386 30584532 +	0	0	0
SINE	Alu	B1_Mm	chr10	12781178 12781193 -	0	0	0
SINE	Alu	B1_Mus2	chr2	31074800 31074946 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	31074947 31074970 +	0	0	0
LTR	MaLR	MTA_Mm	chr2	31181852 31182247 -	0	0	0
SINE	Alu	B1_Mus2	chr2	31407169 31407315 -	0	0	0
SINE	Alu	PB1D9	chr2	31484280 31484427 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	31540855 31540898 +	0	0	0
SINE	Alu	B1_Mus2	chr2	31985852 31985999 -	0	0	0
SINE	Alu	B1_Mm	chr2	32131259 32131405 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	32131406 32131462 +	0	0	0.342623
LTR	MaLR	MTB_Mm	chr2	32260946 32261348 +	0	0	0
SINE	Alu	B1_Mus2	chr2	32314881 32315027 +	0	0	0
SINE	Alu	B1_Mus2	chr2	32339271 32339417 -	0	0	0
SINE	Alu	B1_Mus2	chr2	32952829 32952975 -	0	0.708814	0
SINE	Alu	B1_Mus2	chr2	33329746 33329892 -	0	0	0
SINE	Alu	B1_Mm	chr10	12799254 12799266 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	12799266 12799268 +	0.332194	0	0
SINE	Alu	B1_Mus2	chr10	12803614 12803629 -	0	0	0
SINE	Alu	B1_Mus2	chr2	34270572 34270718 +	0	0	0
SINE	B2	B2_Mm1a	chr10	12805311 12805330 -	0	0	0
SINE	Alu	B1_Mus1	chr2	34530110 34530221 -	0.591053	0	0.615324
SINE	Alu	B1_Mus2	chr2	34682611 34682757 -	0	0	0
SINE	Alu	B1_Mus2	chr2	34704026 34704171 -	0	0	0
SINE	Alu	B1_Mus2	chr2	35361679 35361824 -	0	0	0
SINE	Alu	B1_Mus2	chr10	12813408 12813422 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	12813422 12813425 +	0	0	0
LINE	L1	L1Md_F2	chr2	36318816 36324107 +	0	0	0
LTR	MaLR	MTA_Mm	chr2	36583643 36584036 -	0	0	0
LTR	MaLR	MTA_Mm	chr2	36585133 36585528 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	36727353 36727380 +	0	0	0
LTR	MaLR	MTA_Mm	chr2	36944566 36944960 -	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr2	37045703 37045754 +	0	0	0
SINE	Alu	B1_Mm	chr2	37590106 37590244 -	0	0	0
SINE	B2	B2_Mm1t	chr2	38190104 38190296 -	0	0	0
LTR	MaLR	MTA_Mm	chr2	38522241 38522637 +	0	0	0
SINE	Alu	B1_Mm	chr2	38849302 38849447 -	0	0	0
LINE	L1	L1Md_T	chr2	39096000 39102538 -	0	0	0
LTR	MaLR	MTA_Mm	chr2	39592935 39593314 -	0	0	0
SINE	Alu	B1_Mus1	chr2	39771964 39772109 -	0	0	0
LINE	L1	L1_Mus1	chr2	40268705 40274952 +	0	0	0
SINE	Alu	B1_Mm	chr2	40290357 40290503 -	0	0	0
SINE	Alu	B1_Mus2	chr2	40308786 40308932 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	42733048 42733126 +	0	0	0
SINE	Alu	B1_Mus2	chr2	42954132 42954278 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	43052043 43052082 +	0	0	0.342623
Low_complexity	Low_complexity	A-rich	chr2	44364201 44364253 +	0	0	0
SINE	Alu	B1_Mm	chr2	44773401 44773546 -	0.591053	3.42939	3.41924
Simple_repeat	Simple_repeat	(A)n	chr2	45168811 45168837 +	0	0	0.615324

Simple_repeat	Simple_repeat	(A)n	chr2	45245759 45245799 +	0	0	0
SINE	Alu	B1_Mus2	chr2	45583728 45583873 -	0	0	0
LTR	MaLR	MTA_Mm	chr2	45662432 45662827 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	45960948 45960987 +	0	0	0
LINE	L1	Lx	chr2	46065352 46071469 +	0	0	0
SINE	Alu	B1_Mus2	chr2	46258723 46258869 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	46438228 46438291 +	0	0	0
LTR	MaLR	MTA_Mm	chr2	46644108 46644502 +	0	0	0
LINE	L1	L1Md_T	chr2	46978311 46984999 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	47444296 47444372 +	0	0	0
SINE	Alu	B1_Mus2	chr2	47900533 47900678 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	48183114 48183141 +	0	0	0
LTR	MaLR	MTA_Mm	chr2	48207153 48207547 -	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr2	48253409 48253447 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	12915602 12915607 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	48269214 48269251 +	0	0	0
SINE	Alu	B1_Mus2	chr2	48285677 48285791 +	0	0	0
LTR	MaLR	MTA_Mm	chr2	49852366 49852761 +	0	0	0
LINE	L1	L1_Mus2	chr2	50093544 50098384 +	0	0	0
SINE	Alu	B1_Mus2	chr2	50859728 50859870 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	50859871 50859916 +	0	0	0
LTR	MaLR	MTA_Mm	chr2	51650234 51650632 +	0	0	0
SINE	Alu	B1F2	chr2	52708737 52708874 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	52708875 52708898 +	0	0	0
SINE	Alu	B1_Mm	chr10	12970117 12970131 -	0	0	0
SINE	Alu	B1_Mm	chr2	53803224 53803370 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	53803371 53803394 +	0	0	0
LINE	L1	L1Md_Gf	chr10	12974136 12974622 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	54279650 54279684 +	0	0	0
SINE	Alu	B1_Mus2	chr2	55522730 55522860 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	55800233 55800279 +	0	0	0.342623
LINE	L1	L1Md_A	chr2	55902154 55908318 -	0	0	0
LINE	L1	L1Md_F	chr2	56195494 56200714 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	56372732 56372772 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr10	12997844 12997848 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	56944017 56944051 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	58115573 58115614 +	0	0	0
SINE	Alu	B1_Mus2	chr2	59206251 59206397 -	0	0	0
SINE	Alu	B1_Mus1	chr11	3148936 3149084 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	3149085 3149108 +	0	0	0
LTR	MaLR	MTB	chr2	59444099 59444506 -	0	0	0
SINE	Alu	B1_Mm	chr2	60013074 60013219 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	60013220 60013247 +	0.664388	0	0
SINE	Alu	B1_Mm	chr2	60035799 60035945 -	1.77316	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	60975299 60975334 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	61065448 61065487 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	61108608 61108690 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	61495503 61495539 +	0	0	0
SINE	Alu	B1_Mus2	chr2	61760922 61761066 -	0	0	0
LTR	MaLR	MTA_Mm	chr2	61859480 61859875 +	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr2	64359226 64359254 +	0	0	0
SINE	Alu	B1_Mus1	chr11	3451341 3451487 +	0	0	0
SINE	Alu	B1_Mus1	chr2	64593170 64593319 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	64593320 64593348 +	0	0	0
SINE	Alu	B1_Mus1	chr2	64876634 64876773 -	0	0	0
SINE	Alu	B1_Mus2	chr11	3487094 3487238 +	0	0	0
SINE	Alu	B1_Mus2	chr2	65152663 65152809 -	0.591053	0	1.23065
Simple_repeat	Simple_repeat	(GAAAA)n	chr11	3487239 3487296 +	0	0	0
SINE	Alu	B1_Mus2	chr11	3514977 3515112 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	65986615 65986653 +	0	0	0
SINE	Alu	B1_Mus1	chr2	65991120 65991267 -	0	0	0
SINE	Alu	B1_Mus1	chr2	69045129 69045267 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	69045268 69045317 +	0	0	0
SINE	B2	B2_Mm1a	chr2	69675563 69675750 -	1.8711	0	1.10507
SINE	Alu	B1_Mm	chr2	72117938 72118084 -	0	0	0
SINE	Alu	B1_Mm	chr2	72456529 72456669 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	72515787 72515829 +	0	0	0.342623

SINE	Alu	B1_Mus1	chr2	72891845 72891982 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	72891983 72892008 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	73104912 73104955 +	0	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr2	73873861 73873937 +	0	0	0
SINE	Alu	B1_Mm	chr2	74022372 74022519 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	74022520 74022599 +	0	0.380568	0.342623
Simple_repeat	Simple_repeat	(A)n	chr2	74987483 74987553 +	0	0	0
LINE	L1	L1Md_T	chr2	75064941 75071646 -	0	0	0
SINE	Alu	B1_Mus1	chr2	75457827 75457974 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	75457975 75458000 +	0	0	0
SINE	Alu	B1_Mm	chr2	75509888 75510034 +	0	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr2	75510035 75510060 +	0	0.380568	0
SINE	B2	B2_Mm2	chr2	75616894 75617078 -	0	0	0
SINE	Alu	B1_Mus1	chr2	76207358 76207505 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	4502051 4502085 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	78179694 78179734 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	78351205 78351235 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	78706556 78706578 +	0	0	0
SINE	Alu	B1_Mur3	chr2	78848527 78848636 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	79272503 79272538 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	79523881 79523928 +	0	0	0
SINE	Alu	B1_Mus1	chr11	4598527 4598675 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	4598676 4598699 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	80339974 80340002 +	0	0	0
SINE	Alu	B1_Mus1	chr2	80524282 80524419 -	0	0	0
SINE	Alu	B1_Mus1	chr2	80596000 80596147 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	81278827 81278879 +	0	0	0
LINE	L1	L1Md_T	chr2	82329221 82336204 -	0	0	0
SINE	Alu	B1_Mus1	chr11	4753427 4753572 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	4753573 4753596 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	4760996 4761040 +	0	0	0
LTR	MaLR	MTB	chr2	83536245 83536646 +	0	0	0
LTR	MaLR	MTA_Mm	chr2	83842446 83842829 +	0.332194	0	0
LTR	MaLR	MTA_Mm	chr2	83981174 83981569 +	0	0	0
LTR	MaLR	MTA_Mm	chr2	84005610 84006005 -	0	0	0
SINE	B2	B2_Mm1a	chr2	84421684 84421875 -	0	0	0.615324
SINE	Alu	B1_Mus1	chr2	84424716 84424863 -	0	0	0
SINE	B2	B2_Mm1a	chr2	84673658 84673847 -	0	0	0
SINE	Alu	B1_Mm	chr2	85124221 85124364 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr2	85124365 85124416 +	0.332194	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr2	85406802 85406824 +	0	0	0
SINE	Alu	B1_Mus2	chr2	86047210 86047348 -	0	0	0
SINE	Alu	B1_Mus1	chr2	86198269 86198415 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	86198416 86198460 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	86261295 86261330 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	86445850 86445884 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr2	86594905 86595027 +	0	0	0
SINE	Alu	B1_Mm	chr2	86836376 86836522 -	0	0	0
LTR	MaLR	MTA_Mm	chr2	86901012 86901404 +	0.591053	0	0
SINE	Alu	B1_Mm	chr2	87335028 87335174 -	0	0	0
SINE	Alu	B1_Mus2	chr2	87337192 87337337 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	87826766 87826818 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	87929669 87929752 +	0	0	0
LINE	L1	L1Md_F3	chr2	88006664 88012110 +	0	0	0
LINE	L1	L1Md_T	chr2	88545232 88551738 -	0	0	0
SINE	Alu	B1_Mm	chr2	88566862 88567002 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	88567003 88567031 +	0	0	0.342623
LINE	L1	L1Md_T	chr2	89355875 89359530 -	0	0	0
LINE	L1	L1Md_T	chr2	89733231 89739899 -	0	0	0
LTR	MaLR	MTA_Mm	chr2	90431303 90431690 -	0	0	0.957947
SINE	Alu	B1_Mus1	chr2	90466500 90466647 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	90466648 90466680 +	0	0	0
SINE	Alu	B1_Mus1	chr2	90582071 90582215 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	90582216 90582235 +	0.591053	0	0
SINE	Alu	B1_Mus1	chr2	91564369 91564512 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	91569820 91569872 +	0	0	0
SINE	B2	B2_Mm1a	chr11	5393491 5393681 -	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr2	93438493 93438529 +	0.591053	0	0
LTR	MaLR	MTA_Mm	chr2	93759975 93760370 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	94103205 94103249 +	0	0.380568	0
SINE	Alu	B1_Mus2	chr11	5599008 5599134 -	0	0	0
SINE	Alu	B1_Mus2	chr2	94568900 94569046 -	0	0	0
LTR	MaLR	MTA_Mm	chr2	94685320 94685708 +	0	0	0
LINE	L1	L1Md_F2	chr2	94929034 94935257 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	95158669 95158713 +	0	0	0
SINE	Alu	B1_Mus1	chr2	95161043 95161186 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	95161187 95161215 +	0	0	0
LINE	L1	L1Md_F2	chr2	95684500 95690724 -	0	0	0
SINE	Alu	B1_Mus2	chr2	96148225 96148370 -	0	0	0
LINE	L1	L1Md_T	chr2	96794845 96801163 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	96885365 96885408 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	96951977 96952012 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	97018336 97018377 +	0	0	0
LINE	L1	L1Md_F2	chr2	97095760 97101107 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	97121797 97121844 +	0	0	0
SINE	Alu	B1_Mus2	chr2	97123424 97123569 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	97123570 97123592 +	0	0	0
SINE	Alu	B1_Mus1	chr2	98737189 98737336 +	0	0	0
SINE	Alu	B1_Mm	chr2	98771918 98772064 -	0	0	0
SINE	Alu	B1_Mur2	chr2	99023465 99023602 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	99237849 99237877 +	0	0	0
SINE	Alu	B1_Mus2	chr2	99241251 99241397 +	0	0	0
SINE	B2	B3	chr11	5996739 5996914 -	1.77316	0	0.615324
SINE	Alu	B1_Mm	chr11	5996947 5997092 -	1.77316	0	0.615324
LTR	MaLR	MTA_Mm	chr2	10034703 10034742 +	0	0	0
LTR	MaLR	MTA_Mm	chr2	10034851 10034891 +	0	0	0
LTR	MaLR	MTA_Mm	chr2	10051354 10051393 +	0	0	0
LTR	MaLR	MTA_Mm	chr2	10051502 10051542 +	0	0.380568	0
SINE	Alu	B1_Mus1	chr2	10085343 10085357 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	10085357 10085359 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	10093980 10093983 +	0	0	0
LTR	MaLR	MTB	chr2	10176126 10176173 +	0	0	0
SINE	Alu	B1_Mm	chr2	10224328 10224343 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	10224343 10224345 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr11	6262050 6262091 +	0	0	0
SINE	Alu	B1_Mus2	chr11	6279738 6279884 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	10365180 10365184 +	0	0	0
SINE	Alu	B1_Mus2	chr11	6324646 6324792 -	0.591053	0	0.342623
Simple_repeat	Simple_repeat	(TTTA)n	chr2	10436202 10436210 +	0	0	0
Simple_repeat	Simple_repeat	(GAAAA)n	chr2	10439869 10439875 +	0	0	0
SINE	Alu	B1_Mus2	chr11	6355358 6355503 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	6355504 6355528 +	0	0	0
SINE	Alu	B1_Mm	chr2	10469850 10469864 -	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr2	10472345 10472352 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	10605853 10605856 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	10677446 10677449 +	0	0	0
SINE	Alu	B1_Mus1	chr2	10727745 10727759 +	0.591053	1.32017	0
Simple_repeat	Simple_repeat	(A)n	chr2	10727759 10727763 +	0.591053	1.32017	0.615324
Simple_repeat	Simple_repeat	(A)n	chr2	10812954 10812958 +	0	0	0
LINE	L1	L1Md_F2	chr2	10832956 10833486 +	0	0	0
Simple_repeat	Simple_repeat	(CAAAAA)n	chr2	10877913 10877926 +	0	0	0
LINE	L1	L1Md_F2	chr2	10892637 10893145 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr11	6688972 6689024 +	0.591053	0	0
LINE	L1	L1Md_F2	chr2	10940716 10941244 +	0	0	0
SINE	Alu	B1_Mm	chr2	10975618 10975633 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	10975633 10975635 +	0	0	0
SINE	Alu	B1_Mus2	chr2	11100152 11100166 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	11100167 11100169 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	11120432 11120434 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	11203186 11203189 +	0	0	0
SINE	B2	B2_Mm1a	chr2	11207661 11207680 +	0	0	0
LTR	MaLR	MTA_Mm	chr2	11229116 11229156 +	0	0	0
SINE	Alu	B1_Mus2	chr2	11290268 11290283 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	11290283 11290285 +	0	0	0

SINE	Alu	B1_Mus1	chr2	11518842 11518858 -	0	0	0
SINE	Alu	B1_Mus1	chr2	11565018 11565033 -	0	0	0
SINE	B2	B2_Mm1a	chr2	11593778 11593797 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	11593797 11593800 +	0.332194	0	0
SINE	Alu	B1_Mus2	chr2	11601715 11601730 -	0	0	0
SINE	Alu	B1_Mus1	chr2	11605683 11605697 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	11605697 11605700 +	0	0	0
Simple_repeat	Simple_repeat	(CCAA)n	chr2	11605700 11605702 +	0	0	0
LTR	MaLR	MTA_Mm	chr2	11627579 11627619 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	11648495 11648499 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	11664922 11664926 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	11667158 11667160 +	0	0	0
LTR	MaLR	MTB	chr2	11678037 11678076 -	0	0	0
SINE	Alu	B1_Mus2	chr11	7527922 7528068 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	11841873 11841877 +	0	0	0
SINE	Alu	B1_Mus2	chr2	11868528 11868543 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	11868543 11868546 +	0.591053	0	0
SINE	Alu	B1_Mus1	chr2	11893766 11893780 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	11893780 11893782 +	0	0	0
SINE	Alu	B1_Mus1	chr11	7690661 7690807 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	11931246 11931250 +	0	0	0
SINE	Alu	B1_Mus2	chr2	11998573 11998586 +	0	0	0
SINE	Alu	B1_Mus2	chr2	12011767 12011782 -	0	0	0
SINE	B2	B2_Mm1t	chr2	12050799 12050818 -	0	0	0
SINE	Alu	B1_Mm	chr2	12130431 12130446 +	0.591053	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr2	12205936 12205941 +	0	0	0
SINE	Alu	B1_Mus2	chr2	12229035 12229050 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	12229051 12229062 +	0	0	0
LINE	L1	L1Md_F2	chr2	12238879 12239505 -	0	0	0
SINE	Alu	B1_Mus2	chr2	12248636 12248651 -	0	0	0
SINE	Alu	B1_Mus1	chr2	12364606 12364621 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	12364623 12364629 +	0	0	0.342623
LTR	MaLR	MTA_Mm	chr2	12396140 12396177 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	12417551 12417555 +	0	0	0
SINE	Alu	B1_Mm	chr2	12420196 12420211 -	0	0	0
LTR	MaLR	MTA_Mm	chr2	12430002 12430041 +	0	0	0
SINE	Alu	B1_Mus2	chr2	12698296 12698311 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	12698311 12698314 +	0	0	0
SINE	Alu	B1_Mus2	chr2	12750883 12750897 -	0	0	0
SINE	Alu	B1_Mm	chr2	12851960 12851975 -	0	0	0
SINE	Alu	B1_Mus2	chr2	12863213 12863227 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	12863228 12863230 +	0	0	0
SINE	Alu	B1_Mus1	chr2	12897171 12897186 -	0	0	0.615324
SINE	Alu	B1_Mm	chr2	12898130 12898142 -	0	0	0
SINE	Alu	B1_Mus1	chr2	12898364 12898378 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	12898378 12898384 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	12901325 12901332 +	0	0	0
SINE	Alu	B1_Mm	chr2	12962339 12962353 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	12962353 12962356 +	0	0	0
SINE	Alu	B1_Mm	chr2	12980841 12980855 -	0	0	0
SINE	Alu	B1_Mus2	chr2	13052506 13052519 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	13052519 13052521 +	0	0	0
LTR	MaLR	MTA_Mm	chr2	13068516 13068555 +	0.591053	0	0
LTR	MaLR	MTA_Mm	chr2	13094768 13094808 -	0	0	0
LTR	MaLR	MTB	chr2	13140538 13140578 +	0	0	0
SINE	Alu	B1_Mur4	chr2	13199661 13199676 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	13199676 13199679 +	0	0	0
LTR	MaLR	MTA_Mm	chr2	13224549 13224588 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	9643899 9643933 +	0	0	0
LTR	MaLR	MTB	chr2	13249320 13249360 -	0	0	0
LINE	L1	L1_Mur1	chr2	13414770 13415271 +	0	0	0
SINE	Alu	B1_Mus2	chr2	13448096 13448110 -	0	0	0
LINE	L1	L1_Mus3	chr11	10018107 10023288 -	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr2	13624163 13624166 +	0	0	0
LTR	MaLR	MTA_Mm	chr2	13634418 13634458 -	0	0	0
SINE	Alu	B1_Mm	chr2	13649157 13649171 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	13649172 13649175 +	0	0	0



Simple_repeat	Simple_repeat	(A)n	chr2	13690005	13690009 +	0	0	0
SINE	Alu	B1_Mus2	chr2	13737192	13737207 +	0	0	0
SINE	Alu	B1_Mm	chr2	13740380	13740394 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	13740395	13740400 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	13748758	13748763 +	0	0	0
SINE	Alu	B1_Mus1	chr2	13811494	13811504 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	13815568	13815572 +	0	0	0
LTR	MaLR	MTA_Mm	chr11	10291426	10291815 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	13859401	13859407 +	0	0.708814	0
SINE	Alu	B1_Mus2	chr2	13877274	13877289 -	0	0	0
SINE	B2	B2_Mm1a	chr2	13993125	13993145 +	0	0	0
LTR	MaLR	MTA_Mm	chr11	10540093	10540488 -	0.591053	0	0
SINE	Alu	B1_Mus2	chr11	10550131	10550277 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	10550278	10550305 +	0	0.708814	0
SINE	Alu	B1_Mus2	chr2	14274505	14274520 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	14303284	14303288 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	14315218	14315221 +	0	0	0
LINE	L1	Lx	chr10	17612750	17617503 -	0.591053	1.41763	0.342623
SINE	Alu	B1_Mus2	chr2	14365152	14365166 -	0	0	0
SINE	Alu	B1_Mm	chr2	14402384	14402398 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr2	14402398	14402401 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	10910239	10910272 +	0	0	0
SINE	Alu	B1_Mm	chr2	14416566	14416581 -	0	0	0
SINE	Alu	B1_Mm	chr2	14417469	14417483 -	0	0	0
SINE	Alu	B1_Mus2	chr2	14423528	14423542 -	0	0	0
SINE	Alu	B1_Mus1	chr2	14424160	14424174 -	0	0	0
SINE	Alu	B1_Mus2	chr2	14497841	14497855 -	0	0	0
LTR	MaLR	MTA_Mm	chr2	14558792	14558832 -	0	0	0
SINE	Alu	B1_Mus1	chr2	14661634	14661651 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	14661651	14661655 +	0	0	0
SINE	Alu	B1_Mus2	chr2	14667110	14667125 -	0	0	0
LTR	MaLR	MTA_Mm	chr2	14761006	14761045 +	0	0	0
SINE	Alu	B1_Mur2	chr2	14779754	14779770 -	0	0	0
SINE	Alu	B1_Mm	chr2	14815542	14815557 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	14823546	14823550 +	0.332194	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr2	14846740	14846745 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	14938288	14938292 +	0	0	0
SINE	Alu	B1_Mus2	chr2	14941085	14941099 -	0	0	0.615324
SINE	Alu	B1_Mm	chr2	15011423	15011437 -	0	0	0
SINE	Alu	B1_Mm	chr2	15012728	15012742 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	15012742	15012746 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	11575224	11575262 +	0.332194	0	0
SINE	Alu	B1_Mm	chr2	15019380	15019395 -	0	0	0
SINE	Alu	B1_Mus2	chr2	15021270	15021285 -	0.591053	0	1.23065
SINE	Alu	B1_Mus2	chr2	15031425	15031439 -	0	0	0
SINE	Alu	B1_Mm	chr2	15032348	15032363 -	0.923247	1.08938	0
LINE	L1	L1Md_T	chr2	15032927	15033462 -	0	0	0
SINE	Alu	B1_Mm	chr2	15035541	15035556 -	0	0	0
SINE	Alu	B1_Mm	chr11	11678180	11678329 -	0	0	0
LTR	MaLR	MTA_Mm	chr2	15157071	15157110 +	0	0	0.342623
SINE	Alu	B1_Mm	chr11	11818442	11818587 -	0	0	0
SINE	Alu	B1_Mus2	chr2	15259138	15259151 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	15259152	15259155 +	0	0	0
SINE	Alu	B1_Mur3	chr2	15310696	15310707 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr2	15310707	15310714 +	0	0	0
Simple_repeat	Simple_repeat	(TTTTA)n	chr2	15341450	15341454 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	15341501	15341504 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr2	15344158	15344172 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	15344172	15344175 +	0	0	0
SINE	Alu	B1_Mus2	chr2	15353440	15353455 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	15392021	15392025 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	15412958	15412962 +	0	0	0
SINE	Alu	B1_Mus2	chr11	12416163	12416309 -	0	0	0
SINE	Alu	B1_Mm	chr2	15672103	15672118 -	0	0	0
SINE	Alu	B1_Mus1	chr2	15683367	15683381 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	15683381	15683384 +	0	0	0
SINE	Alu	B1_Mus2	chr2	15742319	15742333 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr2	15742333 15742336 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	15755182 15755188 +	0	0	1.10507
SINE	Alu	B1_Mm	chr2	15779858 15779873 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	15779873 15779876 +	0	0	0
LTR	MaLR	MTA_Mm	chr11	12807604 12807999 -	0	0	0
LTR	MaLR	MTA_Mm	chr11	12957698 12958090 +	0	0	0.615324
SINE	Alu	B1_Mus1	chr2	15921201 15921216 -	0	0	0
SINE	Alu	B1_Mus2	chr11	13068885 13069030 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	13069031 13069058 +	0	0	0
SINE	Alu	B1_Mm	chr2	16013381 16013396 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	16035486 16035490 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	16036974 16036977 +	0	0	0
SINE	Alu	B1_Mus1	chr2	16040315 16040333 -	0	0	0
LTR	MaLR	MTA_Mm	chr10	3125491 3125864 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	16105492 16105496 +	0	0	0
SINE	Alu	B1_Mm	chr2	16108704 16108719 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	16108719 16108725 +	0	0	0
SINE	Alu	B1_Mus2	chr11	13355751 13355897 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	16261470 16261473 +	0	0	0
SINE	Alu	B1_Mus1	chr11	13454360 13454507 -	0	0	0
SINE	Alu	B1_Mus1	chr2	16344712 16344727 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	16344727 16344731 +	0	0	0
SINE	Alu	B1_Mus1	chr2	16347211 16347226 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	16347226 16347229 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	13591820 13591873 +	0	0	0
SINE	Alu	B1_Mm	chr2	16451653 16451668 -	0	0	0
LTR	MaLR	MTA_Mm	chr11	13941232 13941570 +	0	0	0
SINE	Alu	B1_Mm	chr2	16616066 16616080 -	0	0	0
SINE	Alu	B1_Mus2	chr2	16638083 16638098 -	0	0	0
SINE	Alu	B1_Mur4	chr2	16659387 16659400 -	0	0	0
SINE	Alu	B1_Mm	chr2	16659760 16659775 -	0	0	0
SINE	Alu	B1_Mus2	chr2	16661371 16661385 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	16702301 16702303 +	0.923247	0	0.615324
SINE	Alu	B1_Mus2	chr2	16722073 16722088 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	16722088 16722090 +	0	0	0
SINE	Alu	B1_Mus2	chr2	16746072 16746088 -	0	0	0
SINE	Alu	B1_Mus1	chr2	16750737 16750751 -	0	0	0
SINE	Alu	B1_Mur1	chr2	16807432 16807447 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	16813283 16813285 +	0	0	0
SINE	Alu	B1_Mus2	chr2	16813707 16813721 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	16813721 16813724 +	0	0	0
LINE	L1	L1_Mus2	chr11	14359886 14365793 -	0	0	0
SINE	B2	B2_Mm1a	chr2	16861489 16861509 -	0	0	0
SINE	Alu	B1_Mm	chr2	16865160 16865173 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	16865173 16865175 +	0	0	0
SINE	B2	B2_Mm1a	chr2	16873979 16873998 -	0	0.380568	0
SINE	Alu	B1_Mus1	chr2	16878874 16878889 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	16980293 16980300 +	0	0	0
SINE	Alu	B1_Mus2	chr2	16996857 16996871 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	17029551 17029554 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	14699506 14699544 +	0	0	0
LTR	MaLR	MTA_Mm	chr2	17180948 17180988 +	0	0	0
SINE	Alu	B1_Mus2	chr2	17191143 17191158 +	0.664388	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	17191158 17191160 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	17199403 17199407 +	0	0	0
SINE	Alu	B1_Mus2	chr2	17214545 17214560 -	0	0	0
SINE	Alu	PB1D10	chr2	17248138 17248149 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	17248150 17248154 +	0	0	0
SINE	B2	B2_Mm1a	chr2	17279389 17279408 +	0	0	0
SINE	B4	ID_B1	chr2	17291539 17291560 -	0	0	0
LTR	MaLR	MTA_Mm	chr2	17370505 17370543 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	15149916 15149951 +	0	0	0
SINE	Alu	B1_Mm	chr2	17407601 17407615 +	0	0	0
SINE	Alu	B1_Mm	chr2	17458461 17458475 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	17458475 17458477 +	0	0	0
LTR	MaLR	MTA_Mm	chr2	17473586 17473624 +	0	0	0
SINE	Alu	B1_Mus1	chr2	17488092 17488106 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr2	17488106 17488110 +	0	0	0
SINE	Alu	B1_Mus2	chr2	17548966 17548981 -	0	0	0
SINE	Alu	B1_Mus2	chr2	17550934 17550949 +	0.591053	0	0
SINE	Alu	B1_Mus2	chr11	15371734 15371880 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	15371881 15371903 +	0	0	0
SINE	Alu	B1_Mus1	chr2	17687749 17687763 -	0	0	0.615324
SINE	Alu	B1_Mus2	chr2	17713380 17713394 +	0	0	0
SINE	Alu	B1_Mus2	chr2	17720426 17720441 -	0	0	0
SINE	Alu	B1_Mus2	chr2	17745319 17745334 -	0	0	0
SINE	Alu	B1_Mus2	chr2	17747260 17747274 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	17747274 17747278 +	0	0	0
SINE	Alu	B1_Mus2	chr2	17753013 17753027 +	0	0	0
SINE	Alu	B1_Mus2	chr2	17758654 17758668 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	17767373 17767377 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	15628316 15628362 +	0	0	0
LINE	L1	L1Md_F2	chr2	17888592 17889132 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr2	17903889 17903892 +	0	0	0
LTR	MaLR	MTA_Mm	chr2	17905205 17905244 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	15765076 15765127 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	15793213 15793251 +	0	0	0
SINE	Alu	B1_Mus2	chr2	18045409 18045423 -	0	0	0
SINE	B2	B2_Mm1t	chr2	18143466 18143486 +	0	0	0
LTR	MaLR	MTA_Mm	chr10	4599525 4599922 -	0	0	0
LTR	MaLR	MTA_Mm	chr10	18002760 18003137 -	0	0	0
LTR	MaLR	MTA_Mm	chr3	3678720 3679115 -	0	0	0
LINE	L1	L1Md_T	chr3	4164192 4170853 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	4447388 4447436 +	0	0	0
LTR	MaLR	MTA_Mm	chr11	16225227 16225624 -	0	0	0
LINE	L1	L1_Mus1	chr3	6188692 6194794 +	0.591053	0	0
SINE	Alu	B1_Mus2	chr3	6219439 6219585 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	6219586 6219620 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	6265004 6265398 -	0	0	0
SINE	Alu	B1_Mus2	chr3	6436386 6436531 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr3	6436532 6436624 +	0	0	0
SINE	Alu	B1_Mus2	chr3	6760274 6760418 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	6760419 6760499 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	6785486 6785537 +	0	0	0
SINE	Alu	B1_Mus2	chr3	6957704 6957812 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	6957813 6957844 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	6959282 6959328 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	7010425 7010470 +	0	0	0
SINE	Alu	B1_Mm	chr11	16437726 16437869 -	0	0	0
SINE	Alu	B1_Mus1	chr3	8232960 8233105 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	8233106 8233130 +	0	0	0
SINE	Alu	B1_Mus1	chr3	8245571 8245718 +	0	0	0
Simple_repeat	Simple_repeat	(CAAAAC)n	chr3	8245747 8245769 +	0.591053	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr11	16477999 16478036 +	0	0	0
LINE	L1	L1Md_F2	chr3	8330527 8335703 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	8651898 8652278 +	0	0	0
SINE	Alu	B1_Mus2	chr3	9011506 9011617 +	0	0	0
SINE	Alu	B1_Mur4	chr3	9241405 9241544 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	9241545 9241603 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	9416687 9417072 -	0	0.380568	0
LTR	MaLR	MTA_Mm	chr3	10073766 10074161 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	10097278 10097305 +	0	0	0
SINE	Alu	B1_Mus1	chr3	10134119 10134264 -	0	0	0
SINE	Alu	B1_Mus2	chr3	10151987 10152132 -	0	0	0
SINE	Alu	B1_Mus1	chr3	10824527 10824671 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	10824672 10824703 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	11012719 11013112 +	0	0	0
LINE	L1	L1Md_F2	chr3	11367351 11372700 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	11615025 11615049 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	11729266 11729329 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	12113117 12113504 -	0	0	0
LINE	L1	L1Md_F2	chr3	12619618 12625330 +	0	0	0
SINE	Alu	B1_Mus1	chr3	12672844 12672990 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	12672991 12673014 +	0	0	0

LTR	MaLR	MTA_Mm	chr3	12887285 12887679 -	0	0	0
SINE	Alu	B1_Mus1	chr3	12941557 12941703 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	13352572 13352607 +	0	0.380568	0
LTR	MaLR	MTA_Mm	chr3	15189734 15190130 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	15693803 15693843 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	15752839 15752924 +	0	0	0.615324
LINE	L1	L1Md_T	chr3	15822545 15829064 -	0	0	0
SINE	Alu	B1_Mm	chr3	16043126 16043272 +	0	0	0
LTR	ERVVK	RMER13A	chr3	16762314 16763095 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	16881572 16881639 +	0.827951	0	0
LINE	L1	L1Md_F2	chr3	17413654 17419583 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	17810628 17811014 +	0	0	0
SINE	Alu	B1_Mm	chr3	18262571 18262717 -	0	0	0
LINE	L1	L1_Mus1	chr3	18304422 18310872 +	0	0	0
Low_complexity	Low_complexity	CT-rich	chr3	19314608 19314658 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	19702274 19702657 +	0	0	0
LTR	MaLR	MTA_Mm	chr10	18099028 18099424 -	0	0	0
LTR	MaLR	MTA_Mm	chr3	20196444 20196838 -	0	0	0
SINE	Alu	B1_Mm	chr3	20461059 20461205 -	0	0	0
LTR	MaLR	MTA_Mm	chr3	20688518 20688890 -	0	0	0
SINE	Alu	B1_Mus2	chr3	20805125 20805272 -	0	0	0
SINE	Alu	B1_Mm	chr3	20950943 20951099 +	1.18211	0.708814	0.342623
Simple_repeat	Simple_repeat	(A)n	chr3	20951100 20951123 +	1.18211	0	0
LTR	MaLR	MTEa	chr3	21188701 21189042 +	0	0	0
SINE	B4	B4	chr3	21594385 21594677 +	0	0	0
SINE	Alu	B1_Mus1	chr3	21778822 21778969 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	21778970 21778993 +	0	0	0
SINE	Alu	B1_Mus2	chr3	21801481 21801627 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	22120588 22120649 +	0.591053	0.708814	0
LINE	L1	L1Md_A	chr3	22347096 22352613 -	0	0	0
LINE	L1	L1Md_T	chr3	22352630 22359509 -	0.591053	0	0
LINE	L1	L1Md_F2	chr3	22588804 22595023 -	0	0	0
SINE	B2	B2_Mm1t	chr11	17875355 17875546 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	17875547 17875583 +	0.332194	0	1.4477
LINE	L1	L1Md_T	chr3	22906497 22914001 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	23742220 23742244 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	23803971 23804364 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	24179577 24179610 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	24232475 24232509 +	18.3018	11.2056	11.6282
Simple_repeat	Simple_repeat	(A)n	chr3	24446038 24446100 +	0	0	0
SINE	Alu	B1_Mus2	chr3	24472456 24472602 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	24472603 24472626 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	24585102 24585496 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	24808479 24808508 +	0	0	0
SINE	Alu	B1_Mus2	chr3	27755964 27756110 -	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr3	28041748 28041778 +	0	0	0
SINE	Alu	B1_Mus1	chr11	18518995 18519133 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	28639858 28639896 +	0	0	0
LINE	L1	L1Md_T	chr3	29765054 29771712 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	29827882 29827902 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	30413765 30413794 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	30462654 30462977 +	0	0	0
SINE	Alu	B1_Mus1	chr3	30796189 30796326 -	0.591053	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr3	31509047 31509094 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	31738383 31738778 +	0	0	0
SINE	Alu	B1_Mus2	chr3	32294676 32294812 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	32294813 32294838 +	0	0	0
SINE	Alu	B1_Mus1	chr3	33487701 33487849 -	0	0	0
SINE	Alu	B1_Mus2	chr3	34617241 34617387 +	0	0	0
SINE	Alu	B1_Mus1	chr3	34940470 34940616 -	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	34977271 34977331 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	36246418 36246449 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	36656824 36657205 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	36658297 36658678 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	37481266 37481671 +	0	0	0.615324
LTR	MaLR	MTA_Mm	chr10	18284560 18284939 +	0	0	0
SINE	Alu	B1_Mus1	chr3	38764013 38764163 -	0	0	0

SINE	Alu	B1_Mus2	chr3	38769792 38769937 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	39045673 39045699 +	0	0	0
SINE	Alu	B1_Mm	chr3	39303477 39303622 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	39390451 39390485 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	40861330 40861694 +	0	0	0
LINE	L1	L1Md_F2	chr3	41594751 41599914 +	0	0.708814	0.342623
LTR	MaLR	MTA_Mm	chr3	41655818 41656212 -	0.332194	0	0
SINE	Alu	B1_Mm	chr11	19931275 19931419 -	0	0	0
SINE	Alu	B1_Mus2	chr11	19955126 19955272 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	42986291 42986334 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	43098577 43098642 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	43159367 43159402 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	43515394 43515788 +	0.332194	0.708814	0.866667
Simple_repeat	Simple_repeat	(A)n	chr3	43787145 43787175 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	44133411 44133438 +	0	0	0
SINE	Alu	B1_Mur3	chr11	20044190 20044337 -	0.591053	0	0
SINE	Alu	B1_Mus2	chr3	44555441 44555595 -	0	0	0
LINE	L1	L1Md_F2	chr3	44730975 44737160 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	44851511 44851561 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	44991437 44991478 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	45314976 45315367 -	0	0	0
LTR	MaLR	MTA_Mm	chr3	45321646 45322040 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	45468770 45468813 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	46114710 46114744 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	46172366 46172410 +	0	0	0
SINE	Alu	B1_Mus2	chr3	46254464 46254606 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	46670281 46670378 +	0.332194	0	0
SINE	Alu	B1_Mus1	chr3	46801879 46802023 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	46802024 46802046 +	0	0	0
SINE	Alu	B1_Mus1	chr3	47001791 47001937 -	0	0	0
LINE	L1	L1Md_F3	chr3	47105820 47112010 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	47266569 47266609 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	47453364 47453736 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	47460138 47460533 -	0	0	0
LTR	MaLR	MTA_Mm	chr3	47601653 47602046 -	0	0	0
LTR	MaLR	MTA_Mm	chr3	47603143 47603536 -	0	0	0
LTR	ERV1	RLTR24	chr11	20269083 20269531 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	49129863 49130256 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	49131308 49131701 +	0	0	0
LTR	MaLR	MTA_Mm	chr11	20321657 20322048 +	0	0	0
SINE	Alu	B1_Mus2	chr11	20324806 20324951 +	0	0	0
SINE	Alu	B1_Mus1	chr11	20374898 20375040 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	20375041 20375068 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	50653260 50653301 +	0	0	0
SINE	Alu	B1_Mus2	chr11	20408776 20408922 -	0	0	0
SINE	Alu	B1_Mus2	chr3	51928899 51929044 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	51929045 51929067 +	0	0	0
SINE	Alu	B1_Mus2	chr3	52017487 52017634 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	52017635 52017671 +	0	0	0
SINE	B2	B2_Mm1a	chr3	52279434 52279625 -	0	0	0
SINE	Alu	B1_Mm	chr3	53165048 53165194 +	0	0	0
LINE	L1	L1Md_T	chr3	53519032 53525883 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	53603103 53603140 +	0	0	0
SINE	Alu	B1_Mus2	chr3	53619756 53619900 -	0	0	0
SINE	Alu	B1_Mus2	chr3	53786002 53786149 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	53786150 53786172 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr3	53892577 53892665 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	53906636 53906685 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	54253940 54254334 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	54391179 54391213 +	0	0	0
Simple_repeat	Simple_repeat	(CAAAC)n	chr11	20837715 20837742 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	55991948 55991982 +	0.591053	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr3	56058482 56058509 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	56129641 56130037 +	0	0	0
LINE	L1	L1Md_F2	chr3	56352275 56357275 -	0	0	0
SINE	Alu	B1_Mus2	chr3	57174380 57174526 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	57174527 57174569 +	0	0	0.342623

SINE	Alu	B1_Mus2	chr3	57381654 57381800 -	0	0	0
SINE	Alu	B1_Mus2	chr3	57387061 57387207 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	57387208 57387231 +	0	0	0
LTR	MaLR	MTA_Mm	chr10	4653854 4654253 +	0	0	0
SINE	Alu	B1_Mm	chr3	58279183 58279327 +	0	0	0
LINE	L1	L1Md_F2	chr3	58404445 58409468 +	0.591053	0.708814	0.615324
SINE	Alu	B1_Mus1	chr11	21128285 21128430 -	0	0	0
LTR	MaLR	MTA_Mm	chr3	59510019 59510414 -	0	0	0
LTR	MaLR	MTA_Mm	chr3	59511504 59511895 -	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr10	18507499 18507542 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	59781336 59781731 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	60541979 60542018 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	61042411 61042805 -	0	0	0
LTR	MaLR	MTA_Mm	chr3	61043900 61044294 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	61223706 61223748 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	61472475 61472509 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	62439590 62439985 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr3	62725129 62725271 -	0	0	0
SINE	Alu	B1_Mus2	chr3	62911832 62911978 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	63666743 63666769 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	64292471 64292553 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	64370740 64370783 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	65385408 65385444 +	0	0	0
SINE	Alu	B1_Mur3	chr3	65822832 65822982 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	65822983 65823006 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	66406979 66407005 +	0	0	0
SINE	Alu	B1_Mus2	chr3	67673916 67674063 -	0	0	0
SINE	Alu	B1_Mm	chr3	68517389 68517505 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	68517506 68517531 +	0	0	0
SINE	Alu	B1_Mus2	chr3	68603190 68603335 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	68603336 68603362 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	68636270 68636665 +	0	0	0
SINE	Alu	B1_Mur3	chr3	68668723 68668863 -	1.85701	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	68864467 68864493 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr3	69598593 69598712 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	69804905 69805299 +	0	0	0.957947
LTR	MaLR	MTA_Mm	chr11	22205745 22206141 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	70250824 70251217 -	0	0	0
LTR	MaLR	MTA_Mm	chr3	70518923 70519314 -	0	0	0
SINE	Alu	B1_Mus2	chr3	70697709 70697855 -	0	0	0
LTR	MaLR	MTA_Mm	chr11	22339591 22339985 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	71437018 71437057 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	71462058 71462086 +	0	0	0
LINE	L1	L1Md_F2	chr3	71481041 71486238 +	0	0	0
LINE	L1	L1Md_F	chr3	71547793 71553709 -	0	0	0
SINE	Alu	B1_Mus1	chr3	72260310 72260457 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	72260458 72260487 +	0	0	0
SINE	Alu	B1_Mus1	chr3	72309699 72309858 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	72309859 72309966 +	0	0	0
SINE	Alu	B1_Mm	chr3	72344232 72344381 -	0	0	0
SINE	Alu	B1_Mm	chr11	22423833 22423979 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	72636435 72636463 +	0	0	0
SINE	Alu	B1_Mm	chr11	22493129 22493275 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	22509075 22509113 +	0	0	0
SINE	Alu	B1_Mur1	chr11	22572779 22572916 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	22572917 22572950 +	0	0	0
SINE	Alu	B1_Mus2	chr3	74512229 74512368 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	74512369 74512409 +	0	0	0
LTR	MaLR	MTA_Mm	chr11	22600151 22600545 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	74786161 74786211 +	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr3	75042640 75042696 +	0	0	0
SINE	Alu	B1_Mus2	chr3	75296636 75296781 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	76740180 76740228 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	76983842 76983885 +	0	0	0
LINE	L1	L1Md_T	chr3	76987954 76994367 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	77031435 77031477 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	77079903 77079933 +	0	0	0

LTR	MaLR	MTA_Mm	chr3	77355528 77355926 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	77773004 77773399 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	77924322 77924717 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	78066947 78066982 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	78312684 78312723 +	0	0	0
LINE	L1	L1Md_F3	chr3	78370846 78376348 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	78425518 78425546 +	0	0	0
SINE	Alu	B1_Mus1	chr3	78573176 78573323 -	0	0	0.342623
LTR	MaLR	MTA_Mm	chr3	78955450 78955845 +	0	0	0
SINE	Alu	B1_Mus1	chr3	79759513 79759660 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	79759661 79759686 +	0.591053	0	0
LTR	MaLR	ORR1B2	chr3	79930865 79931157 -	0	0	0
LINE	L1	L1Md_F2	chr3	79947917 79954042 -	0	0	0
LTR	MaLR	MTA_Mm	chr3	80238168 80238563 -	0	0	0
LTR	MaLR	MTA_Mm	chr3	80325463 80325857 -	0	0	0
LTR	MaLR	MTA_Mm	chr3	80326953 80327347 -	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr3	80741004 80741047 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	81050701 81050731 +	0.332194	0	0
LINE	L1	L1_Mus4	chr3	81315832 81321762 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	82298085 82298118 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	82568375 82568768 +	0	0	0
SINE	Alu	B1_Mm	chr3	83020239 83020384 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	83020385 83020408 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	83206164 83206234 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr3	84451988 84452133 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	84452134 84452161 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	84589292 84589325 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	85312988 85313026 +	0	0	0
SINE	B4	B4A	chr3	85359870 85360177 -	3.15698	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	85582048 85582101 +	0	0	0
SINE	Alu	B1_Mm	chr3	85745970 85746116 -	0	0	0
SINE	Alu	B1_Mus1	chr3	85790414 85790561 +	1.77316	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	85790562 85790586 +	1.77316	0	0
SINE	Alu	B1_Mus2	chr3	85967221 85967357 -	0	0	0
SINE	Alu	B1_Mus2	chr3	86799109 86799255 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	86799256 86799284 +	0	0	0
SINE	Alu	B1_Mur4	chr3	87690071 87690217 +	0	0.380568	0.342623
Simple_repeat	Simple_repeat	(A)n	chr3	87690218 87690247 +	0	0.380568	0.342623
SINE	B2	B2_Mm1a	chr3	87761374 87761561 +	0	0	0
SINE	Alu	B1_Mm	chr3	87818182 87818328 -	0	0	0
SINE	B2	B2_Mm1a	chr11	23402039 23402232 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	23402233 23402262 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr11	23407488 23407634 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	23407635 23407662 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	88071665 88072046 -	0	0	0
SINE	Alu	B1_Mm	chr11	23416000 23416145 -	0	0	0
SINE	Alu	B1_Mus1	chr3	88282921 88283065 -	0	0	0
SINE	Alu	B1_Mm	chr3	88378663 88378809 -	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr3	88414309 88414339 +	0	0	0
SINE	Alu	B1_Mus2	chr3	88760857 88760989 -	0	0	0
SINE	Alu	B1_Mus1	chr3	88886516 88886660 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	88916988 88917017 +	0	0	0
SINE	Alu	B1_Mus2	chr3	89004305 89004451 -	0	0	0
SINE	B2	B2_Mm1a	chr3	89259677 89259834 -	0	0	0
SINE	Alu	B1_Mus2	chr3	89290497 89290613 -	0	0	0
SINE	Alu	B1_Mus1	chr3	89496393 89496533 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	89496534 89496555 +	0	0	0
SINE	Alu	B1_Mm	chr3	89787240 89787386 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	89787387 89787437 +	0	0	0.342623
SINE	Alu	B1_Mm	chr3	90023725 90023878 -	0	0	0
SINE	Alu	B1_Mus2	chr3	90087665 90087810 +	0	0	0
SINE	Alu	B1_Mm	chr3	90091530 90091679 -	0	0	0
SINE	Alu	B1_Mus2	chr3	90093301 90093446 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	90093447 90093479 +	0	0	0
SINE	Alu	B1_Mus1	chr3	90251584 90251730 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	90251731 90251759 +	0	0	0
SINE	Alu	B1_Mus2	chr3	90274991 90275137 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr3	90275138 90275169 +	0	0	0
SINE	Alu	B1_Mus2	chr3	90279296 90279448 +	0	0	0
LTR	MaLR	MTA_Mm	chr11	23689081 23689480 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	90663576 90663611 +	0	0	0
LINE	L1	L1Md_F3	chr3	90971295 90975562 +	0	0	0.342623
SINE	Alu	B1_Mm	chr11	23719512 23719657 -	0	0	0
LINE	L1	L1_Mus2	chr3	91279558 91285114 -	0	0	0
LINE	L1	L1Md_F2	chr3	91459705 91465170 +	0	0	0
LINE	L1	L1Md_F2	chr3	91559343 91564778 +	0	0	0
LINE	L1	L1_Mus1	chr3	91721211 91726913 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	91808850 91808874 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	91818084 91818124 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	92374727 92374763 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	92602954 92603348 -	0	0	0
LTR	MaLR	MTA_Mm	chr3	93104132 93104526 +	0	0	0
SINE	Alu	B1_Mm	chr3	93253193 93253338 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	93253339 93253362 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	23906926 23906967 +	0	0	0
SINE	Alu	B1_Mus2	chr3	93287123 93287268 -	0	0	0
SINE	Alu	B1_Mus1	chr3	94086204 94086321 -	0	0	0
SINE	Alu	B1_Mus2	chr3	94395714 94395860 -	0	0	0
SINE	Alu	B1_Mus1	chr3	94412768 94412900 -	0	0	0
SINE	Alu	B1_Mus1	chr3	94526435 94526581 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	94526582 94526606 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	94861273 94861668 +	0	0	0
SINE	Alu	B1_Mus1	chr3	95112253 95112400 -	0	0	0
SINE	Alu	B1_Mm	chr3	95230785 95230932 -	0	0	0
SINE	Alu	B1_Mur2	chr3	95315534 95315673 +	0	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr3	95315674 95315699 +	0	0.380568	0
SINE	Alu	B1_Mus1	chr3	95524682 95524830 -	0	0	0
SINE	Alu	B1_Mus1	chr3	95630185 95630328 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	95630329 95630367 +	0	0	0
SINE	Alu	B1_Mus1	chr3	95764019 95764156 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	95764157 95764196 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr3	96021706 96021757 +	0	0	0
SINE	Alu	B1_Mm	chr3	96356292 96356438 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	96356439 96356473 +	0	0	0
SINE	B2	B2_Mm1t	chr3	96889878 96890071 +	0	0	0
SINE	Alu	B1_Mus2	chr3	96926642 96926788 +	0	0	0
SINE	Alu	B1_Mm	chr3	96943885 96944030 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	96944031 96944080 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	97046455 97046483 +	0	0	0
SINE	Alu	B1_Mus2	chr3	97163963 97164108 -	0	0	0
SINE	Alu	B1_Mm	chr3	97421293 97421407 -	0	0	0
LINE	L1	L1Md_T	chr11	24775660 24782253 -	0	0	0
SINE	Alu	B1_Mm	chr3	97997970 97998115 -	0	0	0
SINE	Alu	B1_Mm	chr3	98146680 98146826 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	98146827 98146854 +	0	0	0
SINE	Alu	B1_Mus2	chr3	98163720 98163866 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	98163867 98163888 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	24851086 24851114 +	0	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr3	99043110 99043136 +	0	0	0
LINE	L1	L1Md_F	chr3	99170363 99175652 +	0	0	0
LINE	L1	L1Md_F2	chr3	99410060 99416420 -	0	0	0
SINE	Alu	PB1D9	chr3	10051845 10051857 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	10051857 10051861 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	10056637 10056676 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	10084335 10084372 -	0	0	0
SINE	Alu	B1_Mm	chr3	10183070 10183085 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	10183085 10183091 +	0	0	0
SINE	Alu	B1_Mus2	chr3	10188406 10188421 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	10305795 10305798 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	10349632 10349672 -	0	0	0
SINE	Alu	B1_Mm	chr3	10350811 10350825 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	10350826 10350829 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	10381285 10381287 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	10414024 10414027 +	0	0.380568	0



SINE	Alu	B1_Mus2	chr3	10434372	10434387 -	0.332194	0	0
SINE	Alu	B1_Mus2	chr3	10446882	10446897 -	0	0	0
SINE	Alu	PB1D7	chr3	10489134	10489144 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	10489144	10489147 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	10498754	10498794 -	0	0	0
SINE	Alu	B1_Mur4	chr3	10549318	10549333 +	0	0	0
SINE	Alu	B1_Mus2	chr3	10563959	10563973 -	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr3	10565222	10565225 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	10573919	10573923 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	10646522	10646561 -	0	0	0
LTR	MaLR	MTA_Mm	chr3	10646667	10646706 -	0	0	0
SINE	Alu	B1_Mus1	chr3	10714247	10714262 -	0.332194	0	0
LTR	MaLR	MTA_Mm	chr11	26141137	26141531 -	0	0	0
SINE	Alu	B1_Mm	chr3	10771508	10771520 -	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr3	10781115	10781119 +	0	0	0
SINE	Alu	B1_Mus2	chr3	10804344	10804359 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	10804359	10804362 +	0	0	0
SINE	Alu	B1_Mus1	chr3	10819293	10819307 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	10819307	10819315 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	11026376	11026415 +	0	0	0
LINE	L1	L1Md_F2	chr3	11063086	11063701 -	0	0	0
SINE	Alu	B1_Mus1	chr3	11080280	11080294 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	11080295	11080297 +	0	0	0
SINE	Alu	B1_Mm	chr3	11089918	11089933 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	11089933	11089936 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	11091517	11091557 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	11091663	11091702 +	0	0	0
LTR	MaLR	MTA_Mm	chr11	26562613	26562945 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	11138073	11138078 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	11194867	11194906 +	0	0	0
SINE	Alu	B1_Mm	chr11	26677502	26677648 -	0	0	0
LTR	MaLR	MTA_Mm	chr3	11227044	11227083 +	0	0	0
LINE	L1	L1_Mus2	chr3	11233136	11233587 -	0	0	0
LINE	L1	L1Md_F2	chr3	11233620	11234156 +	0	0	0
LINE	L1	Lx2	chr3	11280539	11281039 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	11283115	11283154 -	0	0	0
SINE	Alu	B1_Mus2	chr11	26755993	26756139 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	26756140	26756176 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	11407001	11407041 -	0	0	0.615324
Simple_repeat	Simple_repeat	(GAAA)n	chr3	11422482	11422492 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	11496681	11496720 +	0	0	0.342623
LINE	L1	L1Md_T	chr3	11512345	11512881 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	11520039	11520043 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	11525787	11525791 +	0	0	0
SINE	Alu	B1_Mus1	chr3	11555645	11555659 -	0	0	0
SINE	Alu	B1_Mus2	chr3	11576718	11576733 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	11576733	11576736 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	11576899	11576939 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	27088440	27088482 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	11583851	11583891 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr3	11596080	11596083 +	0	0	0
LINE	L1	L1Md_T	chr11	27282597	27289454 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	11782868	11782872 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	11783752	11783756 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	11788897	11788902 +	0	0	0
SINE	Alu	B1_Mur4	chr3	11918661	11918673 -	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	11920994	11920997 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	11938772	11938776 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	11988289	11988292 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	12003681	12003721 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	12027021	12027024 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	12049579	12049582 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	12053758	12053762 +	0	0	0
Simple_repeat	Simple_repeat	(GAAAA)n	chr3	12059692	12059700 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	12065403	12065406 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	12089977	12090016 -	0	0	0
SINE	B2	B2_Mm1a	chr3	12126736	12126756 +	0	0	0

LTR	MaLR	MTB	chr10	4726252	4726647	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	12197672	12197675	+	0	0	0.342623
SINE	Alu	B1_Mus1	chr3	12239103	12239117	-	0	0	0
SINE	B2	B2_Mm1t	chr3	12263111	12263129	-	0	0	0
SINE	Alu	B1_Mus1	chr3	12269834	12269849	-	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chr11	27956130	27956231	+	0	0	0
LTR	MaLR	MTA_Mm	chr3	12387264	12387304	+	0	0	0
SINE	Alu	B1_Mus1	chr3	12415655	12415671	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	12415671	12415673	+	0	0	0
SINE	Alu	B1_Mus2	chr3	12419654	12419669	-	0	0.380568	0
Simple_repeat	Simple_repeat	(GAAA)n	chr3	12500902	12500911	+	0	0	0
LINE	L1	L1Md_F2	chr3	12507730	12508327	-	0	0	0
SINE	Alu	B1_Mus1	chr3	12554561	12554575	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	12554575	12554578	+	0.332194	0	0
LTR	MaLR	MTA_Mm	chr3	12582853	12582891	-	0	0	0
SINE	Alu	B1_Mus2	chr3	12660619	12660633	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	12660633	12660636	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	12684337	12684341	+	0	0	0
SINE	Alu	B1_Mm	chr3	12691754	12691769	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	12691769	12691771	+	0	0	0
SINE	Alu	B1_Mm	chr3	12692420	12692434	-	2.89812	1.85052	5.26521
Simple_repeat	Simple_repeat	(A)n	chr3	12692824	12692828	+	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	12696025	12696028	+	0	0	0
SINE	B2	B2_Mm1a	chr3	12714560	12714579	+	0	0	0
SINE	Alu	B1_Mm	chr3	12722940	12722954	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	12796103	12796105	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	12813499	12813504	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	28610173	28610202	+	0	0	0
SINE	Alu	B1_Mus1	chr3	12827113	12827127	-	0	0	0
SINE	Alu	B1_Mm	chr3	12830543	12830557	-	0	0	0
SINE	Alu	B1_Mus2	chr3	12833169	12833184	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	12833184	12833188	+	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	12844478	12844482	+	0	0	0
SINE	Alu	B1_Mus2	chr3	12868165	12868180	-	0	0	0
SINE	Alu	B1_Mus2	chr3	12871180	12871194	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	12871195	12871200	+	0	0	0
SINE	Alu	B1_Mm	chr3	12912458	12912472	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	19229015	19229043	+	0	0	0
LTR	MaLR	MTB	chr3	12947980	12948019	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	28856085	28856119	+	0	0	0
SINE	Alu	B1_Mus1	chr3	13043430	13043445	+	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	13043445	13043450	+	0.591053	0	0
SINE	Alu	B1_Mus2	chr3	13043610	13043624	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	13043625	13043627	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	13051636	13051644	+	0	0	0
SINE	Alu	B1_Mus1	chr3	13061362	13061378	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	13065879	13065884	+	0	0	0.342623
LINE	L1	L1Md_T	chr3	13136962	13137490	+	0	0	0
SINE	Alu	B1_Mm	chr3	13169254	13169268	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	13193610	13193613	+	0	0	0
SINE	Alu	B1_Mus1	chr3	13231687	13231701	-	0	0	0
SINE	Alu	B1_Mm	chr3	13250226	13250240	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	13294923	13294929	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	13322054	13322057	+	0	0	0
Low_complexity	Low_complexity	AT_rich	chr3	13411876	13411880	+	0	0	0
SINE	Alu	B1_Mus2	chr3	13449106	13449121	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	13449121	13449123	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	13470858	13470861	+	0	0	0
SINE	Alu	B1_Mm	chr10	19287667	19287813	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	19287814	19287838	+	0	0	0
SINE	Alu	B1_Mm	chr3	13548560	13548573	-	0	0	0
SINE	Alu	PB1D9	chr3	13620480	13620491	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	13620491	13620493	+	0	0	0
SINE	Alu	B1_Mus2	chr3	13691754	13691769	-	0	0	0
SINE	Alu	B1_Mm	chr11	29497558	29497704	+	0	0	0
SINE	Alu	B1_Mus1	chr3	13752482	13752494	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	13752495	13752497	+	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr3	13766074	13766078 +	0	0	0
SINE	B2	B2_Mm1a	chr11	29566743	29566934 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	13875111	13875149 +	0	0	0
SINE	Alu	B1_Mm	chr3	14092979	14092993 -	0	0	0
SINE	Alu	B1_Mus1	chr3	14100428	14100443 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr3	14100443	14100454 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	14184006	14184045 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr3	14184241	14184245 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	14272344	14272383 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	14377402	14377406 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	14379113	14379117 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	14398471	14398475 +	0	0	0
SINE	Alu	B1_Mus2	chr3	14431410	14431424 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	14442605	14442608 +	0	0	0
SINE	Alu	B1_Mus1	chr3	14521735	14521749 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	14521749	14521752 +	0	0	0
SINE	Alu	B1_Mus2	chr3	14587986	14588000 -	0	0	0
SINE	Alu	B1_Mus2	chr3	14618538	14618553 -	0	0	0
SINE	Alu	B1_Mus1	chr3	14686551	14686566 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	14833253	14833257 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	14875745	14875749 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	14916852	14916857 +	0	0	0
SINE	B2	B2_Mm1t	chr3	14916965	14916984 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	14917351	14917358 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	14925801	14925804 +	0	0	0
SINE	Alu	B1_Mus2	chr3	14933073	14933087 -	0	0	0
SINE	Alu	B1_Mm	chr3	15034674	15034689 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	15175324	15175328 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	15281239	15281278 +	0	0	0
LTR	MaLR	MTA_Mm	chr3	15341545	15341584 +	0	0	0.342623
SINE	B2	B2_Mm1a	chr3	15350215	15350234 +	0	0	0
LTR	MaLR	MTB	chr3	15409632	15409672 +	0	0	0
LTR	MaLR	MTB_Mm	chr3	15409850	15409891 -	0	0	0
LTR	MaLR	MTA_Mm	chr3	15444842	15444882 +	0	0	0
SINE	Alu	B1_Mus1	chr11	31093673	31093821 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	31093822	31093846 +	0	0	0
SINE	Alu	B1_Mm	chr3	15514032	15514044 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	15514044	15514048 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	15523969	15523973 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	15542001	15542005 +	0	0	0
SINE	Alu	B1_Mm	chr3	15748203	15748218 +	0	0	0
LTR	MaLR	MTA_Mm	chr11	31323251	31323647 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr3	15760431	15760434 +	0	0	0
LINE	L1	L1_Mus2	chr3	15826022	15826480 -	0	0	0
LTR	MaLR	MTA_Mm	chr3	15856558	15856590 +	0.591053	0	0
SINE	Alu	B1_Mus2	chr3	15929101	15929116 -	0	0	0
LTR	MaLR	MTA_Mm	chr4	3014062	3014455 -	0	0	0
SINE	Alu	B1_Mus1	chr4	3073799	3073918 -	0	0	0
SINE	Alu	B1_Mus2	chr4	3378663	3378798 -	0	0	0
SINE	Alu	B1_Mus2	chr4	3569715	3569861 -	0	0	0
SINE	Alu	B1_Mus2	chr4	3720228	3720373 -	0	0	0.615324
SINE	Alu	B1_Mus1	chr4	3736676	3736821 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	3736822	3736855 +	0.332194	0	0.615324
SINE	Alu	B1_Mus2	chr4	3781228	3781373 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	5350835	5350860 +	0	0	0.342623
LINE	L1	L1VL4	chr4	5938064	5941310 -	0	0	0
SINE	Alu	B1_Mus1	chr4	6164096	6164243 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	6164244	6164268 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	6234245	6234298 +	0	0	0
SINE	Alu	B1_Mm	chr4	6398368	6398488 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	6398489	6398533 +	0	0	0
SINE	Alu	B1_Mus2	chr4	6452102	6452248 -	0	0	0
LTR	MaLR	MTA_Mm	chr4	6478145	6478544 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	6479640	6480039 +	0	0	0
SINE	Alu	B1_Mur1	chr4	7107751	7107889 -	0	0	0
SINE	Alu	B1_Mm	chr4	7146176	7146325 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	7382583	7382614 +	0	0	0

LTR	MaLR	MTA_Mm	chr4	7446578	7446966	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	7825165	7825215	+	0	0	0
SINE	Alu	B1_Mm	chr4	7888196	7888329	-	0	0	0
SINE	Alu	B1_Mus2	chr4	8026375	8026523	-	0	0	0
LTR	MaLR	MTA_Mm	chr4	8440128	8440523	-	0	0	0
LTR	MaLR	MTA_Mm	chr4	8441621	8442016	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	8442877	8442903	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	8569975	8569997	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	8611681	8611702	+	0	0	0
SINE	Alu	B1_Mus2	chr11	32024718	32024865	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	32024866	32024889	+	0	0	0
LTR	MaLR	MTEa	chr4	8896206	8896583	-	0	0	0
SINE	Alu	PB1D9	chr4	9047076	9047193	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	9047194	9047228	+	0	0	0
SINE	Alu	B1_Mm	chr4	9608727	9608873	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	9608874	9608897	+	0	0	0
LTR	MaLR	MTA_Mm	chr4	9694928	9695323	-	0	0	0
LINE	L1	L1_Mus1	chr4	10055682	10061390	-	0	0	0
LTR	MaLR	MTA_Mm	chr4	10229581	10229962	-	0	0	0
LTR	MaLR	MTA_Mm	chr4	10231059	10231454	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	10800300	10800338	+	0	0	0
LTR	MaLR	MTA_Mm	chr4	10885026	10885421	-	0	0	0
LTR	MaLR	MTA_Mm	chr4	10886514	10886909	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	10915401	10915435	+	0	0	0
SINE	Alu	B1_Mus2	chr4	11397676	11397821	-	0	0	0
SINE	Alu	B1_Mus1	chr4	11669591	11669738	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	11865134	11865163	+	0	0	0
SINE	Alu	B1_Mm	chr4	11947851	11947993	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	11947994	11948021	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	12680998	12681034	+	0	0	0
LINE	L1	L1_Mus2	chr4	12915295	12921364	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	13361888	13361914	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	13634284	13634320	+	0	0	0
SINE	Alu	B1_Mm	chr11	32428918	32429065	+	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	32429066	32429091	+	0	0	0.342623
LTR	MaLR	MTA_Mm	chr4	14000339	14000734	+	0	0	0
LTR	MaLR	MTA_Mm	chr4	14042330	14042723	+	0	0	0
SINE	Alu	B1_Mus1	chr4	14762929	14763076	+	0	0	0
LTR	MaLR	MTA_Mm	chr4	15136900	15137299	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	15652389	15652450	+	0	0.708814	0
SINE	Alu	B1_Mm	chr4	16868135	16868276	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	16868277	16868300	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	18053266	18053324	+	0.332194	0.708814	0.957947
LTR	MaLR	MTA_Mm	chr4	18185860	18186250	-	0	0	0
LTR	MaLR	MTA_Mm	chr4	18232111	18232499	+	0	0	0
SINE	Alu	B1_Mus2	chr4	18770301	18770447	-	0	0	0
LTR	MaLR	MTA_Mm	chr4	18894286	18894680	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	19704047	19704074	+	0	0	0
LINE	L1	L1Md_F2	chr4	19710823	19716403	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	20753732	20753777	+	0	0	0
LTR	MaLR	MTA_Mm	chr4	21143107	21143497	+	0	0	0
LTR	MaLR	MTA_Mm	chr4	21289318	21289709	-	0	0	0
LTR	MaLR	MTA_Mm	chr4	21770608	21771007	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	22627283	22627304	+	0.591053	0	0
LTR	MaLR	MTA_Mm	chr4	22708514	22708909	-	0	0	0
SINE	Alu	B1_Mm	chr11	33074335	33074480	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	22761293	22761331	+	0	0	0
LTR	MaLR	MTA_Mm	chr4	22891566	22891960	+	0	0	0
LINE	L1	L1Md_T	chr4	23021702	23028342	-	0	0	0
Low_complexity	Low_complexity	A-rich	chr4	23205350	23205436	+	0	0	0
LTR	MaLR	MTA_Mm	chr4	23299286	23299680	+	0	0	0
LTR	MaLR	MTA_Mm	chr4	23429776	23430171	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	23532611	23532640	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	23692776	23692813	+	0	0	0
LINE	L1	L1_Mus1	chr4	23969521	23975351	-	0	0	0
LTR	MaLR	MTA_Mm	chr4	24064943	24065332	-	0	0	0
SINE	Alu	B1_Mus1	chr4	24087560	24087706	+	0.332194	0	0

Simple_repeat	Simple_repeat	(A)n	chr4	24128331 24128376 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	24313510 24313905 +	0	0	0
SINE	Alu	B1_Mus2	chr4	24407120 24407266 +	0	0	0
SINE	Alu	B1_Mm	chr4	25323459 25323603 +	0	0	0
SINE	Alu	B1_Mus1	chr4	25349148 25349286 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	25478843 25478910 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	26972586 26972976 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	27022466 27022505 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	27064647 27064704 +	0	0	0
SINE	Alu	B1_Mus2	chr4	27294488 27294634 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	27294635 27294697 +	0	0	0
SINE	B2	B2_Mm2	chr4	27514743 27514905 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	27650342 27650371 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	28093003 28093036 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	28950004 28950402 +	0	0	0
SINE	Alu	B1_Mm	chr4	29564094 29564230 -	0	0	0
SINE	Alu	B1_Mus1	chr4	29620025 29620171 -	0	0	0
LINE	L1	L1Md_F2	chr4	29795118 29801437 -	0	0	0
SINE	Alu	B1_Mus1	chr4	29840790 29840935 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	30382638 30382682 +	0	0	0
SINE	B2	B3A	chr4	30828775 30828972 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	31468380 31468414 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	31783053 31783445 -	0	0	0
SINE	Alu	B1_Mus1	chr4	32049000 32049137 -	0	0	0
SINE	Alu	B1_Mus2	chr4	32158334 32158481 -	0	0	0
Low_complexity	Low_complexity	A-rich	chr4	32742344 32742414 +	0	0	0
SINE	B2	B2_Mm1a	chr4	33063628 33063813 +	0	0	0
SINE	Alu	B1_Mm	chr4	33396534 33396680 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	33396681 33396708 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	33693507 33693902 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	33708858 33709254 +	0	0	0
SINE	B2	B2_Mm1a	chr4	35113758 35113950 +	0	0	0
SINE	B2	B2_Mm1t	chr4	35123347 35123538 -	0	0	0
SINE	Alu	B1_Mm	chr4	35185531 35185683 +	0	0	0.615324
LTR	MaLR	MTA_Mm	chr4	35553759 35554139 +	0	0	0
LTR	ERVK	RMER6C	chr4	37877886 37878552 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	37922299 37922335 +	0	0	0
LINE	L1	L1Md_F2	chr4	38161519 38166913 +	0	0	0
SINE	Alu	B1_Mm	chr4	38328185 38328331 +	0	0	0
SINE	Alu	B1_Mus2	chr4	39679015 39679160 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	39679161 39679188 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	39779380 39779774 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	40316374 40316765 +	0	0	0
SINE	Alu	B1_Mus2	chr4	40596879 40597001 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	40627381 40627447 +	0	0	0
SINE	Alu	B1_Mur2	chr4	41436960 41437108 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	41437109 41437132 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	41696687 41697073 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	42233390 42233422 +	0	0	0
LTR	MaLR	MTB	chr4	42919796 42920185 -	0	0	0
SINE	Alu	B1_Mus1	chr4	43029974 43030120 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	43030121 43030143 +	0	0	0
SINE	Alu	B1_Mus1	chr4	43475439 43475586 -	0	0	0
SINE	Alu	B1_Mus2	chr4	43622574 43622703 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	43622704 43622739 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	43846753 43846791 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	44011056 44011082 +	0	0	0
SINE	Alu	B1_Mm	chr4	44303118 44303263 -	0	0	0
SINE	B2	B2_Mm1a	chr4	44306747 44306938 -	0	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr4	44480689 44480722 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	44970626 44971023 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	44987617 44987649 +	0	0	0
SINE	Alu	B1_Mus1	chr4	45643159 45643305 -	0	0	0
LTR	MaLR	MTA_Mm	chr4	45672824 45673216 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	35649532 35649582 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	47510223 47510616 +	0	0	0
SINE	Alu	B1_Mus2	chr4	47931424 47931570 -	0	0	0

SINE	Alu	B1_Mm	chr4	48517505 48517651 +	0	0	0
SINE	Alu	B1_Mus1	chr4	48540202 48540352 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	48540353 48540377 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	49615940 49615966 +	0	0.708814	0
SINE	Alu	B1_Mus1	chr4	49642383 49642530 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	49931280 49931674 +	0	0	0
LINE	L1	L1_Mus1	chr4	50310116 50315638 -	0	0	0
SINE	Alu	B1_Mm	chr4	51103091 51103237 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	51275579 51275621 +	0	0	0
SINE	Alu	B1_Mus1	chr4	51513640 51513787 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	51513788 51513872 +	0	0	0
SINE	Alu	B1_Mus1	chr4	51773017 51773164 -	0	0	0
SINE	Alu	B1_Mm	chr4	52777304 52777450 -	0	0	0
LINE	L1	L1Md_T	chr4	53399667 53405975 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	53410395 53410426 +	0	0	0
SINE	Alu	B1_Mm	chr4	53952595 53952740 -	0.332194	0	0
LTR	MaLR	MTA_Mm	chr4	53978052 53978440 -	0	0	0
SINE	Alu	B1_Mus1	chr4	53984327 53984474 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	53984475 53984499 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	54040906 54040950 +	0	0	0
LINE	L1	L1Md_T	chr4	54683065 54689756 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	54940717 54940745 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	54940837 54940867 +	0	0	0
SINE	Alu	B1_Mus2	chr4	55249561 55249707 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	55249708 55249747 +	0	0	0
SINE	Alu	B1_Mm	chr4	55320534 55320678 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	55355937 55355977 +	0	0	0
LTR	MaLR	MTB_Mm	chr4	55422717 55423113 +	0	0	0
SINE	Alu	B1_Mm	chr4	55482170 55482317 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	55482318 55482348 +	0	0	0
SINE	Alu	B1_Mm	chr4	57843339 57843485 -	0	0	0
LINE	L1	L1Md_T	chr4	58024858 58031068 -	0	0	0
SINE	Alu	B1_Mm	chr4	59144276 59144422 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	59275417 59275484 +	0	0	0
SINE	Alu	B1_Mm	chr4	59584603 59584746 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	61946739 61946775 +	0	0	0
SINE	Alu	B1_Mus2	chr4	62183082 62183219 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	62183220 62183243 +	0	0	0
SINE	B2	B2_Mm1t	chr11	37686231 37686420 -	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr11	37707494 37707530 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	63483229 63483282 +	0	0	0.342623
LTR	MaLR	MTA_Mm	chr4	64632789 64633184 -	0	0	0
SINE	Alu	B1_Mm	chr11	37913545 37913690 +	0	0	0
LINE	L1	L1Md_T	chr4	66292354 66299296 -	0	0	0
SINE	Alu	B1_Mus2	chr4	66511754 66511923 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	66534598 66534642 +	0	0	0
SINE	Alu	B1_Mus1	chr4	66796993 66797134 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	66797135 66797179 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	67086673 67086700 +	0	0	0
SINE	Alu	B1_Mm	chr4	67236896 67237041 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	67624523 67624549 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	67768927 67768969 +	0	0	0
SINE	Alu	B1_Mm	chr4	67798823 67798969 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	67798970 67799010 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	68173074 68173115 +	0.332194	0	0
SINE	Alu	B1_Mus1	chr4	68274007 68274153 -	0	0	0
SINE	Alu	B1_Mus2	chr4	68348928 68349073 +	0	0.380568	0
LTR	MaLR	MTA_Mm	chr4	68934271 68934665 -	0	0	0.615324
LTR	MaLR	MTA_Mm	chr4	69104219 69104613 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	69489105 69489135 +	0.332194	0	0
LINE	L1	L1Md_F2	chr4	70385447 70390478 +	0	0	0
SINE	Alu	B1_Mm	chr4	70630602 70630747 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	70630748 70630777 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	70648761 70649156 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	72162917 72162965 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	72249869 72249916 +	0	0	0
LTR	ERVK	RLTR13D6	chr4	72380999 72381762 -	0	0	0

SINE	Alu	B1_Mus2	chr4	73068672 73068815 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	73115255 73115293 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	73127559 73127953 -	0	0	0
SINE	Alu	B1_Mus1	chr4	73855309 73855455 +	0.591053	0.708814	0.342623
Simple_repeat	Simple_repeat	(A)n	chr4	74161359 74161402 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	74750548 74750578 +	0	0	0.342623
LTR	MaLR	MTA_Mm	chr4	74963922 74964318 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	74968815 74968862 +	0	0	0
SINE	Alu	B1_Mus1	chr4	75070524 75070670 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	75070671 75070700 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	38943030 38943060 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	38949656 38949682 +	0	0	0
SINE	Alu	B1_Mm	chr11	39035338 39035485 -	0	0	0
LINE	L1	L1Md_F3	chr4	78081594 78087717 -	0	0	0
LINE	L1	Lx3A	chr4	78236009 78242084 -	0	0	0
LTR	MaLR	MTA_Mm	chr4	78330325 78330721 +	0	0	0
LTR	MaLR	MTA_Mm	chr11	39208683 39209077 -	0	0	0.615324
LTR	MaLR	MTA_Mm	chr4	78494880 78495276 +	0	0	0
SINE	Alu	B1_Mus1	chr4	78528151 78528298 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	78617374 78617399 +	0	0	0
LINE	L1	L1Md_F2	chr4	78621922 78626952 -	0	0	0
LINE	L1	L1Md_F2	chr4	79161176 79166486 -	0	0	0
SINE	Alu	B1_Mus1	chr4	79283352 79283479 -	0	0	0
LINE	L1	L1_Mus3	chr4	79307124 79312528 -	0	0	0
SINE	Alu	B1_Mus2	chr11	39380534 39380679 -	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	80191916 80191954 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr4	81332754 81332801 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	81374077 81374112 +	0	0	0
SINE	Alu	B1_Mus1	chr11	39508462 39508605 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	39508606 39508642 +	0	0	0
LINE	L1	L1Md_T	chr4	81833098 81839570 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	82333728 82333758 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	83590327 83590356 +	0	0	0
SINE	Alu	B1_Mm	chr4	84476717 84476862 -	0	0	0
LINE	L1	L1Md_T	chr4	85053278 85059536 -	0.332194	0	0
SINE	Alu	B1_Mm	chr4	85141714 85141860 +	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr4	85141861 85141887 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	85250840 85250918 +	0.591053	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr4	86372602 86372627 +	0	0	0
SINE	Alu	B1_Mm	chr4	86498394 86498532 -	1.58764	0.380568	1.84597
SINE	B2	B2_Mm2	chr4	86509973 86510151 +	0	0.708814	0
SINE	Alu	B1_Mus2	chr11	40105350 40105497 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	87107878 87107914 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	87260185 87260232 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	87330405 87330427 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	88656837 88657232 +	0	0	0
LINE	L1	L1Md_T	chr4	89381050 89386227 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	90034151 90034546 +	0	0	0
LTR	MaLR	MTD	chr4	90047842 90048269 +	0	0	0
SINE	Alu	B1_Mm	chr4	90135846 90135992 +	0	0	0
LINE	L1	L1Md_T	chr4	90220445 90226938 -	0	0	0
SINE	Alu	B1_Mm	chr4	90683429 90683575 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	40453841 40453884 +	0	0	0
LTR	MaLR	MTA_Mm	chr11	40456214 40456608 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	91067056 91067095 +	0.332194	0	0
LINE	L1	L1_Mus3	chr4	92302151 92308336 -	0	0	0
LTR	MaLR	MTA_Mm	chr4	92417264 92417658 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	92418758 92419152 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	92858997 92859021 +	0	0	0
SINE	Alu	B1_Mus2	chr4	93238234 93238380 -	0	0	0
SINE	Alu	B1F2	chr11	40788706 40788827 -	0	0	0
SINE	Alu	B1_Mm	chr4	96367563 96367710 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	96367711 96367760 +	0	0	0
SINE	Alu	B1_Mus2	chr4	97064985 97065133 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	98515058 98515077 +	0	0	0
SINE	Alu	B1_Mus2	chr4	98519091 98519237 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	98519238 98519283 +	0	0	0.342623

SINE	Alu	B1_Mm	chr4	98519284 98519428 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr4	98519429 98519457 +	0	0	0.342623
LTR	MaLR	MTB	chr4	98547444 98547837 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	98590274 98590310 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	99106879 99106904 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	99157705 99158100 +	0.332194	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr4	99314056 99314094 +	0	0	0
SINE	Alu	B1_Mm	chr11	41477511 41477634 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	10097471 10097474 +	0.332194	0	0
SINE	Alu	B1_Mus1	chr11	41564180 41564327 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	41564328 41564354 +	0	0	0
SINE	Alu	B1_Mus1	chr4	10155174 10155189 +	0	0	0
SINE	Alu	B1_Mm	chr4	10160411 10160425 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	10160425 10160429 +	0	0	0
SINE	Alu	B1_Mm	chr4	10163097 10163112 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	10163112 10163117 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	10190176 10190178 +	1.18211	0.708814	3.05526
Simple_repeat	Simple_repeat	(A)n	chr4	10236818 10236821 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	10318152 10318162 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	10338550 10338553 +	3.28746	4.63345	1.82461
SINE	Alu	B1_Mus1	chr4	10347268 10347283 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	10376897 10376901 +	0	0	0
SINE	Alu	B1_Mus2	chr4	10494199 10494213 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	10494214 10494217 +	0	0	0.342623
LINE	L1	L1Md_T	chr4	10496511 10497198 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	10512663 10512668 +	0	0	0
LTR	MaLR	MTA_Mm	chr11	42032811 42033210 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	10581240 10581246 +	0	0	0
SINE	Alu	B1_Mus1	chr4	10640109 10640124 -	0	0	0
SINE	Alu	B1_Mus1	chr4	10650798 10650813 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	10650813 10650816 +	0	0	0
SINE	B2	B2_Mm1a	chr4	10689370 10689389 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	42218746 42218773 +	0	0	0
SINE	Alu	B1_Mus2	chr4	10760834 10760849 -	0	0	0
LTR	MaLR	MTA_Mm	chr4	10809347 10809386 -	0	0	0
LTR	MaLR	MTA_Mm	chr4	10809494 10809534 -	0	0	0
SINE	Alu	B1_Mus1	chr4	10848155 10848169 -	0	0	0
SINE	Alu	B1_Mur2	chr4	10887559 10887573 -	0	0	0
SINE	Alu	B1_Mus2	chr4	10928865 10928880 +	0	0	0
Simple_repeat	Simple_repeat	(GGAA)n	chr4	10928882 10928889 +	0	0	0
SINE	Alu	B1_Mm	chr4	10932102 10932116 +	0	0	0
SINE	Alu	B1_Mm	chr4	10934438 10934453 -	0	0	0
SINE	Alu	B1_Mm	chr4	10963966 10963980 -	0	0	0
SINE	Alu	B1_Mus1	chr11	42710611 42710766 -	0	0	0
SINE	Alu	B1_Mus2	chr4	11138382 11138396 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	11138396 11138399 +	0.332194	0	0
LTR	MaLR	MTA_Mm	chr11	42881138 42881530 +	0	0	0
SINE	Alu	B1_Mm	chr4	11199244 11199258 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	42941225 42941291 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	11296611 11296616 +	0	0	0
SINE	Alu	B1_Mm	chr4	11310856 11310871 -	0	0	0
SINE	Alu	B1_Mus2	chr4	11459379 11459393 +	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chr4	11459394 11459400 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr4	11467069 11467083 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	11471636 11471639 +	0	0	0
SINE	Alu	B1_Mus1	chr4	11572746 11572761 -	0	0	0
SINE	Alu	B1_Mus2	chr4	11585993 11586007 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	43247141 43247176 +	0	0	0.342623
LTR	MaLR	MTB	chr4	11622249 11622288 -	0	0	0
SINE	Alu	B1_Mus1	chr4	11635481 11635496 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	11635496 11635502 +	0	0	0
SINE	Alu	B1_Mus2	chr4	11651061 11651075 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	11651075 11651078 +	0	0	0
SINE	Alu	B1_Mur4	chr4	11668220 11668239 -	0	0	0
SINE	Alu	B1_Mm	chr4	11681757 11681772 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	11681772 11681774 +	0	0	0.342623
SINE	Alu	B1_Mus1	chr4	11747182 11747194 -	0	0	0



SINE	B2	B2_Mm1a	chr4	11797335 11797353 -	0	3.53776	0.342623
SINE	Alu	B1_Mus2	chr4	11811229 11811243 -	0	0	0
SINE	Alu	B1_Mus2	chr4	11820609 11820623 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	11842816 11842819 +	0	0	0
SINE	Alu	B1_Mur4	chr4	11854638 11854653 +	0	0	0
SINE	Alu	B1_Mus2	chr4	11858212 11858226 +	0	0	0
LINE	L1	L1VL4	chr4	11861820 11862301 +	0	0	0
SINE	Alu	B1_Mus2	chr4	11865739 11865753 -	0	0	0
SINE	B2	B2_Mm1a	chr4	11866664 11866683 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	11872163 11872167 +	0	0	0
SINE	Alu	B1_Mus2	chr4	11917548 11917563 -	0	0	0
SINE	Alu	B1_Mm	chr4	11931234 11931246 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	11942627 11942632 +	1.84649	1.08938	2.18859
Simple_repeat	Simple_repeat	(GAAA)n	chr4	11986925 11986938 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	12003052 12003056 +	0	0	0
SINE	B2	B3	chr11	43652429 43652623 -	0	0	0
SINE	Alu	B1_Mus2	chr4	12102621 12102635 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	12118806 12118811 +	0	0	0
SINE	Alu	B1_Mus1	chr4	12144484 12144498 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	12144498 12144501 +	0	0	0
SINE	Alu	B1_Mus1	chr4	12150437 12150452 +	0	0	0
SINE	Alu	B1_Mus1	chr4	12189735 12189750 +	0	0	0
SINE	Alu	B1_Mus1	chr4	12215899 12215914 +	0	0	0
SINE	Alu	B1_Mur3	chr4	12249746 12249761 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	12253641 12253644 +	1.18211	0.761137	1.84597
Low_complexity	Low_complexity	GA-rich	chr4	12266564 12266569 +	0.664388	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr4	12276018 12276020 +	0	0	0
SINE	Alu	B1_Mus2	chr4	12295562 12295576 +	0	0	0
SINE	Alu	B1_Mus1	chr4	12353292 12353306 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr4	12353306 12353310 +	0	0	0
SINE	Alu	B1_Mus1	chr4	12380741 12380755 -	0	0	0
LTR	MaLR	MTB	chr4	12401446 12401485 -	0	0	0.342623
LTR	MaLR	MTA_Mm	chr4	12417267 12417307 +	0	0	0
SINE	Alu	B1_Mm	chr4	12425627 12425642 -	0	0	0.615324
SINE	Alu	B1_Mm	chr4	12441739 12441753 -	0	0	0
SINE	Alu	B1_Mm	chr4	12441788 12441803 -	0	0	0
SINE	Alu	B1_Mus2	chr4	12454139 12454153 +	0	0	0
Simple_repeat	Simple_repeat	(GAA)n	chr4	12454154 12454160 +	0	0.380568	0.342623
SINE	Alu	B1_Mus2	chr4	12476037 12476052 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	12476052 12476054 +	0	0	0
SINE	Alu	B1_Mus2	chr4	12562846 12562861 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr4	12562861 12562863 +	0	0	0.615324
SINE	Alu	B1_Mm	chr4	12567219 12567233 +	0	0	0
LTR	MaLR	MTC	chr4	12567234 12567271 +	0	0	0
SINE	Alu	B1_Mus2	chr4	12665441 12665456 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	12665456 12665458 +	0	0	0.342623
SINE	Alu	B1_Mm	chr4	12675283 12675298 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	12675298 12675301 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	44431278 44431356 +	11.4927	4.9617	3.02097
SINE	Alu	B1_Mm	chr4	12714791 12714806 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	12732805 12732807 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	12758660 12758699 -	0	0	0
LTR	MaLR	MTA_Mm	chr4	12860569 12860608 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	12865038 12865042 +	0	0	0
SINE	Alu	B1_Mus2	chr4	12866295 12866310 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	12866310 12866312 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	12894921 12894923 +	0	0	0
SINE	Alu	B1_Mus2	chr4	12918458 12918472 +	0.591053	0	0.615324
SINE	Alu	B1_Mus2	chr4	12922345 12922357 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	12922357 12922359 +	0	0	0
SINE	Alu	B1_Mm	chr4	12922454 12922469 -	0	0	0
SINE	B2	B2_Mm1t	chr4	12947950 12947969 -	0	0	0
SINE	Alu	B1_Mus2	chr4	12961790 12961805 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	12961805 12961808 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr4	12998041 12998043 +	0	0	0
SINE	Alu	B1_Mus1	chr4	13019337 13019350 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr4	13019350 13019353 +	0	0	0.342623

SINE	Alu	B1_Mm	chr4	13019794 13019807 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	13021836 13021839 +	0	0	0
SINE	Alu	B1_Mus1	chr4	13185256 13185271 -	0	0	0.342623
SINE	B2	B2_Mm1a	chr4	13186228 13186247 -	0	0	0
SINE	Alu	B1_Mm	chr4	13190312 13190327 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	13190327 13190333 +	0	0	0
SINE	Alu	B1_Mus1	chr4	13203830 13203844 -	0	0	0
SINE	Alu	B1_Mus2	chr4	13214013 13214027 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	13214027 13214030 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	13235980 13235983 +	0	0	0
SINE	Alu	B1_Mus1	chr4	13278284 13278299 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	13278299 13278301 +	0	0	0
SINE	Alu	B1_Mus2	chr4	13282016 13282030 -	0	0	0
SINE	Alu	B1_Mus2	chr4	13283652 13283667 +	0	0	0
LTR	MaLR	MTA_Mm	chr4	13310631 13310671 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	13315238 13315240 +	0	0	0
SINE	Alu	B1_Mus2	chr4	13334713 13334728 +	0	0	0
SINE	Alu	B1_Mus2	chr4	13347823 13347838 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	13449506 13449509 +	0.591053	0	0
SINE	Alu	B1_Mus1	chr4	13475466 13475480 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	45778527 45778599 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	20892610 20892653 +	0	0	0
SINE	Alu	B1_Mm	chr4	13518764 13518777 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	13518777 13518779 +	0	0	0
SINE	Alu	B1_Mus2	chr4	13522419 13522433 -	0	0	0
SINE	Alu	B1_Mus2	chr4	13543471 13543486 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	13543486 13543488 +	0	0	0
SINE	Alu	B1_Mus2	chr4	13566235 13566250 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	13568690 13568693 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	13575277 13575279 +	0	0	0
SINE	Alu	B1_Mm	chr4	13583391 13583406 -	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr4	13616149 13616153 +	0	0	0
SINE	Alu	B1_Mus2	chr4	13644442 13644457 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	13645941 13645944 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	13681386 13681389 +	0	0	0
SINE	Alu	B1_Mm	chr4	13714529 13714544 -	0	0	0
SINE	Alu	B1_Mus2	chr4	13810602 13810617 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	13810617 13810620 +	0	0	0
SINE	Alu	B1_Mm	chr11	46204625 46204770 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	13869777 13869783 +	0	0	0
SINE	Alu	B1_Mus2	chr4	14023774 14023789 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	14045550 14045553 +	0	0	0
Low_complexity	Low_complexity	AT-rich	chr4	14070153 14070156 +	0	0	0
SINE	Alu	B1_Mus2	chr4	14111045 14111059 -	0	0	0
SINE	Alu	B1_Mm	chr4	14112007 14112022 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	14122455 14122458 +	0	0	0
SINE	B2	B2_Mm1a	chr4	14128252 14128270 +	0	0	0
Low_complexity	Low_complexity	GA-rich	chr11	46608545 46608701 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	14235316 14235318 +	0	0	0
SINE	Alu	B1_Mus2	chr11	46675490 46675636 -	0	0	0
SINE	Alu	B1_Mus1	chr4	14289478 14289492 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr4	14289492 14289500 +	0	0	0
SINE	Alu	B1_Mm	chr4	14296447 14296459 -	0	0	0
SINE	B2	B3A	chr4	14297047 14297066 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	14297066 14297069 +	0	0	0
SINE	Alu	B1_Mus1	chr4	14298369 14298383 -	0	0	0
SINE	Alu	B1_Mus2	chr11	46714729 46714873 -	0	0	0
LTR	MaLR	MTA_Mm	chr4	14308738 14308778 +	0	0	0
SINE	Alu	B1_Mus2	chr4	14351704 14351718 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	14351718 14351720 +	0	0	0
SINE	Alu	B1_Mus1	chr4	14363840 14363855 -	0	0	0
SINE	Alu	B1_Mm	chr4	14480038 14480053 -	0	0	0
SINE	Alu	B1_Mm	chr4	14491621 14491635 -	0	0	0
SINE	Alu	B1_Mus2	chr4	14499704 14499718 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	14499718 14499721 +	0	0	0
SINE	B2	B2_Mm1t	chr4	14507140 14507159 -	0	0	0
SINE	Alu	B1_Mus2	chr4	14515358 14515373 -	0	0	0.615324

LTR	MaLR	MTA_Mm	chr4	14519889 14519928 -	0	0	0
SINE	Alu	B1_Mm	chr4	14557942 14557956 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	14557956 14557959 +	0	0	0
SINE	Alu	B1_Mm	chr4	14559931 14559945 -	0	0	0
SINE	Alu	B1_Mus2	chr4	14571808 14571823 -	0	0	0
SINE	Alu	B1_Mm	chr4	14622539 14622554 +	0	0	0
SINE	Alu	B1_Mus2	chr4	14623026 14623041 -	0	0	0
SINE	Alu	B1_Mm	chr4	14696813 14696828 -	0	0	0
SINE	Alu	B1_Mm	chr4	14702899 14702913 -	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	14719106 14719108 +	0	0	0
SINE	Alu	B1_Mus2	chr4	14743778 14743792 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	14743792 14743794 +	0	0	0
SINE	Alu	B1_Mus1	chr4	14744297 14744310 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	14744310 14744313 +	0	0	0
SINE	Alu	B1_Mus2	chr4	14796433 14796447 -	2.94048	0.708814	1.55647
SINE	Alu	B1_Mus2	chr4	14800258 14800273 -	0	0	0
SINE	Alu	B1_Mus2	chr4	14804683 14804697 -	0	0	0
SINE	Alu	B1_Mus2	chr4	14854172 14854187 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	14887935 14887939 +	0	0	0.342623
SINE	Alu	B1_Mus1	chr4	14908048 14908058 -	0	0	0
SINE	Alu	B1_Mus1	chr4	14911661 14911674 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	14911674 14911677 +	0	0.380568	0
SINE	Alu	B1_Mus1	chr4	14933185 14933198 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	14933198 14933201 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr4	14965126 14965141 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr4	14965141 14965147 +	0	0	0
SINE	Alu	B1_Mus1	chr4	15026595 15026608 +	0	0	0.342623
SINE	B2	B2_Mm1a	chr4	15033196 15033215 -	0	0	0
SINE	Alu	B1_Mus1	chr4	15169781 15169795 -	0	0	0
SINE	Alu	B1_Mm	chr4	15254341 15254356 -	0	0	0
LTR	MaLR	MTB	chr4	15258388 15258427 +	0	0	0
LINE	L1	L1Md_F2	chr11	48012487 48017719 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	15409431 15409434 +	0	0	0
LTR	ERV_L	MLT2D	chr4	15409434 15409464 +	0	0	0
SINE	Alu	B1_Mus2	chr4	15429603 15429618 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	15429618 15429620 +	0	0	0
SINE	Alu	B1_Mus2	chr4	15514013 15514028 -	0	0	0
LTR	ERV_L	RMER15	chr4	15518379 15518412 -	0	0	0
SINE	Alu	B1_Mus1	chr4	15518413 15518428 -	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr4	15521076 15521079 +	0	0.708814	0
SINE	Alu	B1_Mus1	chr4	15544074 15544088 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	15544088 15544091 +	0	0	0
SINE	Alu	B1_Mm	chr4	15551334 15551349 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr4	15551349 15551352 +	0.332194	0	0
SINE	Alu	B1_Mus2	chr4	15558705 15558719 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	3118537 3118593 +	0	0	0
LINE	L1	L1Md_A	chr5	3175316 3180450 -	0	0	0
SINE	Alu	B1_Mus2	chr5	3538007 3538156 -	0	0	0
LTR	MaLR	MTA_Mm	chr5	3650172 3650566 -	0	0	0.342623
LTR	MaLR	MTA_Mm	chr5	3918469 3918847 -	0	0	0
SINE	Alu	B1_Mus2	chr11	48465741 48465887 +	0	0	1.23065
SINE	Alu	B1_Mus2	chr5	5500825 5500969 -	0.591053	0	0
LTR	MaLR	MTA_Mm	chr5	5903165 5903558 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	5904644 5905037 +	0	0	0
SINE	B4	B4A	chr5	6177374 6177638 -	0.591053	0	0
SINE	Alu	B1_Mus2	chr5	6571122 6571268 -	0.332194	0.380568	0
SINE	Alu	B1_Mus2	chr11	48600270 48600415 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	48600416 48600442 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	48612907 48612942 +	0	0	0
LINE	L1	Lx8	chr5	7400952 7405667 -	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr5	7495409 7495444 +	0	0	0
SINE	Alu	B1_Mm	chr5	7549707 7549853 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	7584265 7584307 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	7845535 7845614 +	0	0	0
SINE	Alu	B1_Mus1	chr5	8794566 8794686 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	8794687 8794714 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr5	9242552 9242580 +	0	0	0

SINE	Alu	B1_Mm	chr5	10702482 10702630 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	10702631 10702666 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	11177436 11177465 +	0	0	0
SINE	B2	B2_Mm2	chr10	21251148 21251337 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	11995027 11995059 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	12358935 12358968 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	12800522 12800913 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	12802009 12802424 +	0	0	0
SINE	Alu	B1_Mus1	chr5	13088007 13088148 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	13088149 13088187 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	13890440 13890833 -	0	0	0
LTR	MaLR	MTA_Mm	chr5	14443586 14443980 -	0	0	0
Low_complexity	Low_complexity	A-rich	chr5	14915889 14915981 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	15252668 15252706 +	0	0.380568	0
SINE	Alu	B1_Mus2	chr5	15897110 15897256 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	15897257 15897324 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr5	15965484 15965549 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	16186837 16186881 +	0	0	0
LINE	L1	L1Md_T	chr5	16538848 16545477 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	16670882 16670932 +	0	0	0
LINE	L1	L1Md_T	chr5	16744089 16753072 -	0	0	0
SINE	Alu	B1_Mus1	chr5	16806219 16806369 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	16806370 16806391 +	0	0	0
SINE	Alu	B1_Mus1	chr5	16975257 16975403 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	18546832 18547226 +	0	0	0
SINE	Alu	B1_Mm	chr5	20483029 20483175 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	20483176 20483215 +	0	0	0
SINE	B2	B2_Mm2	chr5	20487001 20487189 +	0	0	0
SINE	B2	B2_Mm1a	chr5	20578171 20578359 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	22081438 22081467 +	0	0	0
SINE	Alu	B1_Mus2	chr5	22120974 22121122 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr5	22121123 22121149 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr5	22177576 22177613 +	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr5	22237107 22237148 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr5	22795153 22795198 +	0	0	0
SINE	Alu	B1_Mus1	chr5	23230505 23230645 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	23805549 23805587 +	0	0	0
SINE	B2	B2_Mm2	chr5	23815082 23815270 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr5	24174715 24174860 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	24174861 24174884 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	24665602 24665996 +	0	0	0
SINE	Alu	B1_Mus1	chr11	50063738 50063884 -	0	0	0
SINE	Alu	B1_Mus2	chr5	24774817 24774957 +	0.591053	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr5	24774958 24774982 +	0.591053	0	0.615324
SINE	Alu	B1_Mm	chr5	25037318 25037463 -	0	0	0
SINE	Alu	B1_Mm	chr5	25167090 25167235 -	0	0	0
SINE	Alu	B1_Mur3	chr11	50157166 50157321 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	50157322 50157347 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	27123309 27123354 +	0	0	0
SINE	B2	B2_Mm1t	chr10	21433994 21434186 -	0	0	0
LTR	MaLR	MTA_Mm	chr5	29280719 29281114 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	29724076 29724120 +	0	0	0
SINE	Alu	PB1D10	chr5	29795441 29795551 -	0	0	0
SINE	Alu	B1_Mm	chr5	29850258 29850404 -	0	0	0
SINE	Alu	B1_Mm	chr5	30149851 30149997 -	0	0	0
SINE	Alu	B1_Mur2	chr5	30192346 30192496 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	30207683 30207717 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr5	30270786 30270872 +	0	0	0
SINE	Alu	B1_Mus1	chr5	31012576 31012714 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	31012715 31012755 +	0.332194	0	0
SINE	Alu	B1_Mur1	chr5	31301749 31301890 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	31301891 31301916 +	0	0	0
SINE	Alu	B1_Mm	chr5	31381858 31382004 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	31382005 31382036 +	0	0	0
SINE	Alu	B1_Mm	chr5	31403907 31404056 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	31404057 31404081 +	0	0	0
SINE	Alu	B1_Mus1	chr5	31408801 31408948 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr5	31408949 31408990 +	0	0	0
SINE	Alu	B1_Mm	chr5	31535153 31535299 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	32723704 32723740 +	0	0.380568	0
SINE	Alu	B1_Mus2	chr5	33129085 33129231 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	33129232 33129263 +	0	0	0
SINE	Alu	B1_Mm	chr5	33201993 33202138 +	0	0	0
SINE	Alu	B1_Mus2	chr5	33658233 33658379 -	0	0	0
SINE	B2	B3A	chr5	33676978 33677173 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	33677174 33677195 +	0	0.380568	0
SINE	Alu	B1_Mm	chr5	34060226 34060372 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr5	34060373 34060401 +	0	0.761137	0.615324
Simple_repeat	Simple_repeat	(A)n	chr5	34319091 34319128 +	0	0	0
SINE	B4	ID_B1	chr5	35431991 35432214 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	35641675 35641715 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	36357696 36358089 +	0	0	0
SINE	Alu	B1_Mm	chr5	37155726 37155872 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	37155873 37155915 +	0	0	0
SINE	Alu	B1_Mm	chr5	37437229 37437375 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	38108750 38108788 +	0	0	0
SINE	Alu	B1_Mm	chr5	38301451 38301599 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	38301600 38301631 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	39024086 39024113 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	39826534 39826929 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	40743604 40743632 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	40973251 40973649 -	0	0	0
LTR	MaLR	MTA_Mm	chr5	40974740 40975138 -	0	0	0
LTR	MaLR	MTA_Mm	chr5	41330001 41330395 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	41336037 41336071 +	0	0	0
LTR	MaLR	MTB	chr5	41403951 41404344 -	0	0	0
SINE	Alu	B1_Mm	chr5	42297793 42297939 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	42297940 42297980 +	0	0	0.342623
LINE	L1	L1Md_T	chr5	42391767 42397052 -	0	0	0
LTR	MaLR	MTB	chr5	43182415 43182787 -	0	0	0
LTR	ERVK	RMER19B	chr5	44049531 44050478 +	0	0	0
SINE	Alu	B1_Mm	chr11	51477816 51477938 -	0	0	0
SINE	Alu	B1_Mus1	chr5	44305968 44306114 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	44560452 44560490 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	45499344 45499738 -	0	0	0
SINE	B2	B2_Mm1t	chr5	45719065 45719255 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	45728257 45728310 +	0	0	0
SINE	Alu	B1_Mm	chr5	46296351 46296496 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	46296497 46296573 +	0	0	0
scRNA	scRNA	4.5SRNA	chr5	46498263 46498356 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	46726531 46726560 +	0	0	0
SINE	Alu	B1_Mm	chr5	47050350 47050495 +	0	0	1.10507
Simple_repeat	Simple_repeat	(A)n	chr5	47050496 47050521 +	0	0	1.10507
SINE	Alu	B1_Mm	chr11	51614093 51614238 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	51614239 51614264 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	48212764 48212792 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	49726187 49726226 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	50065257 50065304 +	0	0	0
LINE	L1	L1Md_F2	chr5	50466858 50471342 +	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr5	50548444 50548473 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	50571143 50571185 +	0	0	0
SINE	Alu	B1_Mus1	chr5	50583500 50583646 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	50583647 50583673 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	51064755 51064791 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	52041748 52042143 -	1.18211	0	0.957947
Simple_repeat	Simple_repeat	(A)n	chr5	52263344 52263380 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	52333307 52333703 +	0	0	0
SINE	Alu	B1_Mus1	chr5	53109935 53110070 -	0	0	0
SINE	Alu	B1_Mus1	chr5	53473877 53474024 -	0	0	0
SINE	Alu	B1_Mm	chr5	53879646 53879790 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	53879791 53879814 +	0	0	0
SINE	Alu	B1_Mus1	chr5	54071430 54071567 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	55487170 55487224 +	0	0	0
SINE	Alu	B1_Mus2	chr5	55647202 55647344 +	0	0	0

SINE	Alu	B1_Mus1	chr5	55907400 55907547 +	0	0	0
SINE	Alu	B1_Mus2	chr5	56098912 56099057 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	56146741 56146790 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	56200637 56200666 +	0	0	0
LINE	L1	L1Md_T	chr5	56228525 56235445 -	0	0	0
SINE	Alu	B1_Mm	chr5	56252636 56252750 -	0	0	0
SINE	Alu	B1_Mus2	chr5	56256680 56256825 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	56256826 56256847 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	56622432 56622468 +	0	0	0
SINE	Alu	B1_Mus2	chr5	56749775 56749901 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	56749902 56749930 +	0	0	0
SINE	Alu	B1_Mur4	chr5	56879996 56880145 -	0	0	0
SINE	Alu	B1_Mm	chr5	57419783 57419934 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	57419935 57419969 +	0	0	0
LINE	L1	L1Md_F2	chr5	57636519 57641682 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	52247261 52247300 +	0	0	0
SINE	Alu	B1_Mus1	chr11	52263217 52263353 -	0	0	0
SINE	Alu	B1_Mm	chr5	59075751 59075897 -	0	0	0
LINE	L1	L1_Mus2	chr5	59478292 59482906 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	59815412 59815453 +	0	0	0
SINE	Alu	B1_Mus2	chr5	60222719 60222866 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	60222867 60222891 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	60339531 60339580 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr5	60799577 60799644 +	0	0	0
LINE	L1	L1Md_T	chr5	60941757 60948495 -	0	0	0
LTR	MaLR	MTA_Mm	chr5	61032907 61033302 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	61073357 61073752 +	0	0	0
SINE	Alu	B1_Mm	chr5	61154611 61154737 -	0	0	0
SINE	Alu	B1_Mus1	chr5	62370353 62370497 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	62641838 62641878 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	62661189 62661232 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	62858265 62858614 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	63529151 63529192 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	63949103 63949129 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	64035374 64035404 +	0	0	0
SINE	Alu	B1_Mus1	chr5	64363734 64363881 +	0	0	0
SINE	Alu	B1_Mus2	chr5	65093923 65094048 -	0	0	0
SINE	Alu	B1_Mus2	chr5	65325893 65326039 +	0	0	0
SINE	Alu	B1_Mus1	chr5	66091919 66092066 -	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr5	66110272 66110295 +	0	0	0
SINE	Alu	B1_Mus2	chr5	66288482 66288627 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	66288628 66288654 +	0	0	0
SINE	Alu	B1_Mus2	chr5	66345681 66345839 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	66345840 66345865 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	66348009 66348044 +	0	0	0
LTR	MaLR	MTB	chr5	67518841 67519242 +	0	0	0
SINE	Alu	B1_Mus2	chr11	53120850 53120994 -	0	0	0
SINE	B2	B2_Mm1t	chr11	53126638 53126823 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	68394262 68394322 +	0	0	0
SINE	Alu	B1_Mus1	chr11	53132342 53132487 -	0	0	0
LTR	MaLR	MTA_Mm	chr5	68617578 68617972 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	69043634 69044028 -	0	0	0
LINE	L1	L1_Mus3	chr5	70478162 70484368 -	0	0	0
LINE	L1	L1Md_F2	chr5	70906915 70912884 -	0	0	0
LTR	MaLR	MTA_Mm	chr5	70961348 70961743 +	0	0	0
SINE	Alu	B1_Mus1	chr11	53240693 53240847 -	0	0	0
SINE	Alu	B1_Mus1	chr5	71111707 71111854 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	71954550 71954584 +	0	0	0
Simple_repeat	Simple_repeat	(GAA)n	chr5	73270814 73270864 +	0	0.380568	0.342623
SINE	Alu	B1_Mus2	chr5	73296706 73296840 -	0	0	0
SINE	Alu	B1_Mm	chr5	73757110 73757255 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	73757256 73757297 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	73777254 73777648 -	0	0	0
SINE	Alu	B1_Mm	chr5	74279284 74279430 +	0	0	0
LTR	MaLR	MTB	chr5	74351973 74352371 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr5	74468844 74468866 +	0	0	0
LTR	MaLR	MTB	chr5	75332738 75333130 +	0	0	0

LTR	MaLR	MTB	chr5	75765626 75766019 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	76390995 76391036 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr5	76610227 76610267 +	0	0	0
SINE	Alu	B1_Mm	chr5	77400549 77400705 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	77400706 77400729 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	77554999 77555396 +	0	0	0
SINE	Alu	B1_Mur1	chr5	77666037 77666179 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	77666180 77666203 +	0	0	0
SINE	Alu	B1_Mm	chr5	77720339 77720468 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	77720469 77720492 +	0	0	0
LTR	MaLR	MTB	chr5	78069434 78069833 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	79530078 79530137 +	0	0	0
LINE	L1	L1Md_F2	chr5	79629285 79634730 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	79976441 79976537 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	80180014 80180391 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	80317087 80317131 +	0.591053	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr5	80699241 80699277 +	0	0	0
SINE	Alu	B1_Mus2	chr5	80929421 80929567 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	80971609 80971652 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	81028523 81028547 +	0	0	0
SINE	Alu	B1_Mus2	chr5	82457878 82458024 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	82458025 82458064 +	0	0	0
SINE	Alu	B1_Mus1	chr5	82531052 82531198 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	82531199 82531287 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	82643972 82644013 +	0	0	0
SINE	Alu	B1_Mm	chr5	82768806 82768936 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	82769985 82770019 +	0	0	0
SINE	Alu	B1_Mus2	chr5	82921272 82921417 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	82921418 82921478 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	83186090 83186129 +	0	0	0
Simple_repeat	Simple_repeat	(CAAAAC)n	chr5	83348503 83348564 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	83348565 83348588 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	83354749 83354775 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	83416919 83417314 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	83603042 83603078 +	0	0	0
SINE	Alu	B1_Mur3	chr5	83622793 83622932 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr5	83622933 83623037 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	83818988 83819044 +	0	0	0
LTR	MaLR	MTB	chr5	84888623 84889016 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	85645045 85645082 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	85685292 85685323 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	85726294 85726335 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	54516067 54516091 +	0.923247	0.380568	0
LTR	MaLR	MTA_Mm	chr5	89714621 89715015 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	90023587 90023623 +	0	0	0
SINE	Alu	B1_Mus2	chr11	54610982 54611127 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	90357653 90357703 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr5	90610415 90610542 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	90617267 90617311 +	0	0	0
SINE	Alu	B1_Mus1	chr11	54646456 54646596 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	54646597 54646619 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	92157649 92158043 -	0	0	0
SINE	Alu	B1_Mm	chr5	92407018 92407161 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	92407162 92407195 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	93505586 93505980 -	0	0	0.615324
SINE	Alu	B1_Mus1	chr5	93508945 93509084 -	0	0	0
SINE	Alu	B1_Mm	chr5	93783060 93783206 +	0	0	0
SINE	Alu	PB1D9	chr5	93912102 93912210 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	93912211 93912248 +	0	0	0
Simple_repeat	Simple_repeat	(CAAG)n	chr5	93912247 93912267 +	0	0	0
SINE	Alu	B1_Mus2	chr5	93926219 93926354 -	0	0	0
SINE	Alu	B1_Mus2	chr5	94269164 94269299 +	0	0	0
SINE	Alu	PB1D9	chr5	94283295 94283403 -	0	0	0
SINE	Alu	B1_Mus1	chr5	94363164 94363309 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	94363310 94363351 +	0	0	0
SINE	Alu	B1_Mus2	chr5	94635088 94635224 +	0.332194	0	0
SINE	Alu	B1_Mus2	chr5	94951518 94951664 +	0	0	0.342623

SINE	Alu	B1_Mus2	chr5	95290737 95290873 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	95310346 95310382 +	0	0	0
SINE	Alu	B1_Mus2	chr11	54909120 54909265 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	54909266 54909287 +	0	0	0
SINE	Alu	B1_Mus1	chr5	95954363 95954510 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	95954511 95954552 +	0	0	0
SINE	Alu	B1_Mus1	chr5	96028980 96029125 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	96029126 96029163 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	96428828 96428867 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	98608362 98608397 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr5	98733725 98733775 +	0	0	0
LINE	L1	L1Md_T	chr5	99338414 99344953 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	99555181 99555208 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	10015896 10015901 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	10027225 10027264 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	10036511 10036550 +	0.591053	0	0
LTR	MaLR	MTA_Mm	chr10	22138463 22138857 -	0	0	0
SINE	Alu	B1_Mm	chr5	10118768 10118782 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	10118782 10118785 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	10153720 10153723 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	10186717 10186721 +	0	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr5	10211826 10211833 +	0	0	0
SINE	Alu	B1_Mus1	chr5	10217810 10217824 +	0	0	0
LINE	L1	L1Md_T	chr5	10261982 10262645 -	0	0	0
SINE	Alu	B1_Mm	chr5	10364210 10364224 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	10379763 10379766 +	0	0	0.615324
SINE	Alu	B1_Mus2	chr5	10414354 10414369 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	10415779 10415783 +	0	0	0
SINE	Alu	B1_Mus2	chr5	10417251 10417266 -	0	0	0
SINE	Alu	B1_Mus2	chr5	10513426 10513441 -	0	0	0
SINE	Alu	B1_Mm	chr5	10519679 10519692 -	0	0	0
SINE	Alu	B1_Mm	chr5	10582260 10582275 -	0	0	0
SINE	Alu	B1_Mm	chr5	10610757 10610771 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	10610771 10610775 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	10611493 10611496 +	0	0	0
SINE	B2	B2_Mm1t	chr5	10629386 10629405 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	10629406 10629407 +	0	0	0
SINE	B2	B2_Mm1t	chr5	10629407 10629426 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	10629426 10629427 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	10640448 10640451 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	10668780 10668819 +	0	0	0
LTR	MaLR	MTB	chr10	22199858 22200191 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	10725838 10725843 +	0	0	0
SINE	Alu	B1_Mus1	chr5	10795538 10795551 -	0	0	0
SINE	Alu	B1_Mus2	chr5	10813602 10813616 +	0	0	0
Simple_repeat	Simple_repeat	(GAAAA)n	chr5	10813616 10813620 +	0	0	0
SINE	Alu	PB1D9	chr5	10831453 10831464 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	10831464 10831471 +	0	0	0
SINE	Alu	B1_Mus2	chr5	10843055 10843070 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr5	10843070 10843077 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr5	10888388 10888392 +	0	0	0
SINE	Alu	B1_Mus2	chr5	10911801 10911816 -	0	0	0
LTR	MaLR	MTA_Mm	chr5	11001030 11001070 +	0	0	0
SINE	Alu	B1_Mus1	chr5	11003603 11003618 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	11003618 11003622 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	11017650 11017652 +	0	0	0
SINE	B2	B2_Mm1a	chr5	11090510 11090529 -	0	0	0
LTR	MaLR	MTA_Mm	chr5	11094239 11094278 -	0	0	0
SINE	Alu	B1_Mus1	chr5	11174058 11174072 -	0	0	0
LTR	MaLR	MTA_Mm	chr5	11189168 11189207 -	0	0	0
SINE	Alu	B1_Mm	chr11	56285979 56286125 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	56286126 56286166 +	0.332194	0.380568	0
SINE	Alu	B1_Mus2	chr5	11264159 11264174 -	0	0	0
SINE	Alu	B1_Mus1	chr5	11493345 11493360 -	0	0	0
SINE	Alu	B1_Mur4	chr5	11500571 11500586 +	0	0	0
SINE	Alu	B1_Mm	chr5	11501286 11501300 +	0	0	0
SINE	Alu	B1_Mus2	chr5	11515496 11515510 +	0	0	0



LTR	ERVK	RMER4A	chr5	11533828 11533865 -	0	0	0
SINE	Alu	B1_Mm	chr5	11533867 11533882 -	0	0	0
SINE	Alu	B1_Mm	chr5	11544114 11544127 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	11544127 11544132 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	11545899 11545906 +	0	0	0
LINE	L1	L1Md_T	chr11	56613952 56620865 -	0	0	0
SINE	Alu	B1_Mus2	chr5	11575244 11575258 +	0	1.41763	0.615324
Simple_repeat	Simple_repeat	(A)n	chr5	11575258 11575261 +	0	0.708814	0
SINE	Alu	B1_Mus2	chr5	11628057 11628070 -	0	0	0
LTR	MaLR	MTA_Mm	chr11	56753799 56754193 +	0	0	0
SINE	Alu	B1_Mm	chr5	11648397 11648412 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	11648412 11648414 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	11653803 11653806 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	11676275 11676278 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	56815458 56815499 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	11714543 11714546 +	0	0	0
SINE	Alu	B1_Mus1	chr5	11725848 11725863 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	11725863 11725871 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	11748150 11748154 +	0	0	0
SINE	Alu	B1_Mus2	chr10	22329454 22329609 -	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	11893180 11893183 +	0	0	0
SINE	Alu	B1_Mus2	chr5	12010313 12010327 +	0	0	0
SINE	B2	B3	chr5	12054387 12054406 +	0	0	0
SINE	B2	B2_Mm2	chr5	12055451 12055470 -	0	0	0
LTR	MaLR	MTA_Mm	chr11	57289302 57289695 -	0	0	0
LTR	MaLR	MTA_Mm	chr11	57382178 57382563 -	0	0	0
SINE	Alu	PB1D9	chr11	57395550 57395642 -	0.332194	0	0
SINE	Alu	B1_Mus2	chr5	12195798 12195812 +	0	0	0
SINE	Alu	B1_Mus1	chr5	12201038 12201053 +	0	0	0
SINE	Alu	B1_Mus1	chr5	12254008 12254022 -	0	0	0
SINE	Alu	B1_Mm	chr5	12254544 12254559 -	0	0	0
SINE	Alu	B1_Mus1	chr5	12259009 12259024 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	12264242 12264245 +	0	0	0
SINE	Alu	B1_Mm	chr5	12265637 12265652 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	12265652 12265655 +	0.332194	0	0
SINE	Alu	B1_Mus2	chr11	57603254 57603399 -	0	0	0
SINE	B2	B2_Mm1a	chr5	12351133 12351152 -	0	0	0
SINE	Alu	B1_Mm	chr11	57639185 57639330 -	0.332194	0	0
LTR	MaLR	MTB	chr5	12417442 12417478 -	0	0	0
SINE	Alu	B1_Mus2	chr5	12450752 12450766 -	0	0	0
SINE	Alu	B1_Mus1	chr5	12480938 12480953 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	12480953 12480957 +	0	0	0
SINE	Alu	B1_Mm	chr5	12537998 12538012 +	0	0	0
SINE	Alu	B1_Mus2	chr5	12544245 12544260 -	0	0	0
LTR	MaLR	MTA_Mm	chr5	12567835 12567875 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	12582225 12582229 +	0	0	0
SINE	Alu	B1_Mus2	chr5	12583795 12583809 -	0	0	0
LTR	MaLR	MTA_Mm	chr5	12585652 12585691 +	0	0	0
SINE	Alu	B1_Mm	chr11	57792362 57792507 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	57799091 57799121 +	0	0	0
SINE	Alu	B1_Mur4	chr5	12601079 12601088 +	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chr5	12601088 12601093 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	12673716 12673719 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	12718284 12718287 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	12737537 12737542 +	0	0	0
LTR	MaLR	MTB	chr5	12740200 12740238 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	12760821 12760824 +	0	0	0
SINE	Alu	B1_Mm	chr11	57994097 57994242 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	12895650 12895653 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	12917873 12917911 +	0	0	0
SINE	Alu	B1_Mus1	chr5	12958554 12958568 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	12958568 12958572 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	12974722 12974761 -	0	0	0
SINE	Alu	B1_Mm	chr5	12987624 12987637 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	12987637 12987640 +	0	0	0
LTR	MaLR	MTA_Mm	chr5	13007832 13007871 -	0	0	0
SINE	Alu	B1_Mm	chr5	13063677 13063692 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr5	13063692 13063695 +	0.332194	0	0
SINE	Alu	B1_Mus1	chr11	58208775 58208921 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	58208922 58208946 +	0	0	0
SINE	B4	ID_B1	chr5	13190230 13190251 -	0	0	0
LTR	MaLR	MTA_Mm	chr5	13308131 13308171 -	0	0	0
SINE	Alu	PB1D10	chr10	22573493 22573609 -	0	0	0
SINE	Alu	B1_Mm	chr11	58675838 58675982 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr11	58675983 58676005 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr5	13395145 13395151 +	5.90476	1.41763	2.53122
SINE	Alu	B1_Mus2	chr5	13481937 13481952 +	0	0	0
SINE	Alu	B1_Mus2	chr5	13504615 13504629 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	13504629 13504632 +	0	0	0
SINE	Alu	B1_Mm	chr11	59047127 59047278 -	0	0	0
SINE	Alu	B1_Mus1	chr5	13587703 13587718 -	0	0	0
SINE	Alu	B1_Mus1	chr5	13615815 13615829 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	13615830 13615832 +	0.332194	0	0
SINE	Alu	B1_Mm	chr5	13643059 13643073 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	13643073 13643076 +	0	0	0
SINE	B4	ID_B1	chr5	13664905 13664924 -	0.591053	0	0
SINE	Alu	B1_Mm	chr5	13709832 13709847 -	0	0	0
SINE	Alu	B1_Mus1	chr11	59287185 59287332 -	0	0	0
SINE	Alu	B1_Mm	chr5	13753219 13753234 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	59337230 59337265 +	0	0	0
SINE	Alu	B1_Mm	chr5	13792024 13792039 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	13818650 13818654 +	0	0	0
SINE	Alu	B1_Mm	chr5	13898524 13898538 -	0	0	0
SINE	Alu	B1_Mm	chr5	13966478 13966492 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	13966492 13966494 +	0	0	0
Simple_repeat	Simple_repeat	(TAA)n	chr5	13966494 13966497 +	0	0	0
SINE	Alu	PB1D10	chr5	14031586 14031596 -	0	0	0
SINE	B2	B2_Mm1a	chr11	59714965 59715160 -	0.591053	0.380568	0
SINE	Alu	B1_Mus1	chr11	59720335 59720437 -	0	0	0
SINE	Alu	B1_Mm	chr5	14087719 14087731 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr5	14087731 14087738 +	0	0	0
SINE	Alu	B1_Mus2	chr5	14092443 14092458 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	14092458 14092461 +	0	0	0
SINE	Alu	B1_Mm	chr5	14092663 14092677 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	14092677 14092679 +	0	0	0
SINE	Alu	B1_Mus2	chr5	14109164 14109178 +	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr5	14109178 14109183 +	0.332194	0.380568	0.342623
Simple_repeat	Simple_repeat	(A)n	chr5	14283082 14283087 +	0	0	0.342623
SINE	B4	ID_B1	chr5	14317004 14317024 -	0	0	0
SINE	Alu	B1_Mus1	chr5	14362156 14362170 -	0	0.380568	0
SINE	Alu	B1_Mus1	chr5	14388808 14388823 -	0	0	0
SINE	Alu	B1_Mus2	chr5	14483514 14483528 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr5	14483528 14483538 +	0	0	0.615324
SINE	Alu	B1_Mus1	chr5	14519740 14519754 +	0	0	0
SINE	Alu	B1_Mus2	chr5	14574046 14574061 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	14574061 14574064 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	14605469 14605472 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	14625742 14625746 +	0	0	0.615324
SINE	Alu	B1_Mus2	chr5	14701206 14701220 -	1.18211	0.708814	1.23065
SINE	Alu	B1_Mus1	chr5	14719111 14719126 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	60450611 60450650 +	0	0	0.342623
SINE	Alu	B1_Mus1	chr5	14738861 14738876 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	14738876 14738878 +	0	0	0
SINE	Alu	B1_Mm	chr5	14753276 14753290 -	0	0	0
LTR	MaLR	MTB	chr5	14754267 14754307 -	0	0	0.615324
SINE	Alu	B1_Mus1	chr5	14763900 14763914 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	14777545 14777549 +	0	0	0
SINE	Alu	B1_Mus1	chr5	14779089 14779104 -	0	0	0
SINE	Alu	B1_Mm	chr5	14813664 14813679 -	0	0	0
SINE	Alu	B1_Mur4	chr5	14836583 14836597 -	0	0	0
LTR	MaLR	MTA_Mm	chr5	14967052 14967091 -	0	0	0
SINE	Alu	B1_Mm	chr5	14974125 14974139 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	14974139 14974142 +	0	0	0
SINE	Alu	B1_Mus1	chr5	14975644 14975658 -	0	0	0.615324

LTR	MaLR	MTB	chr5	14984086 14984126 +	0	0	0
SINE	Alu	B1_Mus2	chr11	60782836 60782982 -	0	0	0
SINE	Alu	B1_Mus2	chr5	15171229 15171243 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	15171244 15171246 +	0	0	0
SINE	Alu	B1_Mm	chr5	15171584 15171599 -	0	0	0
SINE	Alu	B1_Mm	chr5	15228304 15228318 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	15228318 15228320 +	0	0	0
LINE	L1	L1_Mus4	chr5	15241732 15242262 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr5	15246493 15246497 +	0	0	0.342623
LTR	MaLR	MTA_Mm	chr5	15250322 15250362 -	0	0	0
LTR	MaLR	MTA_Mm	chr5	15250471 15250511 -	0	0	0
LTR	MaLR	MTA_Mm	chr6	3398148 3398543 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	3399641 3400036 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	4388908 4388935 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	5402618 5403012 +	0	0	0
SINE	Alu	PB1D9	chr6	5508419 5508519 -	0	0	0
SINE	Alu	B1_Mus2	chr6	5634784 5634929 -	0	0	0
LTR	MaLR	MTA_Mm	chr6	5635993 5636388 +	0	0	0
SINE	Alu	B1_Mm	chr6	6298543 6298690 -	0.591053	0	0
SINE	Alu	B1_Mus1	chr6	6668116 6668263 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	6668264 6668290 +	0	0	0
SINE	Alu	B1_Mus2	chr6	7223290 7223435 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	7223436 7223472 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	7256755 7256830 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	7689470 7689501 +	0	0	0
LTR	ERV1	RLTR24	chr6	7703849 7704360 +	0	0	0
SINE	Alu	B1_Mus1	chr6	7713447 7713594 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	61328999 61329058 +	1.18211	0	1.84597
Simple_repeat	Simple_repeat	(A)n	chr6	9228117 9228174 +	0	0	0
SINE	Alu	B1_Mus2	chr6	9282774 9282915 -	0	0	0
SINE	Alu	B1_Mus1	chr11	61405169 61405302 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	61405303 61405335 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	9543303 9543341 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	10749687 10750081 -	0	0	0
LINE	L1	L1Md_F2	chr6	10754546 10759533 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	11003273 11003668 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	11142889 11142916 +	0	0	0
LTR	ERVK	BGLII	chr6	11293183 11293638 -	0	0	0
LINE	L1	L1Md_T	chr6	11692603 11699447 -	0	0	0
SINE	Alu	B1_Mus2	chr6	12086938 12087084 +	0.591053	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr6	12087085 12087107 +	0	0	0
LINE	L1	L1Md_F2	chr6	12173286 12179332 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	13349775 13349815 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	13431456 13431847 -	0	0	0
LTR	MaLR	MTA_Mm	chr6	13668024 13668422 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	14191269 14191296 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	14440562 14440598 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	15474851 15474889 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	15505807 15505833 +	0	0	0
SINE	Alu	B1_Mm	chr11	61755420 61755565 -	0	0	0
SINE	Alu	B1_Mus2	chr6	15846488 15846605 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	16281944 16281972 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	16983599 16983641 +	0	0.708814	0.615324
Simple_repeat	Simple_repeat	(A)n	chr6	17536925 17536951 +	0	0	0
SINE	Alu	B1_Mus1	chr6	17655804 17655947 +	0	0	0
SINE	Alu	B1_Mus2	chr6	17694394 17694540 -	0	0	0
LINE	L1	L1Md_T	chr6	18112286 18119018 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	18483469 18483506 +	0	0	0
LINE	L1	L1Md_F2	chr6	18957307 18962717 -	0	0	0
LTR	MaLR	MTA_Mm	chr6	19357503 19357896 -	0	0	0
LINE	L1	L1Md_F	chr6	19616859 19621345 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	19837747 19837775 +	0	0	0
SINE	Alu	B1_Mus2	chr6	19847553 19847698 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	19893176 19893228 +	0	0	0
LINE	L1	L1Md_F3	chr6	19927306 19933485 -	0	0	0
LTR	MaLR	MTA_Mm	chr6	20225638 20226031 +	0	0	0
LINE	L1	L1_Mus3	chr6	20417553 20422686 +	0	0	0

LINE	L1	L1Md_F2	chr6	20557760 20562957 +	0	0	0
LINE	L1	L1Md_F2	chr6	20578973 20585158 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	21806341 21806369 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	21882690 21883084 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	22713729 22714124 -	0	0	0
SINE	Alu	B1_Mus2	chr6	25552394 25552540 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	25552541 25552563 +	0	0	0
SINE	Alu	B1_Mus2	chr6	26241579 26241726 -	0	0	0
SINE	Alu	B1_Mus1	chr6	26343175 26343322 -	0	0	0
LTR	MaLR	MTA_Mm	chr6	26546873 26547268 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	62387588 62387626 +	0	0	0
SINE	Alu	B1_Mus2	chr6	26767094 26767242 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	26767243 26767264 +	0	0	0
LINE	L1	L1Md_T	chr6	26824939 26831447 -	0	0	0
SINE	Alu	B1_Mus1	chr6	26896625 26896771 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	26896772 26896811 +	0	0	0
SINE	Alu	B1_Mm	chr11	62413844 62413989 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	62413990 62414012 +	0.923247	0	0
SINE	Alu	B1_Mm	chr6	28258854 28258999 -	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr6	28881069 28881095 +	0	0	0
SINE	Alu	B1_Mm	chr6	29167671 29167819 +	0	0.380568	0.342623
Simple_repeat	Simple_repeat	(A)n	chr6	29379569 29379608 +	0	0	0
SINE	Alu	B1_Mm	chr6	29575307 29575453 -	0	0	0
SINE	Alu	B1_Mus1	chr6	29670647 29670793 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	29670794 29670816 +	0	0	0
SINE	Alu	B1_Mus2	chr6	29990728 29990874 -	0	0	0
SINE	B2	B2_Mm1a	chr6	29992086 29992277 +	0	0	0
SINE	Alu	B1_Mur3	chr6	30106239 30106379 +	0	0	0
SINE	Alu	B1_Mus1	chr6	30155395 30155540 -	0	0	0
SINE	Alu	B1_Mus2	chr6	30273183 30273322 -	0	0	0
SINE	Alu	B1_Mus2	chr6	30307131 30307257 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	30307258 30307280 +	0	0	0
SINE	Alu	B1_Mus2	chr6	31462125 31462270 +	4.41242	2.17876	3.07662
Simple_repeat	Simple_repeat	(A)n	chr6	31462271 31462300 +	5.07681	2.17876	2.87384
Simple_repeat	Simple_repeat	(A)n	chr6	31890244 31890278 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	32719156 32719547 +	0	0	0
SINE	Alu	B1_Mm	chr11	63142516 63142664 -	0	0	0
SINE	Alu	B1_Mus2	chr11	63170006 63170150 -	0	0	0
LINE	L1	L1Md_T	chr6	35289469 35296397 -	0	0	0
SINE	Alu	B1_Mus1	chr6	35873431 35873557 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	35873558 35873594 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	35944656 35944705 +	0	0	0
SINE	Alu	B1_Mus1	chr11	63321623 63321747 -	0	0	0
SINE	Alu	B1_Mus1	chr6	36481254 36481404 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	36481405 36481446 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	36502702 36503095 +	0	0	0
SINE	Alu	B1_Mm	chr6	36580608 36580752 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr6	36580753 36580923 +	0	0	0
LTR	MaLR	MTB	chr6	37545469 37545856 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	37549963 37550355 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	37666145 37666536 -	0	0	0
SINE	Alu	B1_Mus2	chr6	37710966 37711112 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	37711113 37711153 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr6	37743689 37743835 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	37743836 37743894 +	0.332194	0	0
LTR	MaLR	MTA_Mm	chr6	37772082 37772477 -	0	0	0
SINE	Alu	B1_Mus2	chr6	38256162 38256308 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	38256309 38256332 +	0	0	0
SINE	Alu	B1_Mus2	chr6	38257914 38258060 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	38307100 38307126 +	0	0	0
SINE	Alu	B1_Mus2	chr6	38382693 38382815 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	38382816 38382841 +	0	0	0
SINE	Alu	B1_Mus2	chr6	38463934 38464084 -	0	0	0
SINE	Alu	B1_Mus2	chr6	38478814 38478962 +	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr6	38478963 38478987 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	39700292 39700365 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	39764714 39765109 +	0	0	0

LTR	MaLR	MTA_Mm	chr6	39785203 39785597 +	0	0	0
LTR	MaLR	MTB	chr6	40208450 40208849 -	0	0	0
LTR	MaLR	MTA_Mm	chr6	40272308 40272690 +	0	0	0
SINE	Alu	B1_Mm	chr6	41293714 41293859 -	0	0	0
SINE	Alu	B1_Mm	chr6	42364540 42364685 -	0	0	0
LTR	MaLR	MTA_Mm	chr6	42367216 42367592 -	0	0	0
LTR	MaLR	MTA_Mm	chr6	42387774 42388167 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	42389260 42389653 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	42663259 42663657 +	0	0	0
SINE	Alu	B1_Mus2	chr6	42825413 42825559 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	42825560 42825599 +	0	0	0
SINE	Alu	B1_Mm	chr6	43986930 43987067 -	0	0	0
SINE	Alu	B1_Mus2	chr6	44277313 44277459 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	44413856 44414256 -	0	0	0
LINE	L1	L1Md_F2	chr6	44503082 44509533 -	0	0	0
LTR	MaLR	MTA_Mm	chr11	64502426 64502809 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	47263151 47263546 -	0	0	0
LTR	MaLR	MTA_Mm	chr6	47301197 47301592 -	0	0	0
SINE	Alu	B1_Mm	chr6	47336067 47336212 -	0	0	0
SINE	Alu	B1_Mus1	chr6	47366480 47366625 -	0	0	0
scRNA	scRNA	4.5SRNA	chr6	47600761 47600854 -	0.332194	1.32017	0.342623
scRNA	scRNA	4.5SRNA	chr6	47698497 47698590 -	3.13357	0	2.32725
scRNA	scRNA	4.5SRNA	chr6	47707665 47707758 -	0	0	0
scRNA	scRNA	4.5SRNA	chr6	47713263 47713356 -	0.332194	0.380568	4.24948
scRNA	scRNA	4.5SRNA	chr6	47717586 47717660 -	0	0	0
SINE	Alu	PB1D10	chr6	47717592 47717687 -	0	0	0
scRNA	scRNA	4.5SRNA	chr6	47721993 47722067 -	0	0	0
SINE	Alu	PB1D10	chr6	47721999 47722094 -	0	0	0
scRNA	scRNA	4.5SRNA	chr6	47728245 47728338 -	5.81523	3.69765	3.56423
SINE	Alu	B1_Mus2	chr6	47864837 47864982 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	48239128 48239167 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	48305289 48305332 +	0	0	0
SINE	Alu	B1_Mm	chr6	48624566 48624710 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	48624711 48624738 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	64766419 64766454 +	0	0	0
SINE	Alu	B1_Mus1	chr6	50149669 50149807 -	0	0	0
SINE	Alu	B1_Mur4	chr6	50695445 50695603 +	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chr6	50695604 50695666 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	50980689 50980715 +	0	0	0
SINE	Alu	B1_Mus1	chr6	51448730 51448876 -	0	0	0
SINE	Alu	B1_Mm	chr6	51603437 51603594 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	51603595 51603616 +	0	0	0
SINE	Alu	B1_Mur2	chr6	52018187 52018324 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	52018325 52018358 +	0	0	0
SINE	Alu	B1_Mur1	chr6	52059063 52059207 -	0	0	0
SINE	Alu	B1_Mm	chr6	52084386 52084531 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	52084532 52084566 +	0	0	0
SINE	Alu	B1_Mus1	chr6	52355513 52355661 -	0	0	0
SINE	Alu	B1_Mus1	chr6	52402920 52403067 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	52403068 52403091 +	0.332194	0	0
SINE	Alu	B1_Mm	chr11	65158937 65159082 -	0	0	0
SINE	Alu	B1_Mm	chr6	52627790 52627936 -	0	0	0
LTR	MaLR	MTA_Mm	chr6	53414214 53414596 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	53415694 53416076 +	0.332194	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr6	53488980 53489016 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	53715270 53715295 +	0	0	0
SINE	Alu	B1_Mus1	chr6	54679987 54680134 -	0	0	0
SINE	Alu	B1_Mus2	chr6	55568502 55568647 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	55568648 55568697 +	0	0	0
SINE	Alu	B1_Mus2	chr6	56664136 56664281 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	56664282 56664304 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	57162240 57162281 +	0	0	0
SINE	Alu	B1_Mm	chr6	57197913 57198058 -	0	0	0
SINE	Alu	B1_Mm	chr6	57516260 57516406 +	0	0	0
SINE	Alu	B1_Mus1	chr6	59708695 59708841 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	59708842 59708887 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	60128013 60128050 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr6	60201082 60201147 +	0	0	0
SINE	Alu	B1_Mus2	chr10	23489873 23490019 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	60943833 60943868 +	0	0	0
SINE	Alu	B1_Mus2	chr6	62423244 62423395 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	62551059 62551098 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	23510118 23510150 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	64681543 64681580 +	0	0	0
LINE	L1	L1Md_T	chr6	65152421 65158900 -	0	0	0
SINE	Alu	B1_Mm	chr6	65158929 65159064 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	65159065 65159093 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	65253368 65253400 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	65254918 65254956 +	0.332194	0	0
LINE	L1	L1VL4	chr6	66078485 66083685 -	0	0	0
LINE	L1	L1VL4	chr6	66191313 66194086 +	0	0	0
LINE	L1	L1Md_T	chr6	66408595 66414529 -	0	0	0
SINE	Alu	B1_Mus1	chr6	67209230 67209383 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	67209384 67209421 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	67451058 67451096 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	67487505 67487544 +	0	0	0
LINE	L1	L1Md_T	chr6	67526312 67533627 -	0	0	0
LINE	L1	L1Md_F3	chr6	67544935 67550133 +	0	0	0
SINE	Alu	B1_Mus2	chr6	67581328 67581474 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	67884450 67884845 -	0	0	0
LINE	L1	L1Md_F2	chr6	68223311 68230134 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	68357727 68357774 +	0	0	0
LINE	L1	L1Md_F2	chr6	68362603 68367970 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	68711205 68711597 +	0	0	0
SINE	Alu	B1_Mus2	chr6	69993251 69993397 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	69993398 69993438 +	0	0	0
SINE	Alu	B1_Mm	chr6	70244172 70244317 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	70760433 70760471 +	0	0	0
LTR	MaLR	MTA_Mm	chr10	23550230 23550625 -	0.332194	0	0
LTR	MaLR	MTA_Mm	chr10	23551723 23552118 -	0	0	0
SINE	Alu	B1_Mur2	chr6	70997461 70997603 -	0	0	0
SINE	Alu	B1_Mus2	chr6	71443325 71443470 -	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	71780856 71780908 +	0	0	0
SINE	Alu	B1_Mus2	chr6	72401690 72401835 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	72790988 72791025 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	74392703 74392737 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	74636848 74636895 +	0	0	0
SINE	Alu	B1_Mus2	chr6	74716404 74716548 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	74881963 74882009 +	0	0	0
LINE	L1	L1Md_F2	chr6	75082287 75087307 -	0	0	0
SINE	Alu	B1_Mus1	chr6	75212977 75213122 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	75213123 75213152 +	0	0	0
SINE	Alu	B1_Mus1	chr6	75538495 75538642 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	75773684 75773734 +	0	0	0
SINE	Alu	B1_Mus1	chr6	75926108 75926233 -	0	0	0
LTR	MaLR	MTA_Mm	chr6	75965375 75965756 +	0	0	0
SINE	Alu	B1_Mm	chr6	76096400 76096537 -	0	0	0
SINE	Alu	B1_Mm	chr6	76360814 76360958 -	0	0	0
SINE	Alu	B1_Mus1	chr6	77947778 77947902 -	0	0	0
SINE	Alu	B1_Mm	chr6	78036656 78036801 +	0	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr6	78036802 78036844 +	0	0	0
LTR	MaLR	MTA_Mm	chr10	23627364 23627758 +	0	0	0
LINE	L1	L1Md_T	chr6	78826365 78832760 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	78991474 78991510 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr6	79083948 79083989 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	79084052 79084088 +	0	0	0
SINE	Alu	B1_Mm	chr6	79523318 79523464 +	0	0	0
SINE	Alu	B1_Mus2	chr6	80167517 80167663 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	80167664 80167713 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr6	80341790 80341820 +	0	0	0
SINE	Alu	B1_Mus2	chr6	81175839 81175984 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	81175985 81176013 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	81440105 81440498 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	81705715 81705746 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr6	82566330 82566359 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	82614424 82614819 -	0	0	0
SINE	Alu	B1_Mus1	chr6	82618936 82619083 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	83061853 83061901 +	0.923247	0.380568	0.342623
SINE	Alu	B1_Mm	chr6	83740795 83740941 -	0	0	0
SINE	Alu	B1_Mm	chr6	83754833 83754978 -	0	0	0
SINE	Alu	B1_Mur3	chr6	83802926 83803080 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	83803081 83803108 +	0	0	0
SINE	Alu	B1_Mm	chr6	83810485 83810631 +	0	0	0
SINE	Alu	B1_Mus1	chr6	85341636 85341782 -	0	0	0
SINE	B2	B2_Mm1t	chr10	23711359 23711550 -	0	0	0
SINE	Alu	B1_Mus2	chr6	86252188 86252340 -	0	0	0
SINE	Alu	B1_Mus1	chr6	86471987 86472113 +	0	0	0
SINE	Alu	B1_Mm	chr6	86780428 86780574 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	68463611 68463665 +	0	0	0
SINE	Alu	PB1D9	chr6	87892250 87892338 +	0	0	0.957947
Simple_repeat	Simple_repeat	(A)n	chr6	87892339 87892362 +	0	0	0.957947
SINE	Alu	B1_Mus1	chr6	87990743 87990890 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	87990891 87990917 +	0	0	0
SINE	Alu	B1_Mus2	chr6	88020690 88020835 -	0	0	0
SINE	Alu	B1_Mm	chr6	88061717 88061862 -	0	0	0
SINE	Alu	B1_Mm	chr6	88129619 88129765 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	88547940 88547974 +	0	0	0
SINE	Alu	B1_Mus2	chr6	89042905 89043037 -	0	0	0
SINE	Alu	B1_Mus1	chr6	89815369 89815515 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	89815516 89815545 +	0	0	0
LINE	L1	L1Md_T	chr6	89977512 89985872 -	0	0	0
SINE	Alu	B1_Mm	chr6	90135267 90135414 +	0	0	0
SINE	Alu	B1_Mus2	chr6	90466671 90466817 -	0	0	0
SINE	Alu	B1_Mur4	chr6	91255915 91256064 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	91256065 91256126 +	0	0	0
SINE	Alu	B1_Mur2	chr6	91497608 91497747 -	0	0	0
SINE	B2	B2_Mm1a	chr6	91614161 91614353 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	91614354 91614376 +	0	0	0
SINE	Alu	B1_Mus1	chr6	92185718 92185855 -	0	0	0
SINE	Alu	B1_Mus2	chr6	92201391 92201537 -	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr10	23804521 23804555 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	92678055 92678091 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	92681757 92681811 +	0	0	0
SINE	Alu	B1_Mm	chr6	93186804 93186956 +	0	0.708814	1.23065
Simple_repeat	Simple_repeat	(A)n	chr6	93186957 93186982 +	0	0.708814	1.23065
SINE	B2	B2_Mm1a	chr6	93502380 93502572 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	93502573 93502604 +	0	0.380568	0.957947
Simple_repeat	Simple_repeat	(A)n	chr11	68892677 68892714 +	0.827951	0	0
SINE	Alu	B1_Mus2	chr6	94945418 94945564 -	0	0	0
SINE	Alu	B1_Mus1	chr11	69109954 69110096 -	0.591053	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr6	98012029 98012066 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	98175900 98176294 -	0	0	0
LTR	MaLR	MTA_Mm	chr6	98177393 98177787 -	0	0	0
LTR	MaLR	MTA_Mm	chr6	98621425 98621825 -	0	0	0
SINE	Alu	B1_Mm	chr6	98846898 98847044 -	0	0	0
SINE	Alu	B1_Mm	chr6	98858129 98858273 -	0	0	0
SINE	Alu	B1_Mm	chr6	98863383 98863508 -	0	0.380568	0.342623
SINE	B2	B2_Mm1a	chr6	98863539 98863731 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	98863732 98863756 +	0	0.380568	0.342623
SINE	Alu	B1_Mm	chr6	99422805 99422947 -	1.18211	0.761137	0
SINE	Alu	B1_Mus2	chr6	10030843 10030855 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	10030918 10030923 +	0.332194	0	0
SINE	Alu	B1_Mm	chr6	10031023 10031035 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	10037433 10037436 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	69445885 69445909 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	10278848 10278887 +	0	0	0
SINE	Alu	B1_Mm	chr6	10292689 10292703 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	10292704 10292706 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	10302296 10302300 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	10309917 10309955 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	10310064 10310103 +	0	0	0

SINE	Alu	B1_Mus2	chr6	10368402	10368417	-	0	0	0
LINE	L1	L1Md_T	chr6	10420707	10421441	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	10436064	10436066	+	0	0	0
LINE	L1	L1_Mus2	chr6	10486022	10486490	-	0	0	0
LINE	L1	L1Md_A	chr6	10502225	10502864	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	10544963	10544967	+	0	0	0
SINE	Alu	B1_Mus1	chr6	10758054	10758069	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	10758069	10758073	+	0	0	0
LINE	L1	L1Md_T	chr6	10762080	10762768	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	10792671	10792674	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	10794434	10794437	+	0	0	0
SINE	Alu	B1_Mus1	chr6	10862940	10862954	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	10862955	10862957	+	0	0	0
SINE	Alu	B1_Mm	chr6	10934899	10934908	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	10934908	10934911	+	0	0	0
SINE	Alu	B1_Mm	chr6	10981118	10981133	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	10981133	10981136	+	0	0	0
SINE	Alu	B1_Mus1	chr6	10989704	10989720	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	10989720	10989727	+	0	0	0
SINE	Alu	B1_Mur1	chr6	11058576	11058589	-	4.00689	1.41763	3.07662
SINE	Alu	B1_Mm	chr6	11222162	11222177	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	11222177	11222180	+	0	0	0
LTR	MaLR	MTA_Mm	chr6	11248245	11248284	-	0	0	0
SINE	Alu	B1_Mus2	chr10	24044488	24044632	+	0	0	0
SINE	Alu	B1_Mm	chr6	11297670	11297685	-	0	0	0
Simple_repeat	Simple_repeat	(GAA)n	chr10	24044633	24044730	+	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr6	11315641	11315644	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	11338872	11338875	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	70245551	70245585	+	0	0	0
SINE	Alu	B1_Mus2	chr6	11358601	11358616	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	11358616	11358619	+	0	0	0
SINE	Alu	B1_Mus1	chr6	11366335	11366350	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	11366350	11366353	+	0	0	0
LTR	ERVL	RLTR28	chr6	11434381	11434435	-	0	0	0
SINE	Alu	B1_Mus2	chr11	70337152	70337298	-	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr6	11499446	11499449	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	11526309	11526312	+	0	0	0
SINE	Alu	B1_Mus1	chr6	11544149	11544164	-	0	0	0
SINE	Alu	B1_Mm	chr6	11670711	11670724	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	11670724	11670726	+	0	0	0
SINE	Alu	B1_Mus2	chr6	11678840	11678854	-	0	0	0
Low_complexity	Low_complexity	AT_rich	chr6	11704431	11704435	+	0	0	0
LTR	MaLR	MTA_Mm	chr6	11770437	11770476	-	0	0	0
SINE	Alu	B1_Mus2	chr6	11777948	11777963	-	0	0	0
LTR	MaLR	MTE2b	chr6	11778729	11778760	-	0	0	0
SINE	Alu	B1_Mm	chr11	70566596	70566742	-	0	0	0
LINE	L1	L1Md_T	chr6	11815915	11816580	-	0	0	0
SINE	Alu	B1_Mus2	chr11	70652637	70652782	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	70652783	70652840	+	0	0	0
SINE	Alu	B1_Mus1	chr6	12053471	12053486	+	0	0	0
SINE	Alu	B1_Mus2	chr6	12053989	12054004	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	12054004	12054006	+	0	0	0
SINE	Alu	B1_Mus2	chr6	12078380	12078394	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	12078394	12078397	+	0	0	0
SINE	Alu	B1_Mus1	chr6	12275385	12275397	-	0	0	0
LINE	L1	L1_Mus3	chr6	12385252	12385662	-	0	0	0
SINE	Alu	B1_Mus2	chr6	12434529	12434543	-	0	0	0.615324
LTR	MaLR	MTC	chr6	12443900	12443937	-	0	0	0
SINE	Alu	B1_Mus2	chr6	12453980	12453994	-	0	0	0
SINE	Alu	B1_Mm	chr6	12489692	12489706	+	0	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr6	12489706	12489708	+	0	0.380568	0
SINE	Alu	B1_Mus2	chr6	12515480	12515495	-	0	0	0
SINE	Alu	B1_Mus1	chr6	12524812	12524826	-	0	0	0
SINE	Alu	B1_Mus1	chr6	12538431	12538444	-	0	0	0
SINE	Alu	B1_Mus2	chr6	12539659	12539673	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	12539673	12539676	+	0	0	0
SINE	Alu	B1_Mus2	chr6	12692578	12692593	-	0	0.708814	0



SINE	Alu	B1_Mm	chr6	12707422	12707437 +	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr6	12720783	12720785 +	0	0	0
SINE	B2	B3	chr6	12747352	12747373 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	12747373	12747377 +	0	0	0
SINE	Alu	B1_Mm	chr6	12781264	12781279 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	12781279	12781281 +	0	0	0
LTR	MaLR	MTA_Mm	chr6	12929249	12929288 +	0	0	0
SINE	Alu	B1_Mur4	chr6	12948073	12948088 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr6	12948088	12948093 +	0	0	0
SINE	Alu	B1_Mus2	chr6	12951819	12951833 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	12951833	12951836 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr6	12963208	12963217 +	0	0	0
SINE	Alu	B1_Mus2	chr6	13104694	13104709 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	13144071	13144074 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	13153949	13153953 +	0	0	0
SINE	Alu	B1_Mus2	chr6	13241693	13241708 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	13254354	13254357 +	0	0	0
LINE	L1	L1Md_T	chr6	13256515	13257073 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	13284758	13284761 +	0	0	0
SINE	Alu	B1_Mm	chr6	13324141	13324156 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	13324156	13324163 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr6	13388129	13388134 +	0	0	0
SINE	Alu	B1_Mm	chr6	13474556	13474570 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	13492149	13492152 +	0	0	0
SINE	Alu	PB1D10	chr6	13521964	13521975 -	0	0	0
SINE	Alu	B1_Mm	chr6	13615474	13615486 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	13615486	13615489 +	0	0	0
SINE	B2	B2_Mm1a	chr6	13631667	13631686 +	0	0.708814	0
LTR	MaLR	MTA_Mm	chr6	13820846	13820885 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	13872260	13872263 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	13905718	13905721 +	0	0	0
SINE	Alu	PB1D7	chr6	14304901	14304913 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	14304913	14304918 +	0	0	0
SINE	Alu	B1_Mm	chr6	14309042	14309056 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	14309056	14309059 +	0	0	0
SINE	Alu	B1_Mus2	chr6	14309980	14309995 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr6	14309995	14310003 +	0	0	0
SINE	Alu	B1_Mus1	chr6	14311420	14311435 -	0	0	0
SINE	Alu	B1_Mur3	chr6	14456418	14456433 -	0	0	0
SINE	Alu	B1_Mur4	chr11	72966069	72966205 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	14564588	14564590 +	0	0	0
LTR	MaLR	MTB	chr6	14594302	14594342 -	0.332194	0	0
SINE	Alu	B1_Mus1	chr6	14663575	14663587 -	0.332194	0	0
SINE	Alu	B1_Mus1	chr11	73141557	73141703 -	0	0	0
LTR	MaLR	MTA_Mm	chr6	14861150	14861189 -	0	0	0
LTR	MaLR	MTA_Mm	chr6	14861298	14861337 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	14889687	14889691 +	0.591053	0	0
SINE	Alu	B1_Mus1	chr6	14892199	14892214 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	14892214	14892216 +	0	0	0
SINE	Alu	B1_Mm	chr6	14922941	14922955 -	0	0	0
SINE	B2	B2_Mm1t	chr6	14932651	14932670 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr6	14932671	14932673 +	0	0	0
SINE	Alu	B1_Mus1	chr7	3199359	3199500 -	0	0	0
SINE	Alu	B1_Mus2	chr7	3205507	3205653 -	0	0	0
SINE	Alu	B1_Mus2	chr7	3211949	3212068 -	0	0	0
SINE	Alu	B1_Mus1	chr7	3273079	3273220 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	3273221	3273243 +	0	0	0
LTR	MaLR	MTA_Mm	chr7	3710429	3710825 +	0.332194	0	0
LTR	MaLR	MTA_Mm	chr7	3770490	3770884 +	0	0	0
LINE	L1	L1Md_T	chr7	4137820	4144724 -	0	0	0.342623
LTR	MaLR	MTA_Mm	chr7	4933547	4933886 -	0	0	0
SINE	Alu	B1_Mm	chr7	4971391	4971537 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	4971538	4971560 +	0.332194	0	0
LINE	L1	L1_Mus3	chr7	5371037	5376288 -	0	0	0
SINE	Alu	B1_Mus2	chr7	6118645	6118791 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr7	6296486	6296514 +	0	0	0
SINE	Alu	B1_Mm	chr11	74172177	74172323 -	0	0	0

SINE	Alu	B1_Mm	chr7	7009411	7009543	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	7458415	7458451	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	8002903	8002941	+	0	0	0
LINE	L1	L1Md_F2	chr7	9051349	9056456	-	0	0	0
LINE	L1	L1Md_F2	chr7	9342690	9347717	+	0	0	0
LINE	L1	L1Md_F2	chr7	9709170	9714277	-	0	0	0
LTR	MaLR	MTA_Mm-int	chr7	10789210	10790414	+	0	0	0
SINE	Alu	B1_Mus1	chr11	74420400	74420547	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	74420548	74420569	+	0.332194	0	0
LTR	MaLR	MTA_Mm	chr11	74421947	74422327	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	11492226	11492278	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	11897855	11897893	+	0	0	0
SINE	Alu	B1_Mus1	chr7	13505882	13506029	-	0.591053	0	0.615324
SINE	Alu	B1_Mus2	chr7	13537061	13537207	-	0.591053	0	0
SINE	Alu	B1_Mm	chr7	13654283	13654429	-	0	0	0
SINE	Alu	B1_Mur4	chr11	74569355	74569494	-	0	0	0
LINE	L1	L1Md_A	chr7	14001064	14007285	-	0.332194	0	0
SINE	Alu	B1_Mm	chr10	24560456	24560600	-	0	0	0
LTR	MaLR	MTA_Mm	chr7	15221308	15221703	-	0.591053	0	0
LTR	MaLR	MTA_Mm	chr7	15222799	15223194	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	15550974	15551023	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	16652918	16652958	+	0	0	0
SINE	Alu	B1_Mur4	chr7	17346033	17346175	+	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr7	17346176	17346205	+	0	0	0
LTR	MaLR	MTB	chr7	17511449	17511846	-	0	0	0
SINE	Alu	B1_Mus2	chr7	17541457	17541580	-	0	0	0
LTR	MaLR	MTA_Mm	chr7	18373381	18373776	+	0	0	0
SINE	Alu	B1_Mm	chr7	19339092	19339247	+	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr7	19500050	19500079	+	0	0	0
SINE	Alu	B1_Mus2	chr7	19833492	19833636	+	1.18211	0	1.23065
SINE	Alu	B1_Mm	chr11	74988037	74988189	+	3.0286	1.08938	0.957947
Simple_repeat	Simple_repeat	(A)n	chr11	74988190	74988240	+	3.28746	0.708814	0.685246
Simple_repeat	Simple_repeat	(TTTA)n	chr7	20522015	20522044	+	0	0	0
LINE	L1	L1Md_F2	chr7	20794688	20799926	+	0	0	0.342623
SINE	Alu	B1_Mm	chr11	75109298	75109449	-	0	0	0
LINE	L1	L1Md_F2	chr7	21267808	21273041	+	0	0	0
LINE	L1	L1Md_F2	chr7	21467477	21472711	+	0	0	0
LINE	L1	L1Md_F2	chr7	21672872	21678098	+	0	0	0
LINE	L1	L1Md_F2	chr7	22310342	22315572	+	0	0	0
LINE	L1	L1Md_F2	chr7	22793149	22798379	+	0	0	0
LINE	L1	L1Md_F2	chr7	22990015	22995258	+	0	0	0
LINE	L1	L1Md_F2	chr7	23456574	23461810	+	0	0	0
LTR	MaLR	MTA_Mm	chr7	24557633	24558027	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	25680898	25680918	+	0	0	0
SINE	Alu	B1_Mm	chr7	26158697	26158843	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	26158844	26158877	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	26402309	26402342	+	0	0	0
SINE	Alu	B1_Mus2	chr7	26462393	26462539	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	26462540	26462583	+	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr11	75524658	75524687	+	0	0	0
SINE	Alu	B1_Mus2	chr11	75530419	75530547	-	0	0	0
SINE	Alu	B1_Mus2	chr7	27935981	27936138	+	0	0	0
SINE	Alu	B1_Mus2	chr7	28372272	28372409	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	28372410	28372432	+	0	0	0
SINE	Alu	B1_Mm	chr7	28434546	28434692	-	0	0	0
SINE	Alu	B1_Mus1	chr7	28512030	28512174	+	0	0	0
SINE	Alu	B1_Mus2	chr7	28590471	28590617	-	0	0	0
SINE	Alu	B1_Mus2	chr7	28670184	28670330	-	0	0	0
SINE	Alu	B1_Mus1	chr7	28802318	28802463	-	0	0	0
LTR	MaLR	MTB_Mm	chr7	29060036	29060441	+	0.591053	0.708814	0
SINE	Alu	B1_Mm	chr7	29618160	29618307	-	0	0	0
SINE	Alu	B1_Mus1	chr7	29653004	29653150	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	29653151	29653199	+	0	0	0
SINE	Alu	B1_Mm	chr7	29665026	29665171	+	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr7	29665172	29665197	+	0	0	0.342623
SINE	Alu	B1_Mus1	chr7	30546327	30546473	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	30752202	30752227	+	0	0	0

SINE	Alu	B1_Mus2	chr7	31170101 31170247 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	31170248 31170274 +	0	0	0
SINE	Alu	B1_Mus2	chr7	31594575 31594720 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	31594721 31594748 +	0	0	0
LTR	MaLR	MTA_Mm	chr7	31780211 31780600 -	0	0	0
LTR	MaLR	MTA_Mm	chr7	32771122 32771514 +	0	0	0
LINE	L1	L1Md_A	chr7	32908219 32914554 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	33062185 33062237 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	33089312 33089337 +	0	0	0
LINE	L1	L1_Mus1	chr7	33380908 33387196 -	0	0	0
LTR	MaLR	MTA_Mm	chr7	33503462 33503854 +	0	0	0
LTR	MaLR	MTA_Mm	chr7	33730357 33730749 +	0	0	0
LINE	L1	L1Md_F2	chr7	34321163 34326357 +	0	0	0
LINE	L1	L1Md_T	chr7	34734480 34741392 -	0	0	0
SINE	Alu	B1_Mus2	chr7	34924902 34925048 +	0	0	0
SINE	Alu	B1_Mus1	chr7	35413993 35414138 -	0	0	0
SINE	Alu	B1_Mus1	chr7	35446467 35446621 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	35446622 35446665 +	0	0	0
SINE	Alu	B1_Mm	chr7	35642870 35643016 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	35695657 35695691 +	0	0	0
SINE	Alu	B1_Mm	chr7	36015181 36015327 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	36015328 36015393 +	0	0	0
SINE	B2	B2_Mm1a	chr7	36111336 36111528 -	0	0	0
SINE	Alu	B1_Mus1	chr7	36113264 36113410 -	0	0	0
SINE	Alu	B1_Mus2	chr7	36554612 36554758 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	36554759 36554782 +	0	0	0
scRNA	scRNA	BC1_Mm	chr7	37472719 37472839 +	0	0	0
SINE	Alu	B1_Mus1	chr7	37630675 37630807 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	37630808 37630831 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	37926735 37926773 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	38109583 38109610 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	38651313 38651348 +	0	0	0
SINE	Alu	B1_Mm	chr7	39038123 39038269 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	39038270 39038295 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	47353807 47353845 +	0	0	0
SINE	Alu	B1_Mus2	chr7	47674143 47674288 +	0	0	0
Low_complexity	Low_complexity	GA-rich	chr7	47674289 47674371 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	47790412 47790449 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	48008895 48008933 +	0	0	0
SINE	B2	B3A	chr7	48040351 48040540 +	0	0	0
SINE	B2	B2_Mm1a	chr7	48345754 48345957 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	48603883 48603917 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	48609966 48610013 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	48809135 48809157 +	0	0	0
SINE	Alu	B1_Mm	chr7	48832503 48832649 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	48832650 48832683 +	0	0	0
LTR	ERVK	RLTR18B	chr7	48877616 48878136 -	0	0	0
LTR	MaLR	MTA_Mm	chr7	48914761 48915138 -	0	0	0
LTR	MaLR	MTA_Mm	chr7	48917451 48917823 -	0	0	0
SINE	Alu	B1_Mm	chr11	76746128 76746274 -	0	0	0
LINE	L1	L1Md_F2	chr7	49483419 49488637 +	0	0	0
SINE	Alu	B1_Mus2	chr7	49733650 49733789 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	49733790 49733814 +	0	0	0
SINE	Alu	B1_Mm	chr7	49845941 49846086 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	49846087 49846155 +	0	0	0
SINE	Alu	B1_Mus2	chr7	50266515 50266663 +	0.591053	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr7	50485898 50485934 +	0	0	0.342623
SINE	Alu	B1_Mm	chr7	50517710 50517855 +	0	0	0
SINE	Alu	B1_Mm	chr7	50530063 50530176 -	0	0	0
SINE	Alu	B1_Mm	chr7	50531852 50531997 -	0	0	0
SINE	Alu	B1_Mm	chr7	50575459 50575604 -	0	0	0
SINE	B2	B2_Mm1t	chr7	50811580 50811772 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	50811773 50811794 +	0	0.708814	0
SINE	B2	B2_Mm1a	chr7	50811795 50811987 +	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr7	50811988 50812024 +	0	0.708814	0
SINE	Alu	B1_Mus1	chr7	50953689 50953811 +	0	0	0
SINE	Alu	B1_Mus2	chr7	51550181 51550326 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr7	51550327 51550354 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr7	51737373 51737519 -	0.591053	0.708814	0.615324
SINE	Alu	B1_Mm	chr7	51935182 51935328 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	52361560 52361590 +	0	0	0
SINE	Alu	B1_Mm	chr7	52365118 52365243 +	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chr7	52365245 52365336 +	0	0	0
SINE	Alu	B1_Mm	chr7	52629898 52630034 -	0.923247	0.708814	0.615324
SINE	Alu	B1_Mus2	chr7	52656209 52656354 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	52867463 52867504 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr7	53019133 53019277 +	0	0	0
SINE	Alu	B1_Mus2	chr7	53037139 53037287 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	53037288 53037321 +	0	0	0
SINE	Alu	B1_Mus2	chr7	54084984 54085130 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	54085131 54085157 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	54136462 54136498 +	0	0	0
SINE	Alu	B1_Mus1	chr7	55311486 55311629 -	0	0	0
LINE	L1	L1Md_T	chr7	55553241 55560561 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	55885556 55885591 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	55972511 55972538 +	0	0	0
SINE	Alu	B1_Mm	chr7	56159738 56159884 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	56159885 56159947 +	0	0	0
SINE	Alu	B1_Mus2	chr7	56915897 56916026 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	56916027 56916077 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	57229915 57229942 +	0.923247	0	0
LTR	MaLR	MTA_Mm	chr7	58256476 58256871 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	58262586 58262622 +	0	0	0
LTR	MaLR	MTA_Mm	chr7	58350268 58350654 -	0	0	0
SINE	Alu	B1_Mus2	chr7	58415080 58415226 -	0	0	0
SINE	Alu	B1_Mus1	chr11	77435384 77435526 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	77435527 77435582 +	0	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr7	58931904 58931937 +	0	0	0
SINE	Alu	B1_Mus1	chr11	77461710 77461847 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	77461848 77461884 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	59339564 59339606 +	0	0	0
SINE	Alu	B1_Mm	chr11	77483870 77484015 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	60213788 60213814 +	0	0	0
SINE	Alu	B1_Mus2	chr7	60379635 60379779 -	0	0	0
LINE	L1	L1Md_F3	chr7	60463731 60469337 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	61562308 61562356 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	62786236 62786277 +	0	0	0
SINE	Alu	B1_Mus2	chr11	77692794 77692941 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	77692942 77692967 +	0	0	0
LINE	L1	L1Md_T	chr7	65490466 65497395 -	0	0	0
SINE	Alu	B1_Mus2	chr7	65537904 65538050 -	0	0	0
SINE	Alu	B1_Mus2	chr7	65722804 65722950 +	0	0	0
LINE	L1	L1Md_F2	chr7	67059151 67063852 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	67308826 67308859 +	0	0	0
LTR	MaLR	MTA_Mm	chr7	67328544 67328934 -	0	0	0.615324
SINE	Alu	B1_Mus2	chr7	68186190 68186336 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	68186337 68186426 +	0	0	0
LTR	MaLR	MTA_Mm	chr7	68290388 68290783 +	0	0	0
SINE	Alu	B1_Mm	chr7	68370988 68371116 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	68371117 68371157 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	69384160 69384204 +	0	0	0
SINE	Alu	B1_Mus2	chr7	69801203 69801349 +	0	0	0
LTR	MaLR	MTA_Mm	chr7	70403602 70403996 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	70587937 70587977 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	71567907 71567949 +	0	0	0
SINE	Alu	B1_Mus1	chr7	72553500 72553646 -	0.591053	0	0
SINE	Alu	B1_Mus2	chr7	72650239 72650384 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	72650385 72650419 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	73712584 73712628 +	0.591053	0	0
SINE	Alu	B1_Mus1	chr7	75967124 75967272 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	76132139 76132183 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	76563883 76563910 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	77274209 77274251 +	0	0	0
SINE	Alu	B1_Mm	chr11	78530349 78530495 -	0	0	0

SINE	Alu	B1_Mus2	chr11	78533606 78533752 +	0	0	0
SINE	Alu	B1_Mus1	chr11	78569518 78569664 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	78569665 78569692 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	78289934 78289985 +	0	0	0
LTR	MaLR	MTB_Mm	chr11	78590213 78590540 +	0	0	0
SINE	Alu	B1_Mm	chr7	78714528 78714674 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	78714675 78714704 +	0	0	0
LTR	MaLR	MTA_Mm	chr7	79699433 79699827 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	79947443 79947482 +	0	0	0
LTR	MaLR	MTA_Mm	chr7	79961838 79962218 -	0	0	0
LTR	MaLR	MTA_Mm	chr7	79963318 79963712 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	80567428 80567463 +	4.2269	0.380568	0.685246
SINE	Alu	B1_Mus1	chr7	80764656 80764802 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	80764803 80764843 +	0	0	0
SINE	Alu	B1_Mus1	chr7	82947411 82947556 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	82949904 82949940 +	0	0	0
LTR	MaLR	MTA_Mm	chr7	82977112 82977507 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	83034405 83034440 +	0	0	0
LTR	MaLR	MTB	chr7	83311886 83312283 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	83358255 83358282 +	0	0	0
SINE	Alu	B1_Mus1	chr7	84422635 84422772 -	0	0	0
SINE	Alu	B1_Mus2	chr7	84435361 84435507 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	84435508 84435567 +	0	0	0
LINE	L1	L1Md_F	chr7	84500001 84505115 -	0	0	0
LINE	L1	L1Md_F2	chr7	84601290 84607812 -	0	0	0
LTR	MaLR	MTA_Mm	chr7	84956283 84956677 +	0	0	0
SINE	Alu	B1_Mus2	chr7	85222544 85222690 -	0	0	0
SINE	Alu	B1_Mus2	chr7	86322761 86322899 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	86322900 86322931 +	0	0	0
SINE	ID	ID4_	chr7	86765187 86765272 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	86878177 86878208 +	0	0	0
SINE	Alu	B1_Mus2	chr7	86997322 86997466 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	86997467 86997502 +	0	0	0
SINE	Alu	B1_Mm	chr7	87126379 87126528 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	87127533 87127554 +	0	0	0
SINE	Alu	B1_Mm	chr7	87163954 87164100 -	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	87958328 87958364 +	0.332194	0	0
SINE	Alu	B1_Mus1	chr7	87990219 87990357 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	87990363 87990384 +	0	0	0
SINE	Alu	PB1D9	chr7	88193853 88193969 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	88193970 88194001 +	0	0	0
SINE	Alu	B1_Mus1	chr7	88251184 88251321 -	0	0	0
SINE	Alu	B1_Mus1	chr7	89155465 89155627 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	89155628 89155652 +	0	0	0
LTR	MaLR	MTA_Mm	chr7	90535309 90535703 +	0	0	0
SINE	Alu	B1_Mm	chr7	91039299 91039444 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	91039445 91039472 +	0.664388	0	0
SINE	B2	B2_Mm2	chr11	79529388 79529574 -	0	0	0
SINE	Alu	B1_Mus2	chr11	79535240 79535392 +	0.827951	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	79535393 79535416 +	0.827951	0	0
LINE	L1	L1Md_F2	chr7	91640058 91646344 -	0	0	0
SINE	Alu	B1_Mus2	chr7	91948197 91948344 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	91948345 91948374 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr7	91950339 91950372 +	0	0	0
LINE	L1	L1_Mus1	chr7	92472529 92478269 -	0	0	0
LINE	L1	L1_Mm	chr7	92504588 92509196 +	0	0	0
LTR	ERVK	MMERVK10C-int	chr7	92990535 92997858 -	0	0	0
LINE	L1	L1Md_T	chr7	93645972 93652411 -	0	0	0
SINE	Alu	B1_Mus1	chr7	94349957 94350104 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	94350105 94350128 +	0	0	0
SINE	Alu	B1_Mus2	chr7	94668643 94668789 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	94668790 94668831 +	0.332194	0.380568	0.342623
SINE	Alu	B1_Mus1	chr7	95801776 95801923 -	0	0	0
SINE	Alu	B1_Mm	chr7	95860639 95860784 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	95860785 95860826 +	0	0	0
SINE	Alu	B1_Mus2	chr11	79791562 79791708 -	0	0	0
SINE	Alu	B1_Mus1	chr7	96011138 96011285 -	0	0.708814	0

Simple_repeat	Simple_repeat	(A)n	chr7	96120264 96120305 +	0	0	0
SINE	Alu	B1_Mus1	chr11	79800821 79800967 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	96488438 96488481 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	96589926 96589973 +	0	0	0
SINE	Alu	B1_Mus1	chr7	97097571 97097707 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	97097708 97097731 +	0	0	0
SINE	B2	B2_Mm2	chr7	97370204 97370390 -	0	0	0
SINE	Alu	B1_Mm	chr7	97664080 97664231 +	0	0	0
LTR	MaLR	MTA_Mm	chr7	97861306 97861693 +	0	0	0
LTR	MaLR	MTB_Mm	chr7	99682320 99682699 -	0	0	0
SINE	B2	B2_Mm1a	chr11	79961662 79961855 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	79961856 79961877 +	0	0	0
SINE	Alu	B1_Mm	chr7	99822363 99822508 +	0	0	0
LTR	MaLR	MTA_Mm	chr7	10031210 10031249 +	0	0.380568	0
SINE	Alu	B1_Mus2	chr11	80014448 80014593 -	0	0	0
LTR	MaLR	MTA_Mm	chr7	10083790 10083829 +	0	0	0
SINE	Alu	B1_Mm	chr7	10129095 10129110 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	10132065 10132070 +	0.591053	0	0
SINE	Alu	B1_Mus2	chr7	10138948 10138962 -	0	0	0
SINE	Alu	B1_Mm	chr7	10145612 10145627 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	10145627 10145629 +	0	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr7	10184477 10184480 +	0.332194	0	0
LTR	MaLR	MTA_Mm	chr7	10212013 10212053 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	10247196 10247200 +	0.591053	0	0
SINE	Alu	B1_Mus1	chr7	10262413 10262427 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	10262428 10262430 +	0	0	0
SINE	Alu	B1_Mm	chr7	10294261 10294275 -	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	10298374 10298379 +	0	0	0
SINE	B4	ID_B1	chr10	25445135 25445348 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	10322820 10322823 +	0	0	0
SINE	Alu	B1_Mus2	chr11	80231157 80231302 +	3.31962	0	1.57327
Simple_repeat	Simple_repeat	(A)n	chr11	80231303 80231328 +	2.39637	0	0.957947
SINE	Alu	B1_Mus1	chr7	10491230 10491244 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	10491245 10491250 +	0	0	0
SINE	Alu	B1_Mus1	chr7	10494614 10494626 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr7	10494626 10494628 +	0	0	0.615324
LTR	MaLR	MTA_Mm	chr7	10535795 10535835 +	0	0	0
LTR	MaLR	MTA_Mm	chr7	10535945 10535984 +	0	0	0
SINE	Alu	B1_Mus1	chr7	10552352 10552366 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	10552366 10552368 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	10581386 10581389 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	10584102 10584106 +	0	0	0
SINE	Alu	B1_Mm	chr7	10664218 10664233 -	0	0	0
SINE	Alu	B1_Mus1	chr7	10664413 10664428 -	0	0	0
SINE	Alu	B1_Mm	chr7	10664708 10664722 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	10664722 10664725 +	0	0	0
scRNA	scRNA	4.5SRNA	chr7	10665232 10665242 +	0	0	0
SINE	Alu	B1_Mm	chr7	10665242 10665256 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	10665457 10665461 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	10768237 10768240 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr7	10783401 10783405 +	0	0	0
SINE	Alu	B1_Mus2	chr7	10785256 10785271 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	10785271 10785273 +	0	0	0
SINE	Alu	B1_Mus1	chr7	10801448 10801463 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	10821144 10821147 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	10903340 10903344 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	10999554 10999561 +	0	0	0
SINE	Alu	B1_Mus1	chr7	11037810 11037822 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	11075697 11075702 +	0	0	0
LTR	MaLR	MTA_Mm	chr7	11109880 11109919 +	0	0	0
SINE	Alu	B1_Mus2	chr7	11189768 11189782 -	0	0	0
SINE	Alu	B1_Mus2	chr7	11193028 11193042 +	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chr7	11193044 11193053 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	11238219 11238223 +	0	0	0
LINE	L1	L1_Mus2	chr7	11319969 11320565 +	0	0	0
LINE	L1	L1_Mus2	chr7	11344155 11344752 +	0	0	0
LINE	L1	L1Md_F2	chr7	11350100 11350615 +	0	0	0

SINE	Alu	B1_Mus1	chr7	11383414 11383429 -	0	0	0
SINE	Alu	B1_Mm	chr7	11389750 11389764 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	11389764 11389771 +	0.332194	0	0
SINE	Alu	B1_Mm	chr7	11393291 11393306 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	11393306 11393308 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	11545426 11545431 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	11636434 11636438 +	0	0	0
SINE	Alu	PB1D10	chr7	11677144 11677155 +	0	0	0
SINE	B4	RSINE1	chr7	11677156 11677171 +	0	0	0
SINE	Alu	B1_Mus2	chr7	11679517 11679531 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr7	11977436 11977440 +	0.591053	0	0
SINE	Alu	B1_Mus2	chr7	11981833 11981847 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	11981847 11981850 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	12006431 12006434 +	0	0	0
SINE	Alu	B1_Mm	chr7	12045890 12045905 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	12045905 12045907 +	0	0	0
SINE	B2	B3A	chr7	12126971 12126988 +	0	0	0
SINE	Alu	B1_Mm	chr7	12213935 12213949 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	12255675 12255678 +	0	0	0
SINE	B2	B2_Mm1t	chr7	12336216 12336235 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	12336235 12336238 +	0.332194	0	0.685246
Simple_repeat	Simple_repeat	(A)n	chr7	12375033 12375035 +	0	0	0
SINE	Alu	B1_Mm	chr7	12410065 12410079 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	12410079 12410084 +	0	0	0
SINE	Alu	B1_Mm	chr7	12421433 12421448 -	0	0	0
LTR	MaLR	MTA_Mm	chr11	82282149 82282543 -	0	0	0
LTR	MaLR	MTE2b	chr7	12493021 12493055 +	0	0	0
SINE	Alu	B1_Mm	chr7	12517368 12517383 +	0	0	0
LTR	MaLR	MTD	chr7	12517383 12517419 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	12542299 12542301 +	0	0	0
SINE	Alu	B1_Mm	chr11	82515748 82515902 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	82515903 82515933 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	12707879 12707883 +	0	0	0
SINE	Alu	B1_Mm	chr7	12712337 12712351 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	12712351 12712354 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	12726911 12726915 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	12769478 12769481 +	0	0	0
LTR	MaLR	MTA_Mm	chr7	12867178 12867218 -	0	0	0
SINE	Alu	B1_Mus1	chr7	12896657 12896672 +	0	0	0
SINE	Alu	B1_Mus2	chr7	12897278 12897293 -	0	0	0
LTR	MaLR	MTB	chr11	82858321 82858724 -	0	0	0
SINE	Alu	B1_Mus2	chr7	12992084 12992098 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	12992098 12992104 +	0.332194	0	0
SINE	Alu	B1_Mus1	chr7	13019661 13019675 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	13019677 13019683 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	13254096 13254098 +	0	0	0
SINE	Alu	B1_Mm	chr7	13347232 13347246 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	13347246 13347251 +	0.332194	0	0
SINE	Alu	B1_Mus1	chr7	13349114 13349129 -	0	0	0
SINE	Alu	B1_Mm	chr7	13354631 13354646 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	13354963 13354966 +	0	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr7	13368111 13368114 +	0	0	0
SINE	B2	B2_Mm1a	chr7	13377386 13377405 +	0	0	0
SINE	Alu	B1_Mus1	chr7	13405601 13405616 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	13405616 13405620 +	0	0	0
SINE	Alu	PB1D9	chr7	13420134 13420146 -	0	0	0
SINE	Alu	B1_Mus1	chr7	13421143 13421157 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	13428903 13428908 +	0	0	0
LTR	MaLR	MTA_Mm	chr7	13451271 13451311 +	0	0	0
LTR	MaLR	MTA_Mm	chr7	13451407 13451447 +	0	0	0.342623
SINE	Alu	B1_Mm	chr7	13456061 13456075 +	0	0	0
SINE	Alu	B1_Mus1	chr7	13464077 13464089 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	13464089 13464091 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr7	13467081 13467085 +	0.591053	0.708814	0
SINE	Alu	B1_Mus2	chr7	13479306 13479322 -	0	0	1.23065
Simple_repeat	Simple_repeat	(A)n	chr7	13487171 13487176 +	0.332194	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr7	13488398 13488400 +	0	0	0

SINE	B2	B3A	chr7	13513258 13513277 -	0	0	0
SINE	Alu	B1_Mus1	chr7	13546532 13546547 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	13546547 13546549 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	13548458 13548461 +	0	0	1.23065
SINE	Alu	B1_Mus2	chr7	13563330 13563345 +	0	0	0.342623
SINE	Alu	B1_Mus1	chr7	13570244 13570258 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	25897460 25897506 +	0	0	0
SINE	B2	B2_Mm1a	chr7	13753343 13753364 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	13753364 13753370 +	0.664388	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	13865643 13865646 +	0	0	0
SINE	Alu	B1_Mm	chr11	83891508 83891654 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	83891655 83891700 +	0	0	0
SINE	Alu	B1_Mus1	chr7	14013711 14013726 +	4.74462	2.50701	3.29367
Simple_repeat	Simple_repeat	(A)n	chr7	14013726 14013729 +	1.91983	2.50701	3.02097
SINE	Alu	B1_Mm	chr11	83961408 83961553 +	0	0	0.957947
Simple_repeat	Simple_repeat	(A)n	chr11	83961554 83961588 +	0	0	0.957947
SINE	Alu	B1_Mur2	chr11	83984105 83984248 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	14178322 14178325 +	0	0	0
SINE	Alu	B1_Mus2	chr7	14185174 14185189 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr7	14297447 14297456 +	0	0	0
SINE	Alu	B1_Mur1	chr7	14345894 14345909 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	14345909 14345913 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	14359849 14359853 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	14399906 14399910 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr7	14518143 14518158 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	14518158 14518162 +	0	0	0
SINE	Alu	B1_Mm	chr7	14610962 14610976 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	14683406 14683409 +	0	0	0
LTR	MaLR	MTA_Mm	chr7	14695618 14695658 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	14707763 14707766 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	14727285 14727289 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	14798175 14798178 +	0	0	0
SINE	Alu	B1_Mus1	chr7	14819275 14819288 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	14819288 14819292 +	0	0	0
SINE	Alu	B1_Mus1	chr7	14836667 14836681 -	0	0	0
SINE	B2	B2_Mm1t	chr7	14884514 14884533 -	0	0	0.342623
SINE	Alu	B1_Mus1	chr7	14979876 14979890 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	14979890 14979894 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	14985665 14985673 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr7	15069511 15069513 +	0	0	0
SINE	Alu	B1_Mm	chr7	15078810 15078824 +	0	0	0
SINE	Alu	B1_Mm	chr7	15161772 15161786 -	0	0	0
SINE	Alu	B1_Mm	chr11	84689127 84689260 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	84689261 84689308 +	0	0	0
Low_complexity	Low_complexity	GA-rich	chr7	15201787 15201799 +	0	0	0
SINE	Alu	B1_Mus2	chr11	84769140 84769286 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	84769287 84769309 +	0	0.380568	0
SINE	Alu	B1_Mus1	chr8	3560946 3561092 -	0	0	0
LTR	MaLR	MTA_Mm	chr8	3861211 3861583 +	0	0	0
SINE	Alu	B1_Mm	chr8	4133219 4133360 -	0	0	0
LTR	MaLR	MTA_Mm	chr8	4422710 4423104 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	5119157 5119187 +	0	0	0.342623
SINE	Alu	B1_Mus1	chr8	5637088 5637232 +	0	0	0
Low_complexity	Low_complexity	GA-rich	chr8	5637233 5637368 +	0	0	0
SINE	Alu	B1_Mm	chr8	5885533 5885679 -	0	0	0
SINE	Alu	B1_Mus2	chr8	6169310 6169455 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	6169456 6169475 +	0	0	0
LTR	MaLR	MTA_Mm	chr8	6259817 6260202 -	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr8	6297219 6297259 +	0	0	0
SINE	Alu	B1_Mus1	chr8	6436951 6437104 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	6460216 6460253 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	6644418 6644457 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	6857748 6857768 +	0	0	0
LTR	MaLR	MTB	chr8	7001471 7001867 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	7135552 7135577 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	7492555 7492591 +	0	0	0
SINE	Alu	B1_Mus1	chr11	85055835 85055982 -	0	0	0



Simple_repeat	Simple_repeat	(A)n	chr8	7914883	7914945	+	0	0	0
LINE	L1	L1Md_F2	chr8	8066791	8073201	-	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr8	8314801	8314850	+	0	0	0
SINE	Alu	B1_Mus2	chr8	8614893	8615039	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	8615040	8615076	+	0	0	0
SINE	Alu	B1_Mus2	chr8	8701644	8701790	-	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr8	8723275	8723297	+	0	0	0
LTR	MaLR	MTA_Mm	chr8	8816865	8817255	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	8826065	8826112	+	0	0	0
SINE	Alu	B1_Mus2	chr11	85115720	85115860	-	0	0	0
SINE	Alu	B1_Mus2	chr8	9778463	9778609	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	9778610	9778636	+	0	0	0
SINE	Alu	B1_Mm	chr11	85161428	85161569	-	0	0	0
LTR	MaLR	MTB	chr8	10799864	10800255	-	0	0	0
SINE	Alu	B1_Mus2	chr8	10969701	10969847	-	0	0	0
SINE	Alu	B1_Mm	chr8	11495289	11495433	-	0	0	0
SINE	Alu	B1_Mus2	chr8	12266863	12266998	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	12495752	12495797	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	12673499	12673532	+	0	0	0.615324
SINE	Alu	B1_Mus1	chr8	13690256	13690401	+	0	0	0
Low_complexity	Low_complexity	A-rich	chr8	13690402	13690472	+	0	0	0
SINE	Alu	B1_Mus2	chr8	13855473	13855619	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	13855620	13855655	+	0	0	0
LTR	MaLR	MTA_Mm	chr8	15176455	15176846	+	0	0.708814	0
LTR	MaLR	MTA_Mm	chr8	15198515	15198910	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	15449660	15449687	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	15557450	15557479	+	0	0	0
SINE	ID	ID2	chr8	17560192	17560276	-	0	0	0
LTR	MaLR	MTA_Mm	chr8	17630582	17630961	+	0	0	0
SINE	Alu	B1_Mus2	chr8	18138356	18138493	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	18326938	18326974	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	18894133	18894175	+	0	0	0
LTR	MaLR	MTA_Mm	chr8	19539381	19539776	+	0	0	0.342623
SINE	Alu	B1_Mus1	chr8	22942384	22942530	-	0	0	0
SINE	Alu	B1_Mm	chr8	23025264	23025409	-	0	0	0
SINE	Alu	B1_Mus2	chr8	23301528	23301674	+	0	0	0
LTR	MaLR	MTA_Mm	chr8	25438769	25439166	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	25493432	25493472	+	0	0	0
LTR	MaLR	MTA_Mm	chr8	25518738	25519129	+	0	0	0
SINE	Alu	B1_Mur4	chr8	25556609	25556730	-	0	0	0
LTR	MaLR	MTB	chr8	25611755	25612148	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	26574112	26574140	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	26581100	26581128	+	0	0	0
LTR	MaLR	MTA_Mm	chr8	27053880	27054275	+	0	0	0
SINE	Alu	B1_Mus1	chr11	86256819	86256954	-	0	0	0
SINE	Alu	B1_Mus2	chr8	27735629	27735779	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	27735780	27735811	+	0	0	0
LTR	MaLR	MTA_Mm	chr8	27898119	27898511	-	0	0	0
Low_complexity	Low_complexity	AT-rich	chr8	28320437	28320468	+	0	0	0
LTR	MaLR	MTA_Mm	chr8	28355228	28355622	+	0	0	0
SINE	Alu	B1_Mus2	chr8	28636372	28636521	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	29265964	29266013	+	0	0	0
LTR	MaLR	MTA_Mm	chr8	29659436	29659815	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	30607396	30607432	+	0	0	0
SINE	Alu	B1_Mus1	chr8	30981921	30982041	-	0	0	0
LTR	MaLR	MTA_Mm	chr8	30988457	30988850	+	0	0	0
SINE	Alu	B1_Mm	chr8	31064433	31064579	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	31522711	31522739	+	0	0	0
LTR	MaLR	MTA_Mm	chr8	31885947	31886327	+	0.591053	0	0
LTR	MaLR	MTA_Mm	chr8	32059179	32059548	+	0	0	0
LTR	MaLR	MTA_Mm	chr8	32095674	32096065	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	33115317	33115358	+	0	0	0
LTR	MaLR	MTA_Mm	chr8	33543311	33543698	+	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	34090225	34090256	+	0	0	0
SINE	Alu	B1_Mus1	chr8	34268624	34268770	+	0	0	0
SINE	Alu	B1_Mus1	chr8	34332837	34332964	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	34332965	34332985	+	0	0	0

SINE	Alu	B1_Mus2	chr8	34560826 34560972 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	34560973 34561017 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	35043079 35043115 +	0	0	0
SINE	Alu	B1_Mus2	chr8	35109423 35109569 -	0	0	0
SINE	Alu	B1_Mus1	chr8	35214861 35215010 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	35422964 35422993 +	0	0	0
SINE	Alu	B1_Mus2	chr8	36130404 36130552 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	36271709 36271760 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	37103832 37103873 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	39751601 39751633 +	0	0	0
SINE	Alu	B1_Mm	chr8	39872774 39872920 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	39872921 39872944 +	0	0	0
LTR	MaLR	MTA_Mm	chr8	40476505 40476894 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	41867776 41867810 +	0	0	0
LTR	MaLR	MTA_Mm	chr8	42271490 42271877 -	0	0	0
SINE	Alu	B1_Mus1	chr8	42543775 42543923 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr8	42543935 42543967 +	0	0	0
LINE	L1	L1Md_F2	chr8	42587006 42592293 -	0	0	0
LTR	MaLR	MTA_Mm	chr8	43716956 43717349 -	0	0	0
LTR	MaLR	MTA_Mm	chr8	43718429 43718822 -	0	0	0
SINE	Alu	B1_Mm	chr8	44287862 44288006 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	44288007 44288032 +	0	0	0
SINE	Alu	B1_Mm	chr8	44426680 44426826 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	44426827 44426854 +	0	0	0
SINE	Alu	B1_Mus1	chr8	44823905 44824052 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	44824053 44824078 +	0	0	0
SINE	Alu	B1_Mus2	chr8	44876437 44876583 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	44876584 44876717 +	0	0	0
LTR	MaLR	MTA_Mm	chr8	45089812 45090186 +	0	0	0
LINE	L1	L1Md_F2	chr8	45379230 45385341 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	45536722 45536766 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	46549539 46549565 +	0	0	0
SINE	Alu	B1_Mm	chr8	48251306 48251452 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	48251453 48251483 +	0	0	0
SINE	Alu	B1_Mus1	chr8	48543668 48543814 -	0	0	0
SINE	Alu	B1_Mus1	chr8	48648542 48648688 -	0	0	0
SINE	Alu	B1_Mus1	chr8	48726136 48726253 +	0	0	0
SINE	Alu	B1_Mus2	chr8	48783816 48783960 -	0	0	0
SINE	Alu	B1_Mus1	chr8	49100980 49101115 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	49101116 49101138 +	0	0.380568	0
LTR	MaLR	MTB	chr8	49241649 49242047 -	0	0.708814	0.615324
Simple_repeat	Simple_repeat	(A)n	chr8	49796393 49796431 +	0	0	0
LTR	ERVK	RMER6D	chr8	50515401 50516187 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	50926890 50926918 +	0	0	0
LTR	MaLR	MTA_Mm	chr8	51359765 51360156 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	51412196 51412222 +	0	0	0
LTR	MaLR	MTA_Mm	chr8	51443621 51444013 +	0	0	0
SINE	Alu	B1_Mm	chr8	51512988 51513133 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	51513134 51513166 +	0	0	0
SINE	Alu	B1_Mus2	chr8	51824363 51824509 -	1.18211	0	0.615324
LINE	L1	L1Md_F2	chr8	52060475 52065552 +	0	0	1.10507
LTR	MaLR	ORR1D1-int	chr8	52429191 52430442 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	53006005 53006047 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	53046760 53046803 +	0	0	0
SINE	Alu	B1_Mus1	chr8	53464191 53464337 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	53900981 53901037 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	54506778 54506817 +	0	0.380568	0.957947
Simple_repeat	Simple_repeat	(GAAA)n	chr8	54520446 54520619 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr8	54563175 54563287 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	55016041 55016077 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	55091961 55091988 +	0	0	0
SINE	Alu	B1_Mm	chr8	55277062 55277207 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	58922436 58922482 +	0	0	0
SINE	Alu	B1_Mus1	chr8	59104641 59104785 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	59104786 59104809 +	0	0	0
SINE	Alu	B1_Mus2	chr11	88055784 88055917 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	88055918 88055941 +	0	0	0

LTR	MaLR	MTA_Mm	chr8	59524362 59524757 -	0	0	0.342623
LTR	MaLR	MTA_Mm	chr8	59756679 59757060 +	0	0	0
SINE	Alu	B1_Mus2	chr8	59882899 59883024 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	59883025 59883074 +	0	0	0
SINE	Alu	B1_Mur2	chr8	59926214 59926359 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr8	59926362 59926426 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	61420247 61420328 +	0.591053	0	0
SINE	B4	ID_B1	chr8	61585647 61585860 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	61585861 61585885 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	26491840 26491882 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	62066868 62066902 +	0	0	0
LINE	L1	L1Md_F2	chr8	62339122 62344820 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	63172433 63172474 +	0	0	0
LTR	MaLR	MTA_Mm	chr8	63186018 63186413 -	0	0	0
SINE	Alu	B1_Mus1	chr8	63288522 63288668 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	63288669 63288707 +	0	0	0
SINE	Alu	B1_Mm	chr8	63359483 63359628 -	0	0	0
SINE	Alu	B1_Mm	chr8	63683984 63684115 +	0	0	0
Simple_repeat	Simple_repeat	(GAAAA)n	chr8	63684116 63684183 +	0	0	0
LTR	MaLR	MTB	chr8	63934211 63934606 -	0	0	0
LINE	L1	L1Md_F2	chr8	64649394 64655284 -	0	0	0.342623
SINE	Alu	B1_Mus2	chr8	64899392 64899540 -	0	0	0
SINE	Alu	B1_Mm	chr11	88630165 88630311 -	0	0	0
SINE	Alu	B1_Mus1	chr8	65331953 65332099 +	0	0	0
LTR	MaLR	MTA_Mm	chr8	65409137 65409532 +	0	0	0
LTR	MaLR	MTA_Mm	chr8	65943005 65943398 +	0	0	0
LINE	L1	L1_Mus3	chr8	66079146 66085069 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	66093597 66093623 +	0	0	0
SINE	Alu	B1_Mm	chr11	88739080 88739225 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr11	88739226 88739256 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr8	67423729 67423876 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	67423877 67423933 +	0	0	0
LINE	L1	L1Md_F2	chr8	67755269 67761636 -	0	0	0.615324
LINE	L1	L1Md_F2	chr8	69045686 69050636 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	69131143 69131177 +	0	0	0
SINE	B2	B2_Mm1t	chr8	69237567 69237757 +	0	0	0
SINE	Alu	B1_Mus2	chr8	69586444 69586590 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	69586591 69586615 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	69889516 69889537 +	0	0	0
SINE	Alu	B1_Mus2	chr8	70456423 70456570 +	0	0	0
SINE	Alu	B1_Mus1	chr8	70875587 70875734 -	0.591053	0	0.342623
SINE	Alu	B1_Mus2	chr8	71387446 71387591 -	0	0	0
SINE	Alu	B1_Mus1	chr8	71553053 71553199 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	71553200 71553255 +	0	0	0
LTR	MaLR	MTA_Mm	chr8	72209932 72210329 +	0	0	0
SINE	B2	B2_Mm1t	chr8	72212760 72212944 -	0	0	0
SINE	Alu	B1_Mm	chr11	89021077 89021222 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	89021223 89021244 +	0	0	0
SINE	Alu	B1_Mus1	chr8	73391442 73391589 -	0	0	0
SINE	Alu	B1_Mus2	chr8	73405805 73405951 +	0	0	0
SINE	Alu	B1_Mm	chr8	73763998 73764144 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	73764145 73764174 +	0	0	0
SINE	B2	B2_Mm2	chr8	74017526 74017714 -	0	0	0
SINE	Alu	B1_Mus1	chr8	74056019 74056158 -	0	0	0
LTR	MaLR	MTA_Mm	chr8	74444007 74444401 -	0	0	0
SINE	Alu	B1_Mm	chr8	74628346 74628495 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	74628496 74628547 +	0	0.380568	0
SINE	Alu	B1_Mm	chr8	74710869 74711016 -	0	0	0
LINE	L1	L1Md_F2	chr8	76029152 76035614 -	0	0	0
LTR	MaLR	MTA_Mm	chr8	76367664 76368059 +	0	0	0
LINE	L1	Lx	chr8	76683556 76687980 -	0	0	0
LINE	L1	L1Md_F2	chr8	77033597 77039284 -	0.332194	0	0.342623
SINE	Alu	B1_Mm	chr8	77730886 77731031 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	77759633 77759659 +	0	0	0
LTR	MaLR	MTA_Mm	chr11	89619604 89619999 +	0	0	0
LINE	L1	L1Md_F3	chr8	78687786 78693976 -	0	0	0
LINE	L1	L1_Mus1	chr8	78759451 78765556 -	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr8	78949564 78949592 +	0	0	0
SINE	Alu	B1_Mm	chr8	80159529 80159675 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	80614288 80614323 +	0.591053	0	0
SINE	Alu	B1_Mm	chr8	80693450 80693596 +	0	0	0
SINE	Alu	B1_Mm	chr8	82002684 82002829 -	0	0	0
LINE	L1	Lx	chr8	82331942 82338062 -	0	0	0
SINE	Alu	B1_Mus2	chr8	82898210 82898347 -	0	0	0
SINE	Alu	B1_Mur4	chr8	83271368 83271515 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	83271516 83271562 +	0	0	0
SINE	Alu	B1_Mus2	chr8	83281797 83281943 -	0.591053	0.708814	0
LTR	MaLR	MTA_Mm	chr11	90164611 90165009 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	85175047 85175114 +	0	0	0
SINE	Alu	B1_Mm	chr8	86641105 86641236 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	86641237 86641263 +	0	0	0
SINE	Alu	B1_Mus2	chr8	86802241 86802389 -	0	0	0
SINE	Alu	B1_Mm	chr11	90583153 90583299 -	0	0	0
SINE	Alu	B1_Mus2	chr11	90619878 90620024 -	0	0	0
SINE	Alu	B1_Mus2	chr8	87419075 87419220 -	0	0	0
SINE	Alu	B1_Mus1	chr8	87461716 87461862 -	0	0.708814	0
LTR	MaLR	MTA_Mm	chr8	87628089 87628469 +	0	0	0
SINE	Alu	B1_Mm	chr8	87660541 87660687 -	0	0	0
LTR	ERV1	RMER21A	chr8	87744729 87745752 +	0	0	0
SINE	Alu	B1_Mus1	chr8	87979208 87979349 -	0	0	0
SINE	Alu	B1_Mm	chr8	88013735 88013873 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	88013874 88013897 +	0	0	0
LTR	MaLR	MTA_Mm	chr11	90778234 90778629 +	0	0	0
SINE	Alu	B1_Mm	chr8	90494380 90494529 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	90494530 90494551 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	90508017 90508051 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	91001680 91001712 +	0	0	0
SINE	Alu	B1_Mm	chr8	90703388 90703516 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr8	90703517 90703540 +	0	0	0.957947
SINE	Alu	B1_Mm	chr8	90720235 90720381 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	90720382 90720422 +	0	0	0
SINE	Alu	B1_Mm	chr8	90791429 90791546 +	0	0	0
SINE	Alu	PB1D7	chr8	90854945 90855058 -	2.69641	0	1.57327
SINE	Alu	PB1D10	chr8	92718421 92718508 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	92718509 92718535 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	95078434 95078473 +	0	0	0
LTR	MaLR	MTA_Mm	chr8	96257382 96257776 +	0	0	0
SINE	Alu	B1_Mus2	chr8	97221630 97221775 -	0	0	0
SINE	Alu	B1_Mm	chr8	98378650 98378796 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	98378797 98378822 +	0	0	0
SINE	Alu	B1_Mus1	chr8	98380193 98380339 -	0	0	0
LTR	MaLR	MTD	chr8	98923439 98923851 -	0	0	0
LTR	MaLR	MTA_Mm	chr8	99478308 99478703 -	0	0	0
SINE	Alu	B1_Mus2	chr8	99811309 99811454 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	99885489 99885510 +	0	0	0
LINE	L1	L1Md_T	chr8	10006921 10007584 -	0	0	0
LTR	MaLR	MTA_Mm	chr8	10118281 10118320 +	0	0	0
SINE	Alu	B1_Mus1	chr8	10121508 10121521 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	10216197 10216201 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	10299117 10299120 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	10331432 10331436 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	10342302 10342308 +	0	0	0
SINE	Alu	B1_Mm	chr8	10366987 10367002 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	10367002 10367005 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	10387771 10387775 +	0	0	0
LTR	MaLR	MTA_Mm	chr8	10434075 10434115 +	0	0	0
LINE	L1	L1Md_T	chr8	10466535 10467248 -	0	0	0
LINE	L1	L1Md_F2	chr8	10484053 10484626 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	10542351 10542356 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	10652139 10652142 +	0	0	0
SINE	Alu	B1_Mm	chr8	10715091 10715106 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	10715106 10715113 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	10762592 10762595 +	0	0	0
SINE	Alu	B1_Mus2	chr8	10895255 10895269 -	0	0	0

SINE	Alu	B1_Mus1	chr8	10910849 10910863 -	0	0	0
SINE	Alu	B1_Mm	chr8	10937630 10937644 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	10937645 10937647 +	0.923247	0	0
SINE	Alu	B1_Mm	chr8	10963553 10963567 -	0	0	0
SINE	Alu	B1_Mus2	chr8	10974110 10974125 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	10994992 10994995 +	0	0	0
SINE	B2	B2_Mm1t	chr8	11011517 11011536 +	0	0	0
SINE	Alu	PB1D10	chr8	11012684 11012695 -	0	0	0
SINE	Alu	B1_Mm	chr8	11024829 11024843 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	11024843 11024846 +	0	0	0
SINE	Alu	B1_Mm	chr8	11032623 11032636 -	0	0	0
SINE	Alu	B1_Mus2	chr8	11037079 11037094 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	11059637 11059641 +	0	0	0
SINE	Alu	B1_Mus2	chr8	11112737 11112751 -	0	0	0
SINE	Alu	B1_Mus2	chr8	11175831 11175846 -	1.77316	0	0
SINE	Alu	B1_Mur4	chr8	11181913 11181927 +	0	0	0.615324
SINE	Alu	B1_Mus2	chr8	11211317 11211332 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	11211332 11211335 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr8	11253238 11253253 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	11271872 11271874 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr8	11416345 11416348 +	0	0	0
SINE	Alu	B1_Mus2	chr8	11446222 11446237 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	11446237 11446245 +	0	0	0
SINE	Alu	B1_Mus2	chr8	11459308 11459323 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr8	11459323 11459329 +	0	0	0
SINE	Alu	B1_Mm	chr8	11485893 11485907 -	10.9481	12.0857	16.4115
Simple_repeat	Simple_repeat	(A)n	chr8	11502283 11502294 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	11503074 11503079 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	11640966 11640970 +	0	0	0
LTR	MaLR	MTA_Mm	chr8	11814801 11814840 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	11869772 11869775 +	0	0	0
SINE	Alu	B1_Mm	chr8	11877369 11877384 +	0	0	0
Simple_repeat	Simple_repeat	(GAA)n	chr8	11877384 11877392 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	11887135 11887138 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	11939972 11939976 +	0	0	0
LTR	MaLR	MTB	chr8	12041555 12041595 +	0	0	0
SINE	Alu	B1_Mus2	chr11	94255060 94255211 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	12187645 12187648 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	12240632 12240635 +	0	0	0
SINE	Alu	B1_Mus2	chr8	12243119 12243133 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	12243133 12243136 +	0	0	0
LTR	MaLR	MTA_Mm	chr8	12257773 12257812 -	0	0	0
LTR	MaLR	MTA_Mm	chr8	12259379 12259418 +	0	0	0
SINE	B2	B2_Mm1t	chr8	12276899 12276918 -	0	0	0
SINE	Alu	B1_Mus2	chr8	12341458 12341472 -	0	0	0
SINE	Alu	B1_Mus2	chr8	12395968 12395982 -	0	0	0
SINE	Alu	B1_Mm	chr8	12497589 12497603 -	0	0	0
SINE	B4	ID_B1	chr11	94543115 94543335 -	0	0	0
SINE	Alu	B1_Mm	chr8	12508733 12508747 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	12508748 12508750 +	0	0	0
SINE	Alu	B1_Mus2	chr8	12557504 12557519 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	12557519 12557522 +	0	0	0
SINE	Alu	B1_Mus2	chr11	94596615 94596762 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	94596763 94596788 +	0	0	0
LTR	MaLR	MTB	chr8	12615026 12615066 -	0	0	0
SINE	Alu	B1_Mus2	chr8	12639395 12639409 -	0	0	0
SINE	Alu	B1_Mus2	chr8	12639417 12639432 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	12639432 12639434 +	0	0	0
LTR	MaLR	MTA_Mm	chr8	12707709 12707748 +	0	0	0
SINE	Alu	B1_Mus1	chr8	12756930 12756944 -	0	0	0.615324
SINE	Alu	B1_Mm	chr11	94733525 94733665 -	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr8	12790073 12790077 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	12810565 12810569 +	0	0	0
LTR	MaLR	MTA_Mm	chr8	12842368 12842407 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	12866146 12866149 +	0	0	0
LTR	MaLR	MTA_Mm	chr8	12868505 12868545 -	0	0	0.342623
SINE	Alu	B1_Mus1	chr8	12943801 12943815 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr8	12943815	12943818 +	0	0	0
SINE	Alu	B1_Mus2	chr8	12950230	12950245 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	12950245	12950257 +	0	0	0
SINE	Alu	B1_Mus1	chr8	13029316	13029329 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	13029329	13029337 +	0	0	0
LINE	L1	L1Md_F2	chr8	13039370	13039838 +	0	0	0
LTR	MaLR	MTB	chr8	13059788	13059828 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr8	13118066	13118069 +	0	0	0
SINE	Alu	B1_Mus2	chr11	95035331	95035477 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	95035478	95035500 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	3291564	3291953 -	0	0	0
SINE	Alu	B1_Mus2	chr11	95112300	95112445 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	95112446	95112473 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	5039482	5039521 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	5127787	5127809 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	5166358	5166752 +	0	0	0
SINE	Alu	B1_Mus1	chr9	5189843	5189977 -	0	0	0
SINE	Alu	B1_Mus1	chr9	5245575	5245712 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	5245713	5245754 +	0	0	0
SINE	Alu	B1_Mus2	chr9	5491705	5491849 +	0	0	0.342623
SINE	Alu	B1_Mm	chr9	5577244	5577390 -	0	0	0
SINE	Alu	B1_Mus2	chr9	5645142	5645288 +	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chr9	5645289	5645349 +	0	0	0
SINE	Alu	B1_Mus2	chr9	5848659	5848805 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	27462752	27462806 +	0	0	0
LTR	MaLR	MTB	chr9	7255216	7255613 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	7267408	7267803 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	7268900	7269295 +	0.332194	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr9	7593443	7593483 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	7684180	7684228 +	0	0	0
SINE	Alu	B1_Mus1	chr9	7899052	7899198 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	7899199	7899257 +	0	0	0
SINE	Alu	B1_Mus1	chr9	7911626	7911743 -	0	0	0
LTR	MaLR	MTA_Mm	chr9	8508040	8508435 +	0	0	0
SINE	Alu	B1_Mm	chr9	9277938	9278084 -	0	0	0
SINE	Alu	B1_Mus2	chr11	95468586	95468734 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	95468735	95468771 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	9361024	9361419 +	0.591053	0	0
SINE	Alu	B1_Mm	chr11	95499213	95499358 -	0	0	0
LTR	MaLR	MTA_Mm	chr9	11419025	11419419 +	0	0	0
LTR	ERVK	RLTR45	chr9	11513613	11514038 -	0.332194	0	0
LINE	L1	L1Md_F3	chr9	12533593	12539976 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	12880296	12880328 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	12951421	12951462 +	0	0	0
SINE	Alu	B1_Mm	chr9	13138414	13138560 +	0	0	0
SINE	Alu	B1_Mus2	chr9	13331449	13331594 +	0.923247	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	13331595	13331618 +	0.591053	0	0
SINE	Alu	B1_Mus2	chr9	13515354	13515499 +	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr9	13515500	13515525 +	0	0.708814	0
SINE	Alu	B1_Mus1	chr9	14432424	14432571 +	0	0	0
LINE	L1	L1Md_A	chr10	27532354	27539767 -	0	0	0
SINE	Alu	B1_Mus2	chr9	14684447	14684593 +	0	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr9	14684594	14684618 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	14762485	14762517 +	0	0	0
SINE	Alu	B1_Mus1	chr11	95789649	95789796 -	0	0	0
LTR	MaLR	MTA_Mm	chr9	16786370	16786764 +	0	0	0
SINE	Alu	B1_Mus2	chr9	17779615	17779761 -	0	0	0
LINE	L1	L1_Mus1	chr9	17782484	17787747 +	0	0	0
SINE	Alu	B1_Mus2	chr9	18334960	18335083 -	0	0	0
SINE	Alu	B1_Mus2	chr9	18908476	18908621 -	0	0	0
SINE	Alu	B1_Mus1	chr9	19230144	19230250 -	0	0	0
SINE	Alu	B1_Mm	chr9	19673863	19674008 -	0	0	0
LTR	MaLR	MTA_Mm	chr9	19839545	19839927 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	20440314	20440349 +	0	0	0
LTR	ERV1	RLTR23	chr9	20689507	20689895 -	0	0	0
SINE	Alu	B1_Mm	chr9	20689919	20690065 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	96045752	96045792 +	0	0	0

SINE	Alu	B1_Mus2	chr9	20964695 20964848 -	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr9	21170237 21170286 +	0	0	0
SINE	Alu	B1_Mm	chr9	21401698 21401844 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	21401845 21401879 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	96141884 96141922 +	0	0	0
SINE	Alu	B1_Mm	chr9	21665077 21665223 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	21897352 21897387 +	6.07381	3.52684	6.05776
SINE	Alu	B1_Mm	chr9	21945636 21945781 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	21945782 21945806 +	0	0	0
SINE	Alu	B1_Mus2	chr9	22076163 22076309 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	22076310 22076335 +	0	0	0
SINE	Alu	B1_Mus2	chr9	22119795 22119945 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	22119946 22119970 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	22963723 22964118 -	0	0	0
SINE	Alu	B1_Mus1	chr9	23305963 23306101 +	0	0	0
SINE	Alu	B1_Mus2	chr9	23382872 23383018 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	23383019 23383091 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	23394251 23394649 +	0	0.380568	0
SINE	Alu	B1_Mur3	chr9	23494365 23494502 -	0	0	0
LTR	MaLR	MTA_Mm	chr9	23871584 23871980 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	24212587 24212624 +	0	0	0
SINE	Alu	B1_Mur4	chr9	24641035 24641219 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	24641220 24641249 +	0	0	0
SINE	Alu	B1_Mus1	chr9	24730533 24730679 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	24730680 24730733 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	24733785 24734181 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	24847570 24847607 +	0	0	0
SINE	B2	B2_Mm1t	chr9	25674550 25674746 +	0	0	0
Low_complexity	Low_complexity	AT_rich	chr9	25674747 25674767 +	0	0	0
SINE	Alu	B1_Mus1	chr9	25821210 25821355 -	0	0	0
LTR	MaLR	MTA_Mm	chr9	25889214 25889606 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	25890701 25891094 +	0	0	0.342623
LINE	L1	L1Md_F2	chr9	26149394 26155666 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	26289114 26289152 +	0	0	0
SINE	Alu	B1_Mus2	chr9	26964459 26964604 -	0	0	0
SINE	Alu	B1_Mm	chr11	96778318 96778455 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	96778456 96778479 +	0.923247	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr9	30321323 30321368 +	0	0	0
SINE	Alu	B1_Mur4	chr10	27737512 27737662 +	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chr10	27737663 27737740 +	0	0	0
SINE	Alu	B1_Mus2	chr9	30815700 30815846 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	30815847 30815880 +	0	0	0
SINE	Alu	B1_Mus2	chr9	31098915 31099060 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	31099061 31099086 +	0	0	0
SINE	Alu	B1_Mus1	chr9	31363750 31363897 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr9	31363898 31363920 +	0	0	0
SINE	Alu	B1_Mus2	chr9	31893915 31894060 +	6.21866	3.21582	5.24386
Simple_repeat	Simple_repeat	(A)n	chr9	31894061 31894089 +	3.39388	1.08938	3.14654
SINE	Alu	B1_Mus2	chr9	32659305 32659447 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	32659448 32659471 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	32919547 32919580 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	33241160 33241553 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	35630443 35630469 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	97149068 97149138 +	0	0	0.615324
SINE	B2	B2_Mm1a	chr11	97218889 97219079 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	97219080 97219100 +	0	0	0
SINE	Alu	B1_Mus1	chr9	37288206 37288359 -	0.332194	0	0
SINE	Alu	B1_Mus1	chr9	38147881 38148018 -	0	0	0
SINE	Alu	B1_Mus1	chr9	38284092 38284238 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	38284239 38284271 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	38313931 38314001 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	39221060 39221117 +	0	0	0
SINE	Alu	B1_Mus1	chr9	39323842 39323989 -	0	0	0
SINE	Alu	B1_Mus2	chr9	39481381 39481529 -	0	0	0
SINE	Alu	B1_Mm	chr9	39660495 39660640 -	0	0	0
SINE	Alu	B1_Mm	chr9	39993078 39993223 -	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr9	39998710 39998746 +	0	0	0

Simple_repeat	Simple_repeat	(TTTA)n	chr9	40030457	40030496 +	0	0	0
SINE	Alu	B1_Mus2	chr9	40400353	40400498 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	40400499	40400532 +	0	0	0
DNA	MER1_type	RCHARR1	chr9	41957231	41958174 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	42767936	42768331 +	0.591053	0	0
SINE	Alu	B1_Mm	chr9	42927531	42927676 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	42927677	42927711 +	0.332194	0	0
SINE	Alu	B1_Mus2	chr9	43385020	43385150 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	43385151	43385218 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	43658833	43658878 +	0	0	0
SINE	Alu	B1_Mus2	chr9	43683869	43684014 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	43684015	43684061 +	0	0	0
SINE	Alu	B1_Mm	chr9	43713452	43713598 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	43713599	43713653 +	0	0	0
SINE	Alu	B1_Mus2	chr11	97656665	97656805 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	97656806	97656835 +	0	0	0
SINE	Alu	B1_Mm	chr11	97701056	97701190 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	97701191	97701235 +	0	0	0
SINE	Alu	B1_Mus1	chr9	44133527	44133674 -	0	0	0
SINE	Alu	B1_Mm	chr9	44227857	44227986 +	0	0	0
SINE	Alu	B1_Mm	chr9	44266842	44266988 -	0	0	0
SINE	Alu	B1_Mus1	chr9	44348512	44348659 -	0	0	0
SINE	Alu	B1_Mus1	chr11	97748173	97748318 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	97748319	97748350 +	0	0	0
SINE	Alu	B1_Mus2	chr9	45645966	45646114 -	0	0	0
SINE	B2	B2_Mm1a	chr9	45767447	45767639 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	45767640	45767666 +	0	0.708814	0
SINE	Alu	B1_Mm	chr9	46144661	46144806 -	0	0	0
SINE	Alu	B1_Mus1	chr9	46349981	46350128 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	46369485	46369526 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr9	48296496	48296535 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	48692351	48692745 -	0	0	0
LTR	MaLR	MTA_Mm	chr9	48734513	48734913 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	48738494	48738892 -	0	0	0
SINE	Alu	B1_Mm	chr9	48965994	48966130 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	48966131	48966155 +	0	0	0
SINE	B4	B4	chr9	48984039	48984350 -	0	0	0
LTR	MaLR	MTA_Mm	chr9	49242116	49242510 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	49809174	49809208 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	50115259	50115294 +	0	0	0
SINE	B2	B2_Mm1a	chr9	50208684	50208876 -	0	0	0
SINE	B2	B2_Mm1t	chr9	50437487	50437679 +	0	0.380568	0.615324
SINE	Alu	B1_Mm	chr9	50849961	50850100 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	50850101	50850123 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	51342700	51342741 +	0	0	0
SINE	Alu	B1_Mur3	chr9	51720923	51721063 +	0	0	0
SINE	Alu	B1_Mm	chr9	52146399	52146545 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	52260007	52260057 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	53592235	53592268 +	0	0	0
LTR	MaLR	MTB	chr9	53607181	53607529 +	0	0	0
SINE	Alu	B1_Mm	chr9	54687197	54687343 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	54687344	54687375 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	98458939	98458976 +	0	0	0
LTR	ERV1	RMER5	chr9	54955204	54955685 -	0	0	0
SINE	B2	B2_Mm1t	chr9	54958893	54959082 -	0	0	0
SINE	B2	B2_Mm1t	chr9	54964508	54964702 +	0	0	0
SINE	Alu	B1_Mus1	chr9	54991821	54991968 -	0	0	0
SINE	Alu	B1_Mus1	chr9	55921402	55921548 +	0	0	0.615324
LTR	ERVK	RMER17C	chr9	55921549	55921910 +	0	0	0.615324
SINE	Alu	B1_Mm	chr9	56415397	56415542 +	0	0	0
SINE	B2	B2_Mm1a	chr9	56590269	56590461 -	0.591053	0	0
SINE	Alu	B1_Mm	chr9	56651763	56651908 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	56651909	56651934 +	0	0	0
SINE	Alu	B1_Mus2	chr9	56832323	56832468 +	1.77316	0.380568	1.23065
Simple_repeat	Simple_repeat	(A)n	chr9	56832469	56832511 +	0.591053	0.380568	1.57327
SINE	B2	B2_Mm1a	chr9	57496781	57496984 +	0	0	0
SINE	Alu	B1_Mus2	chr9	57555431	57555577 -	0	0	0



SINE	Alu	B1_Mm	chr9	57618469 57618614 -	0.332194	0	0
SINE	Alu	B1_Mus2	chr9	57818537 57818694 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	57818695 57818717 +	0	0	0
SINE	Alu	B1_Mur2	chr9	58057786 58057927 +	0.591053	0	0
SINE	B2	B2_Mm1a	chr9	58428496 58428689 -	0	0	0
SINE	Alu	B1_Mus2	chr9	59359654 59359801 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	59359802 59359823 +	0.591053	0	0
SINE	B2	B3A	chr9	59359824 59359979 +	0.591053	0	0
SINE	Alu	B1_Mus1	chr9	59376565 59376710 -	0	0	0.342623
SINE	Alu	PB1D9	chr9	59414309 59414425 -	0	0	0
SINE	Alu	B1_Mm	chr11	98889294 98889420 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr11	98889421 98889446 +	0	0	0.342623
Simple_repeat	Simple_repeat	(TTTA)n	chr9	60399934 60399955 +	0	0	0
SINE	Alu	B1_Mur4	chr11	98957602 98957740 -	0	0	0
SINE	Alu	B1_Mus1	chr9	60888404 60888547 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	60888548 60888595 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	61544106 61544171 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	61587495 61587532 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	61599381 61599779 +	0	0	0
SINE	B2	B2_Mm1a	chr9	61721718 61721908 +	0	0	0
Simple_repeat	Simple_repeat	(CAAAA)n	chr9	61721909 61721926 +	0.332194	0.380568	0
SINE	Alu	B1_Mus1	chr9	61763731 61763878 -	0	0	0
SINE	Alu	B1_Mus2	chr9	62058379 62058524 +	0	0	0
SINE	Alu	B1_Mus2	chr9	62158118 62158264 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	62470928 62471323 +	0	0	0
SINE	Alu	B1_Mus1	chr9	62935639 62935774 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	62935775 62935799 +	0	0	0
SINE	Alu	B1_Mm	chr9	63001853 63001978 +	0	0	0
SINE	Alu	B1_Mus2	chr11	99200152 99200298 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	99200299 99200320 +	0	0	0
SINE	Alu	B1_Mus1	chr9	63698734 63698881 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	99259477 99259503 +	0	0	0
SINE	B4	ID_B1	chr9	63954152 63954368 +	0	0	0
SINE	Alu	B1_Mus2	chr9	64112392 64112537 -	0	0	0
SINE	Alu	B1_Mus2	chr9	64196817 64196961 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	64196962 64196992 +	0	0	0
SINE	Alu	B1_Mus1	chr9	64209943 64210095 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	99326369 99326409 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	64643968 64644055 +	0	0	0
Simple_repeat	Simple_repeat	(T)n	chr11	99419732 99419786 +	0	0.380568	0
SINE	Alu	B1_Mus1	chr9	65035204 65035351 -	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr9	65094002 65094043 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	65186181 65186232 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	65434355 65434378 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr9	65473444 65473587 -	0	0	0
SINE	Alu	B1_Mus2	chr9	66150414 66150554 -	0	0	0
LTR	MaLR	MTA_Mm	chr11	99760709 99761105 +	0	0	0
SINE	Alu	B1_Mm	chr9	66846291 66846435 -	0	0	0
SINE	B2	B2_Mm1a	chr9	66898098 66898288 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	66898289 66898324 +	1.18211	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chr9	66957409 66957438 +	0	0	0
LTR	MaLR	MTB	chr9	67013850 67014152 +	0	0	0
SINE	Alu	B1_Mus2	chr11	99829937 99830083 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	99830084 99830109 +	0	0	0
SINE	B4	B4	chr9	68021105 68021344 -	0	0.380568	0
SINE	B2	B2_Mm1t	chr9	68021354 68021543 -	0	0	0
LTR	MaLR	MTB	chr9	69550849 69551254 +	0	0	0
SINE	Alu	B1_Mus2	chr9	69859642 69859793 -	0	0	0
SINE	B2	B2_Mm1a	chr9	69871683 69871867 -	0	0	0
SINE	Alu	B1_Mm	chr9	69934500 69934644 -	0	0	0
LTR	MaLR	MTB	chr9	72118897 72119341 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	72241773 72241817 +	0	0	0
SINE	Alu	B1_Mus1	chr9	72354561 72354707 -	0	0	0
SINE	Alu	B1_Mus1	chr9	72472519 72472666 -	0	0	0
SINE	Alu	B1_Mus2	chr11	10040477 10040491 -	0	0	0
SINE	Alu	B1_Mm	chr9	73293988 73294134 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	73314792 73314824 +	0.591053	0	0.342623

LTR	MaLR	MTA_Mm	chr9	73927013 73927410 -	0	0	0
SINE	Alu	B1_Mm	chr9	74444053 74444196 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	74763531 74763564 +	0.591053	0	0
SINE	Alu	B1_Mus1	chr9	75268415 75268569 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	75620680 75620716 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	77595262 77595659 -	0	0	0
SINE	Alu	B1_Mm	chr11	10081022 10081036 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	10086472 10086479 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	79866111 79866133 +	0	0	0
LINE	L1	L1Md_F2	chr9	81235045 81241032 -	0.591053	0.708814	0.615324
LTR	MaLR	MTA_Mm	chr9	81260388 81260783 -	0.591053	1.41763	1.23065
LTR	MaLR	MTA_Mm	chr9	81568796 81569153 -	0	0	0
SINE	Alu	B1_Mur3	chr11	10100067 10100082 -	0	0	0
SINE	Alu	B1_Mus1	chr11	10100464 10100478 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	10100478 10100481 +	0	0.761137	0
SINE	Alu	B1_Mus2	chr9	82043569 82043714 +	0.591053	0.708814	0.615324
Low_complexity	Low_complexity	A-rich	chr9	82043715 82043814 +	0.591053	0.708814	0.615324
SINE	Alu	B1_Mm	chr9	82202120 82202266 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	82266385 82266419 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	82532544 82532569 +	0	0	0
SINE	Alu	B1_Mus2	chr9	82751554 82751699 -	0	0	0
SINE	Alu	B1_Mus1	chr9	83496508 83496655 +	0.591053	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr9	83523699 83523733 +	0.591053	0	0
SINE	Alu	B1_Mus1	chr9	83526443 83526573 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	84174911 84174938 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	84279596 84279625 +	0	0	0
LINE	L1	Lx3_Mus	chr9	84308217 84312517 -	0	0	0
LTR	MaLR	MTA_Mm	chr9	85015323 85015707 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	85087100 85087491 -	0	0	0
SINE	Alu	B1_Mus1	chr9	85206814 85206966 +	0	0	0
Simple_repeat	Simple_repeat	(GAA)n	chr9	85206968 85207069 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	85298142 85298174 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	85485563 85485591 +	0	0	0
SINE	Alu	B1_Mm	chr9	85540958 85541105 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	85541106 85541125 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	85569376 85569770 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	85727125 85727160 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	85980437 85980832 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	86029465 86029858 -	0	0	0
LTR	MaLR	MTA_Mm	chr9	86030957 86031350 -	0	0	0
LTR	MaLR	MTA_Mm	chr9	87250535 87250928 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	87297754 87297780 +	0	0	0
SINE	B4	B4	chr9	87323697 87323945 -	0	0	0
LINE	L1	Lx	chr9	88196359 88201999 +	0	0	0
SINE	Alu	B1_Mm	chr9	88435354 88435497 -	0	0	0
SINE	Alu	B1_Mus2	chr9	88976380 88976524 +	0	0	0
SINE	Alu	B1_Mm	chr9	88994319 88994468 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	88994469 88994499 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	89201810 89202207 +	0.591053	0	0.615324
LTR	MaLR	MTA_Mm	chr9	89247070 89247457 -	0	0	0
LTR	MaLR	MTA_Mm	chr9	89255768 89256155 -	0	0	0
SINE	Alu	B1_Mus2	chr9	89260866 89261014 -	0	0	0.615324
SINE	Alu	B1_Mm	chr9	89942533 89942678 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr9	89942679 89942763 +	0	0	0
SINE	Alu	B1_Mus2	chr9	89978268 89978404 -	1.18211	0.761137	0
SINE	Alu	B1_Mus2	chr9	90042441 90042586 +	2.26557	1.01984	1.97174
Simple_repeat	Simple_repeat	(A)n	chr9	90042587 90042615 +	2.26557	1.01984	1.97174
SINE	Alu	B1_Mus2	chr9	90463090 90463236 -	0	0	0
SINE	B2	B2_Mm1t	chr11	10157381 10157400 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	92720575 92720616 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	93038574 93038649 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	93336904 93336929 +	0	0	0
LINE	L1	L1Md_F2	chr9	94019344 94023371 -	0	0	0
LTR	MaLR	MTA_Mm	chr9	94512606 94512994 -	0	0	0
SINE	Alu	B1_Mus2	chr9	95351793 95351939 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	95351940 95351962 +	0	0	0
SINE	B2	B2_Mm1a	chr9	96268057 96268246 -	2.06356	0	0

SINE	Alu	B1_Mm	chr9	96288649 96288795 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	96288796 96288821 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	96718666 96718695 +	0	0.708814	0
SINE	Alu	B1_Mus2	chr11	10189010 10189025 -	0	0	0
SINE	Alu	B1_Mm	chr9	97055340 97055487 +	0	0	0
SINE	Alu	B1_Mur3	chr11	10191731 10191745 +	0	0	0
SINE	Alu	B1_Mus1	chr9	98827422 98827572 -	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	98832029 98832065 +	0.332194	0	0
LTR	MaLR	MTA_Mm	chr9	99121751 99122135 +	0	0	0
SINE	B2	B2_Mm2	chr9	99492689 99492862 -	0	0	0
LTR	MaLR	MTA_Mm	chr9	99578774 99579168 -	0	0	0
SINE	Alu	B1_Mus2	chr9	99584245 99584393 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	99584394 99584429 +	0	0	0
SINE	Alu	B1_Mus1	chr11	10214340 10214355 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	10214355 10214358 +	0	0	0
SINE	Alu	B1_Mus2	chr9	10047297 10047311 -	0	0	0
SINE	Alu	B1_Mus2	chr9	10053328 10053343 -	0	0	0
SINE	Alu	B1_Mm	chr11	10216314 10216328 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	10216328 10216332 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	10115842 10115846 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	10220343 10220345 +	0	0	0
SINE	Alu	B1_Mm	chr11	10225156 10225170 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	10225170 10225175 +	0	0	0
SINE	Alu	B1_Mm	chr9	10269506 10269520 -	0	0	0
SINE	Alu	B1_Mus1	chr9	10300548 10300564 -	0.591053	0	0
LTR	MaLR	MTD	chr9	10300685 10300728 +	0	0	0
SINE	Alu	B1_Mus2	chr11	10242123 10242137 -	0	0	0
SINE	Alu	B1_Mm	chr11	10243705 10243720 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	10244890 10244892 +	0.591053	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr11	10245994 10245998 +	0.332194	0	0
LTR	MaLR	MTA_Mm	chr9	10546139 10546178 -	0	0	0
LTR	MaLR	MTA_Mm	chr9	10547731 10547770 +	0	0	0
SINE	Alu	B1_Mm	chr9	10574039 10574053 -	0	0	0
SINE	B2	B2_Mm1a	chr9	10600794 10600813 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr9	10600813 10600815 +	0	0	0
SINE	Alu	B1_Mus2	chr9	10603770 10603784 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr9	10603784 10603789 +	0	0	0
SINE	Alu	B1_Mm	chr9	10612264 10612278 +	0	0	0
LTR	MaLR	MTB	chr9	10625466 10625504 +	0	0	0
SINE	Alu	B1_Mus1	chr10	28724109 28724256 -	0	0	0
SINE	Alu	B1_Mm	chr9	10820404 10820417 -	0	0	1.23065
SINE	Alu	B1_Mus2	chr9	10824805 10824819 +	0	0	0.615324
SINE	Alu	B1_Mm	chr9	10840646 10840661 -	0	0	0
SINE	Alu	B1_Mus2	chr9	10856167 10856181 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	10856181 10856186 +	0	0	0
LINE	L1	L1Md_A	chr9	10914750 10915457 -	0.923247	0	0
SINE	Alu	B1_Mus2	chr9	10961284 10961299 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	10977767 10977771 +	0	0	0
SINE	Alu	B1_Mus1	chr9	10981836 10981851 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	10981851 10981853 +	0	0	0
SINE	Alu	B1_Mus2	chr11	10288580 10288593 -	0	0	0
SINE	Alu	B1_Mus2	chr9	11043111 11043125 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	11043125 11043128 +	0	0	0
SINE	B2	B2_Mm1t	chr9	11069903 11069921 -	0	0	0
SINE	Alu	B1_Mus1	chr9	11142407 11142422 -	0	0	0
Simple_repeat	Simple_repeat	(CCAA)n	chr9	11174160 11174163 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	11174163 11174167 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	11290739 11290778 +	0.591053	0	0
LTR	ERVL	RMER15	chr11	10315778 10315837 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	11363485 11363488 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	11396215 11396217 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	11418057 11418061 +	0	0	0
scRNA	scRNA	4.5SRNA	chr9	11458051 11458060 -	40.5997	21.5213	21.4602
SINE	B2	B2_Mm1a	chr11	10333390 10333410 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	10333410 10333412 +	0.332194	0	0
SINE	Alu	B1_Mus1	chr9	11571403 11571417 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	11571417 11571427 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr9	11577988 11577992 +	0	0	0
SINE	Alu	B1_Mus1	chr9	11599436 11599451 -	0	0	0
SINE	Alu	B1_Mm	chr10	28845289 28845435 +	0	0	0
LTR	ERVK	RLTR15	chr9	11647872 11647900 -	0	0	0
SINE	Alu	B1_Mus2	chr11	10362220 10362235 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	11786902 11786905 +	0	0	0
SINE	Alu	B1_Mus2	chr9	11937524 11937538 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	12038848 12038851 +	0	0.708814	0
SINE	Alu	B1_Mm	chr9	12044404 12044418 -	0	0	0
SINE	Alu	B1_Mus2	chr9	12066168 12066182 +	0	0	0
SINE	B2	B2_Mm1a	chr9	12144150 12144169 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr9	12146979 12146983 +	0	0	0
SINE	B2	B2_Mm2	chr9	12252102 12252117 +	2.06418	1.08938	0
Simple_repeat	Simple_repeat	(A)n	chr9	12255155 12255161 +	1.25544	0.761137	0.957947
SINE	B2	B2_Mm1a	chr9	12306655 12306674 +	0	0	0
LTR	MaLR	MTA_Mm	chr9	12313089 12313128 +	0	0	0
SINE	Alu	B1_Mus1	chr9	12313857 12313872 -	0	0	0
SINE	Alu	B1_Mus2	chr9	12338370 12338385 +	0.332194	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr9	12338385 12338387 +	0.332194	0	0.615324
SINE	Alu	B1_Mm	chrX	3770627 3770772 -	0	0	0
LINE	L1	L1Md_T	chrX	3836687 3843352 -	0	0	0
SINE	Alu	B1_Mus2	chrX	5655774 5655918 +	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chrX	5655919 5656005 +	0	0	0
LTR	MaLR	MTB_Mm	chrX	6933336 6933747 -	0	0	0
SINE	Alu	B1_Mm	chrX	7115294 7115440 -	0	0	0
SINE	Alu	B1_Mus2	chrX	7269834 7269957 -	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chrX	7292459 7292485 +	0	0	0
SINE	Alu	B1_Mus1	chrX	7526856 7527003 -	0	0	0
SINE	Alu	B1_Mus1	chrX	7570994 7571146 +	0	0	0
LTR	MaLR	ORR1C2	chrX	7571162 7571512 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	7614098 7614490 -	0	0	0
SINE	Alu	B1_Mm	chrX	7668042 7668188 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	7668189 7668213 +	0	0	0
SINE	Alu	B1_Mur2	chrX	7669740 7669887 +	0	0	0
Low_complexity	Low_complexity	A-rich	chrX	7669888 7669957 +	0	0	0
SINE	Alu	B1_Mm	chrX	7761042 7761190 -	1.84649	1.41763	0.615324
LTR	MaLR	MTA_Mm	chrX	9134901 9135296 +	0	0	0
SINE	Alu	B1_Mm	chr10	5748221 5748349 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	9904355 9904750 -	0	0	0.615324
SINE	B2	B2_Mm1t	chrX	10204472 10204672 -	0	0	0
SINE	Alu	B1_Mus1	chrX	10383269 10383405 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	10383406 10383438 +	0	0	0
SINE	Alu	B1_Mm	chrX	10407100 10407221 -	0	0	0
LINE	L1	Lx6	chrX	10503178 10509625 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	10817999 10818034 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	11605626 11606020 +	0	0	0
SINE	B4	RSINE1	chrX	11786086 11786243 -	0	0	0
SINE	Alu	B1_Mus2	chrX	11964377 11964499 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	11964500 11964524 +	0	0	0
SINE	B4	ID_B1	chrX	12136314 12136515 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	12136516 12136539 +	0	0	0
SINE	Alu	B1_Mm	chrX	12146322 12146467 -	0	0	0
Low_complexity	Low_complexity	GA-rich	chrX	12406096 12406264 +	0	0	0.685246
SINE	Alu	B1_Mus1	chrX	12588910 12589057 +	0	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chrX	12589058 12589077 +	0	0.380568	0
SINE	Alu	B1_Mus1	chrX	12642007 12642153 -	0	0	0
SINE	Alu	B1_Mus1	chrX	12848109 12848260 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	12848261 12848293 +	0	0	0
SINE	Alu	B1_Mur1	chrX	13434375 13434541 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chrX	13434542 13434567 +	0	0	0
SINE	Alu	B1_Mm	chrX	13473255 13473401 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	13867880 13867925 +	0	0	0
LINE	L1	L1Md_F2	chrX	13978907 13984325 +	0	0	0
LINE	L1	L1_Mus3	chrX	14107344 14113488 -	0	0	0
SINE	Alu	B1_Mm	chrX	14128954 14129099 -	0	0	0
LINE	L1	L1_Mus1	chrX	14442771 14447679 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	14602431 14602825 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chrX	15062422 15062462 +	0	0	0
SINE	Alu	B1_Mm	chrX	17007823 17007952 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	17007953 17007981 +	0.591053	0	0
SINE	Alu	PB1D9	chrX	17547820 17547917 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	17547918 17547937 +	0	0	0
LINE	L1	MusHAL1	chrX	18073917 18078596 -	0	0	0
LINE	L1	L1Md_T	chrX	18763899 18770362 -	0	0	0
SINE	Alu	B1F	chrX	18813069 18813172 +	0	0	0
SINE	Alu	B1_Mus1	chrX	19051267 19051414 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	19051415 19051445 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	19086841 19086884 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	19117062 19117460 +	0	0	0
SINE	Alu	B1_Mus2	chrX	19391514 19391659 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	19391660 19391684 +	0	0	0
SINE	Alu	B1_Mm	chrX	19929425 19929582 -	0	0	0
SINE	Alu	B1_Mus1	chrX	20158965 20159104 -	0	0	0
SINE	Alu	B1_Mus1	chrX	20164798 20164932 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	20164933 20164962 +	0	0	0
SINE	Alu	B1_Mus2	chrX	20337050 20337196 -	0	0	0
SINE	Alu	B1_Mm	chrX	20363814 20363938 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	20363939 20363986 +	0	0	0
SINE	Alu	B1_Mus1	chrX	20573732 20573879 -	0	0.708814	0
SINE	Alu	B1_Mm	chrX	20990285 20990431 +	0.591053	0	0
SINE	Alu	B1_Mm	chr11	10592422 10592435 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	10592435 10592437 +	0	0	0
SINE	B2	B2_Mm1t	chrX	21049478 21049674 +	0	0.708814	0
SINE	Alu	PB1D9	chrX	21178408 21178529 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chrX	21518832 21518870 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	22422718 22423114 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	22671925 22671959 +	0	0	0
LINE	L1	L1Md_F2	chrX	23425380 23431778 -	0	0	0
LINE	L1	L1_Mus1	chrX	24200568 24205958 -	0	0	0
LINE	L1	L1_Mus1	chrX	24307301 24313511 +	0	0	0
SINE	Alu	B1_Mur3	chrX	24615743 24615888 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	24615889 24615921 +	0	0	0
LINE	L1	L1_Mus1	chrX	24947154 24952958 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	25145525 25145565 +	0	0	0
LINE	L1	L1_Mus1	chrX	25222976 25228779 +	0	0	0
LINE	L1	L1Md_F	chrX	25510300 25516367 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	26547155 26547196 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	27315557 27315611 +	0	0	0
LINE	L1	L1Md_F3	chrX	27397079 27402517 +	0	0	0
SINE	Alu	B1_Mur3	chrX	27483384 27483529 -	0	0	0
LINE	L1	L1Md_F2	chrX	27492770 27497960 +	0	0	0
LINE	L1	L1Md_F3	chrX	28397342 28402766 +	0	0	0
SINE	Alu	B1_Mur3	chrX	28483635 28483780 -	0	0	0
LINE	L1	L1Md_F2	chrX	28493021 28498206 +	0	0	0
SINE	Alu	B1_Mur3	chrX	29246009 29246154 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	29246155 29246181 +	0	0	0
SINE	Alu	B1_Mur3	chrX	29581740 29581885 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	29581886 29581925 +	0	0	0
LINE	L1	L1_Mus1	chrX	29591169 29597144 -	0	0	0
SINE	Alu	B1_Mur3	chrX	29837398 29837543 -	0	0	0
SINE	Alu	B1_Mus2	chrX	29839263 29839409 -	0	0	0
LINE	L1	L1Md_F2	chrX	29846793 29852177 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	30009446 30009488 +	0	0	0
SINE	Alu	B1_Mm	chrX	30620851 30620996 -	0	0	0
SINE	Alu	B1_Mm	chrX	31723097 31723242 +	0	0	0
LINE	L1	L1Md_F2	chrX	32366460 32372830 -	0	0	0
SINE	Alu	B1_Mus2	chrX	32380214 32380360 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	32380361 32380383 +	0	0.380568	0
SINE	Alu	B1_Mur3	chrX	32382080 32382225 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	32382226 32382253 +	0	0	0
LINE	L1	L1_Mus1	chrX	32987990 32994830 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	33286170 33286576 +	0	0	0
SINE	Alu	B1_Mm	chr11	10658035 10658049 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	10658049 10658051 +	0	0	0

LTR	MaLR	MTA_Mm	chrX	34078088 34078483 +	0	0	0
SINE	Alu	B1_Mus1	chrX	34343866 34343994 -	0	0	0
SINE	ID	ID4	chrX	35225096 35225167 -	0	0	0
SINE	Alu	B1_Mus1	chrX	35941676 35941878 +	0	0	0
LINE	L1	L1_Mus2	chrX	36094670 36100852 -	0	0	0
LINE	L1	L1_Mus2	chrX	37142544 37147876 -	0	0	0.342623
LINE	L1	L1_Mus1	chrX	38126202 38130691 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	38357175 38357224 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	38598062 38598457 -	0	0	0
LINE	L1	L1Md_T	chrX	39052200 39059066 -	0	0	0
SINE	Alu	B1_Mm	chr11	10687144 10687158 -	0	0	0
SINE	Alu	B1_Mus1	chr11	10687548 10687562 -	0	0	0
LINE	L1	L1Md_T	chrX	39342191 39348979 -	0	0	0
SINE	Alu	B1_Mus1	chrX	39396760 39396910 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	39396911 39396934 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	39709152 39709197 +	0	0	0
SINE	Alu	B1_Mm	chr11	10702019 10702034 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	41802739 41803120 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	42599160 42599199 +	0	0	0
LINE	L1	L1Md_F2	chrX	42925892 42931989 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	43109274 43109317 +	0	0	0
SINE	Alu	B1_Mus2	chrX	43856437 43856582 +	0	0	0
SINE	Alu	B1_Mus2	chrX	43888567 43888711 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	44329039 44329429 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	44337179 44337574 +	0	0	0
SINE	Alu	B1_Mus1	chrX	44975345 44975470 -	0	0	0
SINE	Alu	B1_Mus2	chrX	45678722 45678867 -	0	0	0
SINE	Alu	B1_Mus2	chrX	45689743 45689895 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	46080407 46080794 -	0	0	0
LINE	L1	L1Md_T	chrX	46961262 46967207 -	0	0	0
SINE	Alu	B1_Mus2	chrX	47251474 47251615 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	47251616 47251638 +	0	0	0
SINE	Alu	B1_Mus1	chrX	47261607 47261754 -	0	0	0
SINE	Alu	B1_Mm	chr11	10736596 10736611 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	49043878 49044271 -	0	0	0
SINE	Alu	B1_Mus1	chrX	49204450 49204596 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	49288228 49288623 +	0	0	0
SINE	Alu	B1_Mus2	chrX	49617011 49617156 -	0	0	0
SINE	Alu	B1_Mus2	chrX	50205098 50205244 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	50264906 50264934 +	0.332194	0	0
LTR	MaLR	MTA_Mm	chrX	50386753 50387148 +	0	0	0
SINE	Alu	B1F	chrX	50737008 50737155 -	0	0	0
LINE	L1	L1Md_T	chrX	50961676 50968149 -	0	0	0
SINE	Alu	PB1D7	chrX	51027592 51027696 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	51027697 51027742 +	0	0	0
SINE	Alu	B1_Mm	chr11	10759158 10759172 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	10759172 10759175 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	53135811 53136205 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	53588379 53588776 +	0	0	0
SINE	Alu	B1_Mus1	chr10	29583105 29583238 -	0	0	0
SINE	Alu	PB1D9	chrX	54102827 54102929 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	54813850 54814245 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	55319184 55319219 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	55473941 55474336 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	56176249 56176281 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	56477498 56477880 +	0	0	0
LINE	L1	L1Md_T	chrX	56713325 56719726 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	57179252 57179646 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	57436692 57437081 -	0	0	0.615324
LTR	MaLR	MTA_Mm	chrX	57438158 57438503 -	0	0	0
SINE	Alu	B1_Mm	chrX	57866645 57866781 -	0	0	0
LINE	L1	L1Md_T	chrX	58512383 58519652 -	0	0	0
LINE	L1	L1Md_T	chrX	59215611 59222799 -	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	59876571 59876616 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	59965929 59966323 -	0	0	0
LINE	L1	L1Md_F	chrX	59972643 59977359 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	60230004 60230399 +	0	0	0

SINE	Alu	B1_Mus2	chrX	60427427 60427572 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	60813562 60813605 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	60837569 60837604 +	0	0	0
SINE	Alu	B1_Mus2	chrX	61237437 61237559 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chrX	61237560 61237602 +	0	0	0
SINE	Alu	B1_Mm	chr10	29692060 29692191 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	29692192 29692215 +	0	0	0
LINE	L1	L1_Mus1	chrX	61717733 61724157 -	0	0	0
LINE	L1	L1Md_T	chrX	61879182 61886544 -	0	0	0
SINE	Alu	B1_Mur3	chr11	10872521 10872536 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	10872536 10872539 +	0.332194	0	0
SINE	Alu	B1_Mur4	chr11	10873767 10873781 -	0	0	0
SINE	Alu	B1_Mm	chrX	63079770 63079915 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	63079916 63079944 +	0	0	0
LINE	L1	L1_Mus3	chrX	63131420 63135933 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	63755098 63755134 +	0	0	0
SINE	Alu	B1_Mus1	chrX	63928201 63928351 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	63928352 63928374 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	64128922 64128985 +	0	0	0
LTR	ERVK	RMER19A	chrX	64297207 64297972 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	64680136 64680531 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	64715038 64715072 +	0	0	0
LINE	L1	L1_Mus3	chrX	64734841 64741556 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	64856642 64856675 +	0	0	0
SINE	Alu	B1_Mus2	chrX	64861314 64861460 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	64861461 64861486 +	0	0	0
LINE	L1	L1Md_F2	chrX	65145337 65151454 -	0.591053	0	0
SINE	Alu	B1_Mus2	chr11	10890723 10890738 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	10890738 10890740 +	0	0	0
SINE	Alu	B1_Mus2	chrX	65977044 65977190 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	66259233 66259270 +	0	0	0.615324
LINE	L1	L1Md_F2	chrX	66524249 66530303 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	67662882 67663277 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	67783413 67783808 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	67808526 67808920 +	0.332194	0	0
SINE	Alu	B1_Mus2	chrX	68767453 68767599 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	68767600 68767693 +	0	0	0
SINE	Alu	B1_Mus1	chrX	68784359 68784493 -	0	0	0
SINE	Alu	B1_Mm	chrX	68828398 68828544 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	68828545 68828604 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	68881198 68881585 +	0	0	0
SINE	Alu	B1_Mm	chr11	10916289 10916303 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	10916303 10916306 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	69274495 69274534 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	70285603 70285632 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	70308217 70308258 +	0	0	0
SINE	Alu	B1_Mus2	chrX	70996175 70996320 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	70996321 70996344 +	0	0	0
SINE	Alu	B1_Mus2	chrX	71508204 71508350 -	0	0	0
SINE	Alu	B1_Mus1	chrX	71537784 71537937 +	1.18211	0.380568	1.23065
Simple_repeat	Simple_repeat	(A)n	chrX	71537938 71537973 +	1.18211	0.380568	0.615324
SINE	Alu	B1_Mm	chrX	71808214 71808360 -	0	0	0
SINE	Alu	B1_Mus1	chrX	72024375 72024496 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	72024497 72024519 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	72720343 72720378 +	0	0	0
SINE	Alu	B1_Mm	chrX	72951365 72951511 +	0	0	0
LINE	L1	L1Md_F	chrX	73500002 73504695 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	73880287 73880681 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	75061481 75061879 -	0	0	0
SINE	Alu	B1_Mus1	chrX	75178080 75178227 +	0	0	0
Low_complexity	Low_complexity	A-rich	chrX	75178228 75178289 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	75438289 75438323 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	75583441 75583477 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	76045662 76046056 +	0	0.380568	0
LTR	MaLR	MTA_Mm	chrX	76229316 76229699 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	76231748 76232142 -	0	0	0
SINE	Alu	B1_Mus2	chrX	76253395 76253539 +	0	0	0

Simple_repeat	Simple_repeat	(GAAA)n	chrX	76253540 76253605 +	0	0	0
SINE	Alu	B1_Mus1	chrX	76461738 76461884 -	0	0	0
SINE	Alu	B1_Mus2	chrX	76651152 76651298 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	76843725 76844122 +	0	0	0
SINE	Alu	B1_Mus1	chrX	76951577 76951724 -	0	0	0
SINE	Alu	B1_Mus1	chrX	77024945 77025092 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	77642740 77642796 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	77953157 77953551 -	0	0	0
LINE	L1	L1Md_T	chrX	78204440 78212164 -	0	0	0
LINE	L1	L1Md_F2	chrX	78423161 78429108 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	78548413 78548803 -	0	0	0
SINE	Alu	B1_Mm	chr10	29869878 29870023 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	78652855 78653250 +	0	0	0
SINE	Alu	B1_Mm	chrX	78986725 78986871 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	79158298 79158693 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chrX	79247284 79247320 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	79374472 79374867 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	80023819 80024213 -	0	0	0
SINE	Alu	B1_Mm	chrX	80140508 80140654 +	2.10535	2.50701	2.4613
Simple_repeat	Simple_repeat	(A)n	chrX	80140655 80140675 +	2.10535	1.41763	2.4613
LTR	MaLR	MTB	chrX	80144189 80144567 +	0.591053	0	1.91589
LINE	L1	Lx3_Mus	chr10	29889781 29895175 -	0	0	0.342623
LTR	MaLR	MTA_Mm	chrX	83205945 83206339 -	0	0	0
SINE	Alu	B1_Mm	chrX	83331486 83331627 +	0	0	0
SINE	Alu	B1_Mus1	chrX	83505415 83505560 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	85403345 85403392 +	0	0	0
LINE	L1	L1_Mus1	chrX	85801837 85806940 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	85936259 85936653 -	0	0	0
SINE	Alu	B1_Mm	chrX	85960298 85960443 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	86362277 86362314 +	0	0	0
SINE	Alu	B1_Mus1	chrX	86635814 86635960 -	0	0.708814	0
SINE	Alu	B1_Mus2	chrX	86680092 86680238 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	86680239 86680301 +	0	0	0
SINE	Alu	B1_Mus1	chrX	87031589 87031735 -	0	0	0
SINE	Alu	B1_Mm	chrX	87038185 87038330 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	87433908 87433951 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	87442213 87442248 +	0	0	0
SINE	Alu	B1_Mus1	chrX	88463756 88463902 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	88463903 88463928 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	88603630 88604027 +	0	0	0
SINE	Alu	B1_Mus2	chrX	88750534 88750626 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	88750627 88750661 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	11075235 11075239 +	0	0	0
SINE	Alu	B1_Mus2	chrX	90392811 90392957 -	0	0	0
SINE	Alu	B1_Mus2	chrX	91288957 91289103 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	91309011 91309050 +	0	0.380568	0
SINE	Alu	PB1D7	chrX	91386993 91387110 -	0	0	0
SINE	Alu	B1_Mur3	chrX	91396885 91397031 -	0	0	0
SINE	Alu	B1_Mus2	chrX	91459263 91459409 -	0	0	0
SINE	Alu	B1_Mur1	chrX	91535037 91535176 -	0	0	0
SINE	Alu	B1_Mus1	chrX	91904744 91904890 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	11111208 11111210 +	0	0	0
SINE	Alu	B1_Mus1	chrX	92833285 92833431 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	92833432 92833473 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	11120949 11120952 +	0	0	0
SINE	Alu	B1_Mus1	chrX	93184887 93185033 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	93185034 93185081 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	93940612 93941006 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	93982441 93982836 +	0	0	0
LINE	L1	L1VL4	chrX	94975429 94978333 -	0	0	0
LINE	L1	L1Md_T	chrX	95164412 95171295 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	95731169 95731563 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	96666216 96666258 +	0	0.380568	0.342623
LINE	L1	L1Md_T	chrX	97954332 97960794 -	0	0	0
SINE	Alu	B1_Mm	chrX	98549137 98549289 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	98549290 98549314 +	0	0	0
SINE	Alu	B1_Mus1	chrX	98678722 98678869 +	0.591053	0	0



Simple_repeat	Simple_repeat	(A)n	chrX	98678870 98678898 +	0.591053	0	0
SINE	Alu	B1_Mus2	chrX	98721081 98721227 -	0	0	0
SINE	Alu	B1_Mus1	chrX	98727146 98727293 -	0	0	0
SINE	Alu	B1_Mus2	chrX	98826874 98827020 -	0	0	0
SINE	Alu	B1F	chrX	99000096 99000245 -	0	0	0
SINE	Alu	B1_Mm	chrX	99356474 99356620 -	0	0	0
SINE	Alu	B1_Mm	chrX	10046250 10046264 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	10047744 10047748 +	0	0	0
SINE	Alu	B1_Mus2	chrX	10052782 10052797 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	10052797 10052799 +	0	0	0
SINE	Alu	B1_Mm	chrX	10054123 10054135 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	10060565 10060568 +	0	0	0
SINE	Alu	B1_Mm	chrX	10070424 10070439 +	1.05163	0.380568	0
Low_complexity	Low_complexity	A-rich	chrX	10070439 10070449 +	0	0.380568	0
SINE	Alu	B1_Mus2	chrX	10075420 10075434 -	0.591053	0	0
LTR	MaLR	MTA_Mm	chr11	11215109 11215148 -	0	0	0
SINE	Alu	B1_Mm	chr11	11215765 11215777 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	11215777 11215781 +	0	0.380568	0
SINE	Alu	B1_Mus2	chrX	10143576 10143591 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	10143591 10143594 +	0	0	0
SINE	Alu	B1_Mus2	chrX	10166401 10166416 -	0	0	0
SINE	Alu	B1_Mus2	chrX	10187851 10187866 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	10221520 10221523 +	0	0	0
LINE	L1	L1Md_F2	chrX	10245206 10245725 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	10264049 10264088 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	10285325 10285333 +	0	0	0
SINE	Alu	B1_Mus2	chrX	10338026 10338041 -	0	0	0
LINE	L1	L1_Mus2	chrX	10354308 10354887 +	0	0	0
SINE	B2	B2_Mm2	chrX	10411166 10411185 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	10493383 10493422 -	0	0	0
SINE	Alu	B1_Mm	chrX	10520670 10520685 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	10527906 10527910 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	10542883 10542887 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	10548623 10548629 +	0	0	0
LTR	MaLR	MTB	chrX	10554752 10554796 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	10665685 10665724 -	0	0	0
SINE	Alu	B1_Mm	chrX	10670172 10670189 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	10712103 10712105 +	0	0	0
SINE	Alu	B1_Mm	chrX	10745892 10745906 -	0	0	0
SINE	Alu	B1_Mm	chrX	10836966 10836981 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	10836981 10836983 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chrX	10882031 10882035 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	10931748 10931788 +	0	0	0
Simple_repeat	Simple_repeat	(GAA)n	chr11	11289099 11289111 +	0	0	0
LINE	L1	L1_Mus1	chrX	10983472 10984064 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	11101654 11101658 +	0	0	0
LINE	L1	L1_Mus1	chrX	11127549 11128171 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	11130627 11130666 -	0	0	0
LINE	L1	L1Md_T	chrX	11137429 11138163 -	0	0	0
LINE	L1	L1VL4	chrX	11196169 11196664 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	11381326 11381330 +	0	0	0
SINE	Alu	B1_Mm	chrX	11390320 11390335 -	0	0	0
LINE	L1	L1Md_F2	chrX	11402905 11403545 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	11422895 11422902 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	11517966 11517969 +	0	0	0
LINE	L1	L1_Mus2	chrX	11554416 11554876 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	11575523 11575563 +	0	0	0
LINE	L1	L1Md_F2	chrX	11586980 11587588 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	11607090 11607093 +	0	0	0
SINE	Alu	B1_Mm	chr11	11348949 11348963 +	1.5143	0	0.615324
SINE	Alu	B1_Mm	chrX	11713125 11713140 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	11715772 11715775 +	0	0	0
LINE	L1	L1Md_T	chrX	11807278 11807926 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	11822297 11822302 +	0.332194	0	0
SINE	Alu	B1_Mus1	chrX	11846744 11846759 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	11846759 11846762 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	11863382 11863386 +	0	0	0

SINE	Alu	B1_Mm	chr11	11361402	11361416 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	11361416	11361418 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	11902958	11902962 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	30343343	30343377 +	0	0	0.342623
SINE	Alu	B1_Mus1	chrX	11928538	11928553 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	11928553	11928560 +	0	0	0
LINE	L1	L1Md_F2	chrX	11934612	11935158 +	0	0	0
SINE	Alu	B1_Mus1	chrX	11938701	11938717 +	0	0	0
Low_complexity	Low_complexity	AT-rich	chrX	11938717	11938720 +	0	0	0
SINE	Alu	B1_Mus2	chr11	11363572	11363586 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	11363586	11363589 +	0	0	0
SINE	Alu	B1_Mus1	chrX	11993797	11993812 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	11993812	11993816 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	12016307	12016309 +	0	0	0
Simple_repeat	Simple_repeat	(GGAA)n	chrX	12016309	12016319 +	0	0	0
Low_complexity	Low_complexity	GA-rich	chrX	12018698	12018707 +	0	0	0
LTR	MaLR	MTA_Mm	chr10	30353161	30353551 -	0	0	0
SINE	Alu	B1_Mus1	chrX	12044259	12044274 -	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chrX	12059983	12059997 +	0	0	0
Low_complexity	Low_complexity	A-rich	chrX	12060564	12060572 +	0.591053	0	0
SINE	Alu	B1_Mus1	chrX	12067209	12067223 -	0	0	0.615324
SINE	Alu	B1_Mm	chrX	12074235	12074249 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	12074249	12074254 +	0	0	0
SINE	Alu	B1_Mus1	chrX	12082075	12082090 +	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chrX	12082090	12082103 +	0	0	0
Low_complexity	Low_complexity	A-rich	chrX	12083258	12083266 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	12084894	12084897 +	0.332194	0.380568	0
Simple_repeat	Simple_repeat	(A)n	chrX	12090357	12090362 +	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chrX	12104027	12104040 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	12105187	12105191 +	0	0	0
SINE	Alu	B1_Mus1	chrX	12132437	12132451 +	0	0	0
Low_complexity	Low_complexity	GA-rich	chrX	12132451	12132465 +	0	0	0
SINE	Alu	B1_Mus1	chrX	12133787	12133801 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	12133802	12133804 +	0	0	0
SINE	Alu	B1_Mus2	chrX	12142364	12142379 -	0	0	0
SINE	Alu	B1_Mus1	chrX	12161183	12161198 -	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chrX	12185339	12185352 +	0	0	0
Low_complexity	Low_complexity	A-rich	chrX	12186535	12186544 +	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chrX	12197314	12197327 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	12198245	12198250 +	0	0	0
SINE	Alu	B1_Mus2	chrX	12203903	12203918 +	0	0	0
Low_complexity	Low_complexity	A-rich	chrX	12203918	12203926 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	12207733	12207736 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	12258899	12258903 +	0	0	0
SINE	Alu	B1_Mm	chrX	12269768	12269783 -	0.332194	0	0.615324
LTR	MaLR	MTA_Mm	chrX	12303050	12303090 -	0	0	0
LINE	L1	L1_Mus3	chrX	12311133	12311724 -	0	0	0
SINE	Alu	B1_Mm	chrX	12364054	12364068 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	12364068	12364071 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	12367979	12367982 +	0	0	0
LTR	ERVK	RMER17D2	chrX	12376591	12376682 -	0	0	0
LINE	L1	L1_Mus2	chrX	12377677	12378252 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	12380100	12380139 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	12428413	12428452 -	0	0	0
LINE	L1	L1Md_F2	chrX	12439704	12440227 +	0	0	0
SINE	Alu	B1_Mus2	chr11	11409151	11409165 -	0	0	0
SINE	Alu	B1_Mus1	chrX	12536123	12536137 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	12602919	12602924 +	0.591053	0	0
SINE	Alu	B1_Mus2	chrX	12705050	12705064 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	12719865	12719869 +	0	0	0
SINE	Alu	B1_Mm	chrX	12751221	12751236 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chrX	12833580	12833586 +	0	0	0
SINE	Alu	B1_Mm	chrX	12846020	12846034 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	12846034	12846038 +	0	0	0
SINE	Alu	B1_Mm	chrX	12851886	12851900 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	12859188	12859192 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	12948744	12948748 +	0	0	0

SINE	Alu	B1_Mus2	chrX	12954976 12954991 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	12954991 12954995 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	12972357 12972360 +	0	0	0
SINE	Alu	B1F1	chr11	11446734 11446748 -	0.332194	0	0
SINE	B2	B2_Mm1a	chr11	11446778 11446797 -	0	0	0
LINE	L1	L1Md_F	chrX	13056866 13057404 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	11452898 11452901 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	13092913 13092953 -	0	0	0
SINE	Alu	B1_Mus1	chr11	11453608 11453622 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	11453622 11453625 +	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chrX	13101170 13101172 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	13138925 13138965 -	0	0	0
LTR	MaLR	MTB	chrX	13239302 13239340 -	0	0	0
LTR	MaLR	MTB	chrX	13280012 13280050 -	0	0	0
SINE	Alu	B1_Mus1	chrX	13280873 13280887 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	13280887 13280889 +	0	0	0
SINE	Alu	B1_Mm	chrX	13320716 13320731 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	13320731 13320734 +	0	0	0
SINE	Alu	B1_Mus2	chrX	13322542 13322557 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	13322557 13322560 +	0	0	0
SINE	Alu	B1_Mus1	chr11	11470984 11470999 -	0	0	0
SINE	Alu	B1_Mus2	chrX	13374475 13374490 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	13374490 13374496 +	0	0	0
SINE	Alu	B1_Mus1	chrX	13592512 13592527 -	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chrX	13733528 13733536 +	0	0	0
LINE	L1	L1Md_T	chrX	13825530 13826156 -	0	0	0
SINE	Alu	B1_Mus2	chrX	13828480 13828495 -	0	0	0
LINE	L1	L1Md_F2	chrX	13846608 13847226 -	0	0	0
SINE	Alu	B1_Mus2	chrX	13882910 13882925 -	0	0	0
SINE	Alu	B1_Mus1	chr10	30548091 30548236 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	30548237 30548272 +	0	0.708814	0
SINE	Alu	B1_Mus2	chrX	13949225 13949240 +	1.05163	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	13949240 13949246 +	0	0	0.685246
Simple_repeat	Simple_repeat	(A)n	chrX	13967094 13967100 +	0.591053	0	0
SINE	Alu	B1_Mus2	chrX	14071482 14071497 -	0	0	0
Low_complexity	Low_complexity	AT_rich	chrX	14072383 14072386 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	14073316 14073355 -	0	0	0
SINE	Alu	B1_Mus2	chrX	14164145 14164159 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	14164159 14164165 +	0	0	0
SINE	Alu	PB1D10	chrX	14204665 14204675 -	0.591053	0	0
LINE	L1	L1Md_F2	chrX	14258308 14258827 +	0	0	0
LTR	MaLR	MTD	chrX	14263548 14263591 -	0.591053	0	0
SINE	Alu	B1_Mm	chr11	11541286 11541301 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	11541301 11541303 +	0	0	0
SINE	Alu	B1_Mm	chr11	11541879 11541893 +	0	0	0
LINE	L1	L1Md_F2	chrX	14318003 14318646 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	14387704 14387710 +	1.49234	1.08938	2.29225
LTR	MaLR	MTA_Mm	chrX	14388900 14388939 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	14401103 14401141 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	14435337 14435376 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	14435530 14435534 +	0	0	0
LTR	MaLR	MTA_Mm	chrX	14453421 14453460 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	14502244 14502283 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	14521425 14521464 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	14543466 14543505 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	14580602 14580641 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	14601306 14601313 +	0	0	0
SINE	B2	B2_Mm2	chr11	11558649 11558666 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	14614186 14614225 -	0	0	0
SINE	B2	B3A	chr11	11558667 11558685 -	0	0	0
LTR	MaLR	MTA_Mm	chrX	14650921 14650960 -	0	0	0
SINE	Alu	B1_Mm	chrX	14731590 14731605 -	0	0	0
Low_complexity	Low_complexity	A-rich	chrX	14757958 14757969 +	0	0	0
SINE	Alu	B1_Mus2	chr11	11574620 11574634 -	0	0	0
SINE	Alu	B1_Mus1	chrX	14854515 14854529 -	0	0.708814	0
SINE	Alu	B1_Mus2	chrX	14872220 14872234 +	0	0	0
SINE	Alu	B1_Mus1	chr11	11587730 11587744 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr11	11587744 11587746 +	0	0	0
SINE	Alu	B1_Mus1	chr11	11588010 11588025 -	0	0	0
SINE	Alu	B1_Mus1	chrX	14961957 14961971 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	14961972 14961974 +	0	0	0
SINE	Alu	B1_Mus1	chr10	30692962 30693116 -	0	0	0
SINE	Alu	B1_Mus1	chrX	14999472 14999487 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	14999487 14999489 +	0	0	0
SINE	Alu	B1_Mus1	chr11	11595726 11595740 -	0	0	0
LTR	MaLR	MTB	chrX	15023086 15023126 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	15125927 15125931 +	0	0	0
SINE	Alu	B1_Mus1	chrX	15220092 15220105 -	0	0	0.342623
SINE	Alu	B1_Mus1	chr11	11610218 11610232 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	11610232 11610236 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	15349261 15349267 +	0	0	0
SINE	Alu	B1_Mm	chrX	15449087 15449102 +	0	0	0
LINE	L1	L1Md_F2	chrX	15468543 15469046 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	15502188 15502192 +	0	0	0
SINE	Alu	B1_Mus1	chr11	11632991 11633004 -	0	0	0
LINE	L1	L1Md_F2	chrX	15659235 15659773 +	0	0	0
SINE	Alu	B1_Mm	chrX	15659860 15659875 -	0	0	0
SINE	Alu	B1_Mus1	chr11	11636532 11636547 -	0.591053	0	0
SINE	Alu	B1_Mm	chr11	11637809 11637824 -	0	0	0
LINE	L1	L1Md_T	chrX	15749425 15750000 -	0.332194	0	0
SINE	Alu	B1_Mm	chr11	11649491 11649506 +	0	0	0
LTR	MaLR	MTA_Mm	chr11	11649663 11649701 -	0	0	0
SINE	Alu	B1_Mus2	chr11	11649722 11649736 -	0	0	0
SINE	Alu	B1_Mus2	chrX	15865757 15865772 -	0	0	0
SINE	Alu	B1_Mus2	chrX	15926851 15926866 -	0	0	0
LTR	MaLR	MTA_Mm	chr11	11660440 11660479 +	0	0	0
LINE	L1	L1_Mus1	chrX	15951394 15951901 -	0	0	0
SINE	Alu	B1_Mm	chr11	11663996 11664010 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	11664010 11664013 +	0	0	0
SINE	Alu	B1_Mus1	chrX	15977820 15977834 -	0	0	0
scRNA	scRNA	BC1_Mm	chrX	16055790 16055802 -	0	0	0
SINE	Alu	B1_Mus2	chrX	16087286 16087300 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	16087301 16087303 +	0	0	0
SINE	Alu	B1_Mus1	chrX	16128536 16128551 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	16128551 16128554 +	0	0	0
SINE	Alu	B1_Mm	chr11	11677128 11677143 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	11677143 11677145 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	16189651 16189654 +	0	0	0
SINE	Alu	B1_Mm	chrX	16196589 16196604 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	16197557 16197561 +	0	0	0
SINE	Alu	B1_Mur3	chrX	16256984 16257002 +	0	0	0
SINE	Alu	B1_Mm	chr11	11692252 11692267 -	0	0	0
SINE	Alu	B1_Mur3	chr11	11702582 11702596 -	0	0	0
SINE	Alu	B1_Mus1	chrX	16589731 16589747 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrX	16589747 16589753 +	0	0	0
LINE	L1	L1_Mus2	chrX	16596766 16597288 +	0.332194	0	0
SINE	Alu	B1_Mus2	chr11	11744371 11744386 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	11744386 11744388 +	0	0	0
SINE	Alu	B1_Mus1	chr11	11751316 11751327 -	0	0	0
SINE	Alu	B1_Mm	chr11	11783899 11783912 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	11783912 11783918 +	0	0	0
SINE	Alu	B1_Mm	chr11	11824201 11824216 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	11824216 11824220 +	0	0	0
SINE	Alu	B1_Mus1	chr11	11832857 11832871 +	0	0	0
LTR	ERVK	RLTR16	chr11	11832872 11832915 +	0	0	0
LTR	MaLR	MTA_Mm	chr11	11833041 11833081 -	0	0	0
LTR	MaLR	MTB	chr11	11887292 11887331 -	0	0	0
SINE	B2	B2_Mm1a	chr11	12001506 12001525 -	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr11	12016920 12016927 +	0	0	0
SINE	Alu	B1_Mur4	chr10	5887709 5887856 -	0	0	0
SINE	Alu	B1_Mus2	chr11	12129228 12129243 -	0	0	0
LINE	L1	L1Md_A	chrY	805303 810787 -	0	0	0
LINE	L1	L1Md_A	chrY	885815 891300 -	0	0	0
LTR	MaLR	MTA_Mm	chrY	990967 991380 +	0	0	0

LINE	L1	L1Md_T	chrY	1102872	1109956	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrY	1253690	1253732	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chrY	1994334	1994367	+	0	0	0
SINE	Alu	B1_Mus1	chrY	2007789	2007927	-	0	0	0
LINE	L1	L1Md_A	chrY	2537057	2541534	+	0.664388	0.380568	0.342623
LTR	MaLR	MTA_Mm	chr12	3233234	3233623	-	1.18211	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr12	3347811	3347859	+	0	0	0
SINE	Alu	B1_Mus1	chr12	3511646	3511792	+	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	3511793	3511833	+	0	0	0
SINE	Alu	B1_Mm	chr12	3530796	3530942	-	0	0	0
SINE	Alu	B1_Mus1	chr12	3800718	3800862	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	3800863	3800890	+	0	0	0
LTR	MaLR	MTA_Mm	chr12	4549728	4550121	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	4561680	4561713	+	0.332194	0	0
SINE	Alu	B1_Mus1	chr12	5647165	5647299	+	0	1.01984	0.615324
Simple_repeat	Simple_repeat	(A)n	chr12	5647300	5647336	+	0	1.01984	0
SINE	Alu	B1_Mus2	chr12	5765325	5765470	-	0	0	0
SINE	Alu	B1_Mus2	chr12	6144701	6144847	-	0	0	0
SINE	Alu	B1_Mus1	chr12	6169239	6169386	+	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr12	6169387	6169413	+	0	0	0.342623
LTR	MaLR	MTA_Mm	chr12	6508038	6508429	-	0	0	0
LTR	MaLR	MTA_Mm	chr12	6567743	6568124	-	0	0	0
LTR	MaLR	MTA_Mm	chr12	6641419	6641811	+	0	0	0
LINE	L1	L1Md_F2	chr12	6650652	6655660	+	0	0	0
LTR	MaLR	MTA_Mm	chr12	6804138	6804531	+	0	0	0
SINE	Alu	B1_Mus2	chr12	7517420	7517567	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	8490422	8490457	+	0	0	0
LTR	MaLR	MTB	chr12	8529916	8530386	+	0	0	0
SINE	Alu	B1_Mm	chr12	8625459	8625604	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	8625605	8625629	+	0	0	0
LTR	MaLR	MTA_Mm	chr12	8635000	8635397	-	0.332194	0.380568	0.615324
SINE	Alu	B1_Mus1	chr12	8902711	8902857	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	8902858	8902887	+	0	0	0
SINE	Alu	B1_Mur1	chr12	8951826	8951972	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	8951973	8951997	+	0	0	0
LTR	MaLR	MTA_Mm	chr12	9088668	9089063	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	10023552	10023592	+	0	0	0
LTR	MaLR	MTA_Mm	chr12	10267196	10267593	+	0	0	0
SINE	Alu	B1_Mus2	chr12	10350818	10350963	+	0	0	0
SINE	Alu	B1_Mus1	chr12	10408818	10408965	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	11528674	11528700	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	12470827	12470854	+	0	0	0
LTR	MaLR	MTA_Mm	chr12	12673467	12673861	-	0	0	0
SINE	Alu	B1_Mm	chr12	12720986	12721108	-	0	0	0
SINE	Alu	B1_Mm	chr12	12831961	12832106	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	12832107	12832136	+	0	0	0
SINE	Alu	B1_Mm	chr12	12865689	12865835	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	12865836	12865860	+	0	0	0
SINE	Alu	B1_Mus1	chr12	12910170	12910324	-	0	0	0
LTR	MaLR	MTA_Mm	chr12	13069297	13069691	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	14153912	14153951	+	0.332194	0.708814	0.342623
SINE	Alu	B1_Mm	chr12	14591894	14592039	-	0	0	0
SINE	Alu	B1_Mus2	chr12	15124881	15125024	-	0	0	0
LINE	L1	L1Md_F2	chr12	15672344	15676981	-	0	0	0
SINE	Alu	B1_Mus2	chr12	17520134	17520279	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	17520280	17520317	+	1.05163	0	0
LTR	MaLR	MTA_Mm	chr10	33429352	33429747	-	0	0	0
SINE	Alu	B1_Mm	chr12	19054679	19054824	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	19054825	19054846	+	0	0	0
SINE	Alu	B1_Mus2	chr12	20833632	20833777	-	0	0	0
SINE	Alu	B1_Mus2	chr12	21392697	21392841	+	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	21392842	21392881	+	0	0	0.342623
SINE	Alu	B1_Mm	chr12	21523893	21524010	+	0	0	0
SINE	Alu	B1_Mm	chr12	22319833	22319978	-	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	23068473	23068502	+	0	0	0
SINE	Alu	B1_Mus2	chr12	23432127	23432275	-	0	0	0
SINE	Alu	B1_Mus2	chr12	23644845	23644991	+	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr12	23644992 23645031 +	0	0	0
SINE	Alu	B1_Mm	chr12	23782861 23783006 -	0	0	0
SINE	Alu	B1_Mus2	chr12	23868499 23868645 -	0	0	0
SINE	Alu	B1_Mm	chr12	24107656 24107801 -	0	0	0.342623
SINE	Alu	B1_Mus2	chr12	24597763 24597911 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	24597912 24597934 +	0	0	0
SINE	Alu	B1_Mus2	chr12	25316877 25317024 -	0	0	0
LTR	MaLR	MTA_Mm	chr12	25451960 25452354 -	0	0	0.615324
LTR	MaLR	MTA_Mm	chr12	26484494 26484879 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	34537130 34537163 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	34587040 34587080 +	0	0	0
LTR	MaLR	MTA_Mm	chr12	27633293 27633686 -	0	0	0
SINE	Alu	B1_Mm	chr10	34654164 34654310 -	0	0	0
LTR	MaLR	MTA_Mm	chr12	28309116 28309497 +	0	0	0
LTR	MaLR	MTA_Mm	chr10	34774847 34775222 -	0	0	0
LTR	MaLR	MTA_Mm	chr12	28997443 28997839 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	29943129 29943162 +	0	0	0
LTR	MaLR	MTA_Mm	chr12	30083705 30084100 -	0	0	0
LTR	MaLR	MTA_Mm	chr12	30085196 30085591 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	31283657 31283743 +	0	0	0
LTR	MaLR	MTA_Mm	chr12	31380953 31381350 -	0	0	0
SINE	Alu	B1_Mus2	chr10	35003976 35004122 -	0	0	0
SINE	Alu	B1_Mus2	chr12	32102993 32103139 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr12	32103140 32103163 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr12	33194645 33194703 +	0.591053	0	0
LTR	MaLR	MTA_Mm	chr12	33497715 33498114 -	0	0	0
LTR	MaLR	MTA_Mm	chr10	35317848 35318240 -	0	0	0
SINE	Alu	B1_Mm	chr12	34279377 34279527 -	0	0	0
LINE	L1	L1Md_F2	chr10	35332766 35339246 -	0	0	0
LTR	MaLR	MTA_Mm	chr12	34634155 34634547 +	0	0	0
LTR	MaLR	MTA_Mm	chr12	34706725 34707118 -	0	0	0
SINE	Alu	B1_Mm	chr10	6082720 6082866 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	6082867 6082936 +	0	0	0
SINE	Alu	B1_Mus1	chr12	35627536 35627683 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	35536181 35536232 +	0	0	0
SINE	Alu	B1_Mus2	chr10	35555519 35555664 -	0	0	0
SINE	Alu	B1_Mus2	chr12	37100994 37101137 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	37101138 37101166 +	0	0	0.342623
LTR	MaLR	MTA_Mm	chr12	37527071 37527465 -	0	0	0
SINE	Alu	B1_Mus1	chr12	37646652 37646799 -	0	0	0
SINE	Alu	B1_Mm	chr12	37663351 37663514 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	37663515 37663560 +	0	0	0
SINE	Alu	B1_Mus2	chr12	37697506 37697652 -	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr10	35681421 35681471 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	35826981 35827015 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	39483572 39483606 +	0	0	0
SINE	Alu	B1_Mm	chr12	40026568 40026713 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	40026714 40026738 +	0	0	0
LTR	MaLR	MTA_Mm	chr12	40059374 40059752 -	0	0	0
LTR	MaLR	MTA_Mm	chr12	40362042 40362437 -	0	0	0
SINE	Alu	B1_Mus1	chr10	6107133 6107269 -	0	0	0
LINE	L1	L1Md_T	chr12	44825237 44831659 -	0	0	0
SINE	Alu	B1_Mm	chr12	45321350 45321496 -	0	0	0
LTR	MaLR	MTA_Mm	chr12	45936339 45936737 +	0	0	0
LTR	MaLR	MTB	chr12	46203408 46203806 +	0	0	0
SINE	Alu	B1_Mus2	chr12	46561988 46562134 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	46562135 46562164 +	0	0	0
SINE	Alu	B1_Mus2	chr12	46743803 46743950 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	46743951 46743979 +	0	0	0
SINE	B2	B2_Mm1a	chr12	48387467 48387656 -	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	48577748 48577792 +	0	0	0
SINE	Alu	B1_Mus1	chr10	6126524 6126667 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	6126668 6126713 +	0	0	0
SINE	Alu	B1_Mm	chr12	48975926 48976057 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	49215230 49215267 +	0	0	0
SINE	Alu	B1_Mus2	chr12	49333765 49333892 -	0	0	0
SINE	Alu	B1_Mus1	chr10	36804181 36804324 -	0	0	0

LTR	MaLR	MTA_Mm	chr12	49591008 49591402 -	0	0	0
SINE	Alu	B1_Mm	chr10	36816790 36816917 -	0	0	0
LINE	L1	Lx	chr12	50017821 50022093 +	0	0	0
LTR	MaLR	MTA_Mm	chr12	50224067 50224435 +	0	0	0
LINE	L1	L1Md_F2	chr12	50536723 50543087 -	0	0	0
SINE	Alu	B1_Mus1	chr12	50900820 50900969 -	0	0	0
SINE	Alu	B1_Mus2	chr12	50920296 50920442 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	51011420 51011463 +	0	0	0
LTR	MaLR	MTA_Mm	chr12	51176537 51176932 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	52362320 52362385 +	0	0	0
LTR	MaLR	MTA_Mm	chr12	52621033 52621424 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	53180721 53180756 +	0	0	0.342623
SINE	Alu	B1_Mus1	chr12	53679436 53679598 +	0.923247	3.34916	2.18859
Simple_repeat	Simple_repeat	(A)n	chr12	53719814 53719843 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	53738902 53738950 +	0	0	0
LINE	L1	L1Md_F2	chr10	37226800 37231846 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	55392544 55392589 +	0	0	0
SINE	Alu	B1_Mus2	chr12	55406737 55406882 -	0	0	0
LTR	MaLR	MTA_Mm	chr12	55649987 55650381 -	0	0	0
SINE	Alu	B1_Mm	chr12	55682251 55682395 -	0	0	0.615324
SINE	Alu	B1_Mus2	chr12	55801494 55801640 -	0	0	0
LTR	MaLR	MTA_Mm	chr10	37378634 37379028 -	0	0	0
SINE	Alu	B1_Mus2	chr12	55900580 55900709 -	0	0.708814	0
SINE	Alu	B1_Mus2	chr12	55969479 55969625 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	55969626 55969728 +	0	0	0
SINE	Alu	B1_Mus1	chr12	55988233 55988403 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	55988404 55988436 +	0	0	0
SINE	Alu	B1_Mm	chr12	56219474 56219620 +	0.591053	0.380568	0.342623
Simple_repeat	Simple_repeat	(A)n	chr12	56219621 56219647 +	0	0.380568	0.342623
LTR	ERV1	RMER21A	chr12	56223649 56224525 +	0	0	0
LTR	ERV1	RMER21A	chr12	56373233 56373972 +	0	0	0
SINE	Alu	B1_Mm	chr12	57338326 57338472 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	57338473 57338499 +	0	0	0
LTR	MaLR	MTA_Mm	chr12	59393250 59393645 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	60020970 60020998 +	0	0	0
SINE	Alu	B1_Mur3	chr12	60058234 60058380 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	60058381 60058409 +	0	0	0
LINE	L1	L1Md_T	chr10	37977914 37984384 -	0	0	0
SINE	B4	ID_B1	chr12	60755680 60755897 -	0	0	0
SINE	Alu	B1_Mm	chr12	61790995 61791134 -	0	0	0
LTR	ERVK	RMER6C	chr12	61917368 61918147 -	0	0	0
SINE	Alu	B1_Mus1	chr12	63144502 63144647 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	63364314 63364377 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	63585768 63585805 +	0	0	0
SINE	Alu	B1_Mus2	chr12	64051846 64051991 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	64051992 64052028 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	64118647 64118687 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	38405400 38405427 +	0	0.380568	0
SINE	Alu	B1_Mus1	chr12	64488745 64488890 -	0	0	0
SINE	Alu	B1_Mus1	chr12	64924861 64924999 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	64925000 64925023 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	65108165 65108202 +	0	0	0
LTR	MaLR	MTA_Mm	chr12	65166688 65167082 +	0	0	0
SINE	Alu	B1_Mus2	chr12	65712491 65712640 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	65712641 65712669 +	0	0	0
SINE	Alu	B1_Mus2	chr12	66353088 66353224 -	0	0	0
LTR	MaLR	MTA_Mm	chr12	66607137 66607532 +	0	0	0
SINE	Alu	B1_Mm	chr12	67175772 67175918 -	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr12	67184073 67184112 +	0	0	0
SINE	Alu	B1_Mus1	chr12	67285574 67285711 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	67458147 67458183 +	0	0	0
SINE	Alu	B1_Mm	chr12	67545878 67545979 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	68433604 68433646 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr12	68499219 68499253 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	68612327 68612374 +	81.3778	38.4938	50.0266
SINE	Alu	B1_Mm	chr12	68756631 68756774 -	0.827951	0	0
SINE	Alu	B1_Mm	chr12	68868852 68868993 -	0	0	0

SINE	Alu	B1_Mm	chr12	69267045 69267191 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr12	69267192 69267270 +	0	0	0
SINE	Alu	B1_Mm	chr12	69362713 69362858 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	69560081 69560130 +	0	0	0
SINE	Alu	B1_Mus2	chr12	69687776 69687922 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	69687923 69687945 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	69706019 69706058 +	0	0	0
SINE	B2	B2_Mm2	chr12	70050666 70050838 +	0	0	0
SINE	Alu	B1_Mus2	chr12	70145998 70146128 -	0	0	0
SINE	Alu	B1_Mus1	chr12	70339733 70339879 +	0.591053	0	0
SINE	Alu	B1_Mus2	chr12	70564578 70564714 -	0	0	0
SINE	Alu	PB1D7	chr12	70671932 70672056 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	70672057 70672077 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	71888551 71888588 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	71901009 71901037 +	0	0	0
SINE	Alu	B1_Mus2	chr12	72075174 72075320 -	0	0	0.957947
Simple_repeat	Simple_repeat	(A)n	chr12	72398712 72398739 +	0	0	0
SINE	Alu	B1_Mus2	chr12	73709365 73709509 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	73709510 73709539 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	73722735 73722762 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	73730032 73730072 +	0	0	0
SINE	Alu	B1_Mus2	chr12	73749656 73749802 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	73749803 73749842 +	0	0	0
LTR	MaLR	MTB	chr12	73778348 73778758 +	0	0	0
SINE	Alu	B1_Mus2	chr12	73976684 73976830 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	73976831 73976865 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	74039857 74039905 +	0	0	0
SINE	Alu	B1_Mur4	chr12	74189359 74189496 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	74638590 74638614 +	0	0	0
SINE	Alu	B1_Mm	chr12	75060381 75060526 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	75060527 75060553 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	75681540 75681590 +	0.332194	0	0
LTR	MaLR	MTA_Mm	chr12	76309258 76309653 +	0	0	0
scRNA	scRNA	BC1_Mm	chr12	76796967 76797095 +	0	0	0
SINE	Alu	B1_Mus2	chr12	77439098 77439245 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	77439246 77439277 +	0	0	0
SINE	Alu	B1_Mus2	chr12	77444918 77445064 -	0	0	0.615324
SINE	Alu	B1_Mm	chr12	78074473 78074618 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	78074619 78074644 +	0	0	0
LTR	MaLR	ORR1B2	chr12	78187941 78188222 -	0	0	0
SINE	Alu	B1_Mus1	chr12	78188241 78188386 -	0	0	0
LTR	MaLR	MTA_Mm	chr12	78247223 78247614 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	78890202 78890266 +	0.332194	0	0
SINE	Alu	B1_Mus1	chr10	39903332 39903478 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	39903479 39903502 +	0	0	0
SINE	Alu	B1_Mus2	chr12	79268579 79268725 -	0	0	0
SINE	Alu	B1_Mus2	chr12	80984639 80984748 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	80984749 80984774 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	81032358 81032382 +	0	0	0
SINE	B2	B2_Mm2	chr12	81544171 81544357 -	0	0	0
LTR	MaLR	MTA_Mm	chr12	81730585 81730963 -	0	0	0
SINE	Alu	B1_Mus2	chr12	81829544 81829689 -	0	0	0
SINE	Alu	B1_Mus1	chr12	81987851 81987994 -	0	0	0
SINE	Alu	B1_Mus1	chr12	82658880 82659027 -	0	0	0
SINE	Alu	B1_Mm	chr12	82806992 82807120 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	40264457 40264539 +	0	0	0
SINE	Alu	B1_Mm	chr12	84838543 84838692 -	0	0	0
SINE	Alu	B1_Mus2	chr12	85289350 85289496 -	0	0	0
LTR	MaLR	MTA_Mm	chr12	85390451 85390846 -	0	0	0.342623
SINE	Alu	B1_Mus2	chr12	85572555 85572693 -	0	0	0
SINE	B2	B2_Mm2	chr12	85992682 85992855 -	0	0	0
SINE	Alu	B1_Mm	chr12	86336312 86336457 -	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr12	86621868 86621903 +	0.591053	0.708814	0
SINE	Alu	B1_Mus1	chr12	86718253 86718407 -	0	0	0.342623
SINE	Alu	B1_Mus2	chr12	87063532 87063678 -	0	0	0
SINE	Alu	B1_Mus2	chr12	87077606 87077751 -	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr12	87918789 87918830 +	0	0	0



SINE	Alu	B1_Mus1	chr12	88011031 88011178 -	0	0	0
SINE	Alu	B1_Mus2	chr12	88123916 88124056 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	88124057 88124085 +	0	0	0
SINE	Alu	B1_Mus1	chr12	88187774 88187901 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr12	88187902 88187946 +	0	0	0
SINE	Alu	B1_Mm	chr12	88307650 88307796 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	88307797 88307822 +	0	0	0
SINE	Alu	B1F1	chr12	88625419 88625556 -	0	0	0
SINE	Alu	B1_Mur4	chr12	88738426 88738571 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	88761967 88761996 +	0	0	0
SINE	Alu	B1_Mus1	chr12	91753335 91753482 -	0	0	0
SINE	Alu	B1_Mm	chr10	41186346 41186491 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	93116168 93116208 +	0	0	0
SINE	Alu	B1_Mus2	chr12	93509024 93509171 -	0	0	0
SINE	Alu	B1_Mm	chr12	93545359 93545505 -	0	0	0
Low_complexity	Low_complexity	A-rich	chr12	94504054 94504143 +	0	0	0
SINE	Alu	B1_Mus2	chr12	94656113 94656259 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	95045784 95045821 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	95238418 95238454 +	0	0	0
SINE	Alu	B1_Mus1	chr12	95416229 95416374 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	95466711 95466860 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	95482469 95482520 +	0	0	0
LTR	MaLR	MTA_Mm	chr12	95486837 95487231 -	0	0	0
SINE	Alu	B1_Mm	chr12	95803657 95803803 -	0	0	0
LINE	L1	L1Md_T	chr12	95855838 95861110 -	0	0	0
SINE	Alu	B1_Mus2	chr12	96312581 96312729 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	96572291 96572404 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	96705033 96705078 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	96921996 96922037 +	0	0	0
SINE	Alu	B1_Mm	chr12	97775062 97775198 -	0	0	0
SINE	Alu	B1_Mm	chr12	97999753 97999899 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	97999900 97999951 +	0	0	0
SINE	Alu	B1_Mus2	chr10	41800083 41800233 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	41800234 41800258 +	0	0	0
SINE	B2	B3	chr12	10041043 10041064 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	10041064 10041066 +	0	0	0
LTR	MaLR	MTA_Mm	chr12	10088902 10088941 +	0	0	0
SINE	B2	B2_Mm2	chr12	10131287 10131306 +	0	0	0.342623
SINE	Alu	B1_Mm	chr10	42004353 42004499 +	0	0	0
SINE	Alu	B1_Mm	chr12	10236374 10236389 +	0	0	0
SINE	Alu	PB1D10	chr12	10255283 10255295 -	0	0	0
SINE	Alu	B1_Mus1	chr12	10369535 10369550 -	0	0	0
SINE	Alu	B1_Mus2	chr12	10373900 10373914 -	0	0	0
SINE	Alu	B1_Mus2	chr12	10395966 10395980 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	10395980 10395983 +	0	0	0
LTR	MaLR	MTA_Mm	chr12	10473400 10473439 +	0	0	0
LTR	MaLR	MTB_Mm	chr12	10526007 10526047 -	0	0	0
LTR	MaLR	MTA_Mm	chr12	10557599 10557639 -	0	0	0
LTR	MaLR	MTA_Mm	chr10	6708535 6708928 +	0	0	0.342623
Low_complexity	Low_complexity	A-rich	chr10	42273881 42273978 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	10663251 10663254 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	10776668 10776672 +	0	0	0
LTR	MaLR	MTA_Mm	chr12	10854833 10854872 +	0	0	0
SINE	Alu	B1_Mus1	chr10	42469695 42469842 -	0	0	0
SINE	Alu	B1_Mus1	chr12	10996834 10996849 -	0	0	0
SINE	Alu	B1_Mus2	chr10	42573059 42573205 -	0	0	0
SINE	Alu	B1_Mus2	chr12	11036162 11036177 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	11036177 11036180 +	0	0	0
SINE	Alu	B1_Mus1	chr12	11070832 11070847 +	0	0	0
SINE	Alu	B1_Mm	chr12	11123549 11123564 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	42695503 42695544 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	11236798 11236803 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	11272222 11272226 +	0	0	0
SINE	Alu	B1_Mm	chr12	11275066 11275080 -	0	0	0
SINE	Alu	B1_Mus2	chr12	11289974 11289989 -	0	0	0
SINE	Alu	B1_Mus1	chr12	11293723 11293737 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	11293738 11293740 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr10	42932702 42932736 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr10	42934984 42935027 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	11514587 11514592 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	11516137 11516139 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	11980921 11980924 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	11985890 11985894 +	0	0	0
SINE	Alu	B1_Mm	chr10	43307937 43308085 +	0	0	0
SINE	Alu	B1_Mm	chr12	12031891 12031905 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr12	12050274 12050279 +	0	0.380568	0
scRNA	scRNA	4.5SRNA	chr12	12056363 12056372 -	0	0	0
LTR	MaLR	MTA_Mm	chr10	6875805 6876192 +	0	0	0
SINE	Alu	B1_Mus2	chr13	3074972 3075118 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	3075119 3075142 +	0	0	0
SINE	Alu	B1_Mm	chr13	3256411 3256547 -	0	0	0
SINE	Alu	B1_Mus2	chr13	3393746 3393892 +	0	0	0
LTR	MaLR	MTA_Mm	chr10	43421735 43422129 -	0	0	0
SINE	Alu	B1_Mus2	chr13	3629431 3629572 -	0	0	0
SINE	Alu	B1_Mus2	chr13	3631498 3631640 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	3631641 3631685 +	0	0	0
SINE	Alu	B1_Mus2	chr13	3872925 3873070 -	0	0	0
SINE	Alu	B1_Mm	chr13	4171004 4171150 -	0	0	0
SINE	Alu	B1_Mus2	chr13	4225773 4225919 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	4225920 4225946 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	4962817 4963210 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	43547026 43547085 +	0	0	0
LINE	L1	L1Md_T	chr13	5091424 5098088 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	43571875 43571910 +	0	0	0
SINE	Alu	B1_Mus2	chr13	5471973 5472086 -	0	0	0
LTR	MaLR	MTA_Mm	chr10	43572860 43573255 +	0	0	0
SINE	Alu	B1_Mus2	chr13	5724000 5724145 -	0	0	0
SINE	Alu	B1_Mus1	chr13	6168741 6168882 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	6168883 6168925 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	6210177 6210219 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	6536099 6536492 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	6763760 6763796 +	0	0	0
SINE	Alu	B1_Mus2	chr13	7142048 7142193 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	7142194 7142220 +	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	7338389 7338425 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	7338573 7338597 +	0	0	0
SINE	Alu	B1_Mm	chr13	7349497 7349642 +	0	0	0
Low_complexity	Low_complexity	GA-rich	chr13	7349643 7349780 +	0	0	0
SINE	Alu	B1_Mm	chr13	7581745 7581889 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	7581890 7581913 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	7852506 7852537 +	0	0	0
LTR	MaLR	MTA_Mm-int	chr13	7908604 7909614 -	0	0	0
LTR	MaLR	MTA_Mm	chr13	8066391 8066774 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	8116770 8116796 +	0	0	0
SINE	Alu	B1_Mm	chr10	43766869 43767014 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	8776222 8776256 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	8789329 8789723 +	0	0	0
SINE	Alu	B1_Mm	chr13	8925622 8925767 +	0	0	0
SINE	Alu	B1_Mus2	chr13	8996981 8997126 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	8997127 8997159 +	0	0	0
LTR	MaLR	MTB	chr13	9065833 9066225 -	0	0	0.615324
SINE	Alu	B1_Mm	chr13	9807401 9807541 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	9807542 9807571 +	0	0	0
LINE	L1	L1Md_T	chr13	9831267 9837911 -	0	0	0
SINE	Alu	B1_Mm	chr13	11241806 11241952 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	11241953 11241984 +	0	0	0
SINE	Alu	B1_Mm	chr13	12782709 12782855 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	12782856 12782878 +	0	0	0
SINE	Alu	B1_Mus2	chr10	44207316 44207481 -	0	0	0
LINE	L1	Lx	chr13	13145962 13152191 +	0	0	0.615324
SINE	Alu	B1_Mus1	chr13	13284392 13284535 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr13	13284536 13284614 +	0	0	0
SINE	Alu	B1_Mus2	chr13	13448003 13448154 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	13448155 13448183 +	0	0	0

SINE	Alu	B1_Mus2	chr13	13517876 13518022 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	13518023 13518047 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	13654197 13654590 -	0	0	0
SINE	Alu	B1_Mus1	chr13	14133886 14134033 -	0	0	0.615324
SINE	Alu	B1_Mm	chr13	14139606 14139752 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	14139753 14139783 +	0	0	0.342623
Other	Other	RMER1B	chr13	15249376 15250045 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	16072236 16072273 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	16095834 16095870 +	0	0	0
LINE	L1	L1Md_T	chr13	16477070 16483894 -	0.827951	0	0
LTR	ERV1	RLTR23	chr13	16579226 16579713 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	16872498 16872527 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	17907577 17907973 -	0	0	0
SINE	Alu	B1_Mus2	chr13	18193336 18193482 -	0.591053	0	0
SINE	Alu	PB1D10	chr13	18483657 18483755 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	18590276 18590324 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	18611277 18611671 -	0	0	0
SINE	Alu	B1_Mus2	chr10	44771632 44771778 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	44771779 44771817 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	18707432 18707468 +	0	0	0
SINE	Alu	B1_Mus1	chr13	19961324 19961469 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	20044640 20044704 +	0	0	0
SINE	Alu	B1_Mus1	chr10	44948957 44949101 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	22084589 22084618 +	0	0	0
SINE	Alu	B1_Mm	chr13	22124297 22124443 -	0	0	0
LTR	MaLR	MTA_Mm	chr13	23274139 23274534 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	23500658 23500694 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	23822094 23822148 +	0	0	0
SINE	Alu	B1_Mm	chr13	25642029 25642186 -	0	0	0
LTR	MaLR	MTA_Mm	chr13	25951168 25951561 -	0	0	0
LINE	L1	L1Md_T	chr13	26331199 263337871 -	0	0	0
LTR	MaLR	MTA_Mm	chr10	45467741 45468131 +	0	0	0
SINE	Alu	B1_Mus2	chr13	26859637 26859783 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	26859784 26859824 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	27349371 27349764 -	0	0	0
LINE	L1	Lx	chr13	27839811 27845872 +	0	0	0
SINE	Alu	B1_Mus1	chr13	27994561 27994707 -	0	0	0
LINE	L1	L1VL1	chr13	28194016 28199205 +	0	0	0
SINE	B2	B2_Mm2	chr13	28504805 28504986 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	28504987 28505021 +	0	0	0
LTR	MaLR	MTA_Mm	chr10	45615831 45616225 +	0	0	0
SINE	B2	B2_Mm2	chr10	45666579 45666765 +	0	0	0
Other	Other	RMER1B	chr10	45666766 45668066 +	0	0	0
SINE	Alu	B1_Mm	chr13	29033515 29033661 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	29033662 29033695 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	29172487 29172534 +	0	0	0
SINE	Alu	B1_Mus2	chr13	29955738 29955889 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	30187207 30187236 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	31481522 31481557 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	31596039 31596074 +	0	0	0
SINE	Alu	B1_Mus1	chr13	31599178 31599324 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	31599325 31599392 +	0	0	0
SINE	Alu	B1_Mm	chr13	33362437 33362580 +	0	0	0
SINE	Alu	B1_Mm	chr13	33437000 33437143 +	0	0	0
SINE	Alu	B1_Mm	chr13	33506913 33507056 +	0	0	0
SINE	Alu	B1_Mus2	chr13	33722760 33722911 -	0	0	0
LTR	MaLR	MTA_Mm	chr13	33919288 33919683 -	0	0	0
SINE	Alu	B1_Mus2	chr13	34447082 34447209 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	34447210 34447263 +	0	0	0
SINE	Alu	B1_Mm	chr10	7170108 7170254 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	7170255 7170288 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	35492480 35492876 -	0	0	0
LINE	L1	L1_Mus1	chr13	35681586 35687320 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	35738854 35738890 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	36720144 36720172 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	37214783 37214809 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	37401895 37401937 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr10	46582278 46582345 +	0	0	0
SINE	Alu	B1_Mur3	chr13	37540223 37540346 +	0	0	0
SINE	Alu	B1_Mm	chr13	37828491 37828628 +	0	0	0
SINE	B2	B2_Mm1t	chr13	37828645 37828805 +	0	0	0
SINE	Alu	B1_Mur3	chr13	37867562 37867710 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	37867711 37867733 +	0	0	0
SINE	Alu	B1_Mus2	chr13	38165439 38165586 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	38165587 38165616 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	38178197 38178227 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	39447365 39447418 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	39778342 39778370 +	0	0	0
LTR	MaLR	MTB	chr13	39893579 39893977 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	40041626 40041652 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	40436088 40436123 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	40627378 40627771 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	40771528 40771560 +	0	0	0
SINE	Alu	B1_Mus1	chr10	46989513 46989651 +	0	0	0
Low_complexity	Low_complexity	GA-rich	chr10	46989652 46989737 +	0	0	0
SINE	Alu	B1_Mus1	chr13	40899383 40899523 -	0	0	0
SINE	Alu	B1_Mus2	chr13	41073739 41073885 -	0	0	0
SINE	Alu	B1_Mus2	chr13	41238278 41238415 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	41238416 41238439 +	0	0	0
SINE	Alu	B1_Mus2	chr10	47076019 47076166 -	0	0	0
SINE	Alu	B1_Mus1	chr10	47087340 47087486 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	47105411 47105430 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	42427750 42428138 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	42487162 42487208 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	42754315 42754363 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	42762425 42762819 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	44157084 44157135 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	44227824 44227865 +	0.332194	0	0
SINE	Alu	B1_Mm	chr13	44467305 44467448 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	44579491 44579522 +	0	0	0
SINE	Alu	B1_Mm	chr13	44596213 44596361 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	44596362 44596446 +	0	0	0
SINE	Alu	B1_Mus2	chr13	44640990 44641135 -	0	0	0
LTR	MaLR	MTA_Mm	chr13	44669765 44670160 -	0	0	0
SINE	Alu	B1_Mus1	chr13	44710800 44710946 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	44746982 44747018 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	47511353 47511384 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	47517039 47517065 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	45115589 45115617 +	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr10	47672008 47672069 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	47778583 47778637 +	0	0	0
SINE	Alu	B1_Mus2	chr13	47561767 47561904 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	47993440 47993493 +	0	0.708814	0.957947
SINE	Alu	B1_Mm	chr13	48886106 48886252 +	0	0	0
SINE	Alu	B1_Mus2	chr13	50147655 50147779 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr13	50147780 50147871 +	0	0	0
SINE	Alu	B1_Mus1	chr13	50470512 50470658 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	50470659 50470723 +	0	0	0
SINE	Alu	B1_Mus1	chr13	51000123 51000273 -	0	0	0
SINE	Alu	B1_Mm	chr13	51162146 51162290 -	0	0	0
SINE	Alu	PB1D10	chr13	51252538 51252637 -	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	48380475 48380511 +	0	0	0
SINE	Alu	B1_Mm	chr10	48418109 48418243 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	48418244 48418266 +	0	0	0
SINE	Alu	B1_Mus1	chr13	52031096 52031242 -	0	0	0
Simple_repeat	Simple_repeat	(TTTA)n	chr13	52428439 52428474 +	0	0	0
SINE	Alu	B1_Mus2	chr13	52462170 52462301 -	0	0	0
LTR	MaLR	MTB	chr13	52589609 52590009 -	0	0	0
SINE	Alu	B1_Mus1	chr13	52640744 52640890 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	53078076 53078115 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	53105107 53105145 +	0	0	0
SINE	Alu	B1_Mus2	chr13	53172344 53172490 +	0	0	0
SINE	Alu	B1_Mm	chr13	53198455 53198600 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	53198601 53198632 +	0	0	0

SINE	Alu	B1_Mus2	chr13	54553896 54554042 -	0.923247	0.708814	1.30057
SINE	Alu	B1_Mm	chr13	54570805 54570951 +	0	0	0.615324
Simple_repeat	Simple_repeat	(A)n	chr13	54570952 54570997 +	0	0	0
SINE	Alu	B1_Mus2	chr13	54673779 54673925 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	54673926 54673961 +	0	0	0
LINE	L1	L1Md_F2	chr10	48785219 48790547 +	0	0	0
SINE	Alu	B1_Mm	chr13	54770762 54770909 +	0	0	0
SINE	Alu	B1_Mus2	chr13	54773316 54773460 -	0	0	0
SINE	Alu	B1_Mm	chr13	55274929 55275069 -	0	0	0
SINE	Alu	B1_Mus2	chr13	55277071 55277216 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	55277217 55277243 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	55282446 55282469 +	0	0	0.342623
SINE	Alu	B1_Mus1	chr13	55544392 55544533 -	0	0	0
SINE	Alu	B1_Mm	chr13	55678232 55678377 +	0	1.01984	0
Simple_repeat	Simple_repeat	(A)n	chr13	55678378 55678403 +	0	1.01984	0
SINE	Alu	PB1D10	chr13	56347080 56347195 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	56438131 56438172 +	0	0	0
SINE	Alu	B1_Mus2	chr13	56856130 56856276 -	0	0	0
SINE	Alu	B1_Mus2	chr13	57235429 57235547 -	0	0	0
LTR	MaLR	MTA_Mm	chr13	57461942 57462333 +	0	0.380568	0
SINE	B2	B2_Mm1t	chr13	58273325 58273516 +	0	0	0
SINE	Alu	B1_Mus1	chr13	58716863 58717008 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	58717009 58717041 +	0	0	0
SINE	Alu	B1_Mus1	chr13	59479206 59479353 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	59524902 59524935 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	59824610 59825005 -	0	0	0
LTR	ERV1	MER89	chr13	60623834 60624307 +	0	0	0
SINE	Alu	B1_Mus1	chr13	60691714 60691861 -	0	0	0
SINE	Alu	B1_Mus1	chr13	61564169 61564313 -	0	0	0
LINE	L1	L1Md_F2	chr13	61617493 61622559 +	0	0	0
SINE	Alu	B1_Mus2	chr13	61815325 61815471 -	0	0	0
SINE	Alu	B1_Mm	chr13	61850194 61850339 -	0	0	0
SINE	Alu	B1_Mus2	chr13	61981422 61981567 +	0	0	0
SINE	Alu	B1_Mus2	chr13	62073027 62073173 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	62073174 62073211 +	0	0	0
SINE	Alu	B1_Mus1	chr13	62096694 62096831 -	0	0	0
SINE	Alu	B1_Mus2	chr13	62196835 62196981 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	62196982 62197013 +	0	0	0
SINE	Alu	B1_Mm	chr10	49734284 49734427 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	49734428 49734474 +	0.591053	0	0
SINE	Alu	B1_Mus2	chr13	62593966 62594112 +	0	0	0
SINE	Alu	B1_Mm	chr13	62615786 62615931 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	62615932 62615983 +	0	0	0
SINE	Alu	B1_Mm	chr13	62737425 62737571 -	0	0	1.23065
SINE	Alu	B1_Mm	chr13	62808719 62808864 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	62808865 62808889 +	0	0	0
SINE	Alu	B1_Mus2	chr13	62835842 62835988 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	62835989 62836010 +	0	0	0
SINE	Alu	B1_Mus2	chr13	63030794 63030939 -	0	0	0
SINE	Alu	B1_Mus2	chr13	64389268 64389429 -	0	0	0
LTR	MaLR	MTA_Mm	chr13	64832090 64832485 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	50060516 50060550 +	0.591053	0	0
SINE	Alu	B1_Mus2	chr13	65221347 65221485 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	65221486 65221509 +	0	0	0.615324
LTR	MaLR	MTA_Mm	chr13	65563228 65563622 -	0	0	0
SINE	Alu	B1_Mm	chr13	65720724 65720869 -	0	0	0
SINE	Alu	B1_Mus2	chr10	50160768 50160914 +	0	0	0
SINE	Alu	B1_Mus2	chr13	66087117 66087263 +	0	0	0.615324
LTR	MaLR	MTA_Mm	chr13	66581009 66581403 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	66712376 66712769 -	0	0	0
SINE	Alu	B1_Mus1	chr13	67655089 67655235 +	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chr13	67655236 67655313 +	0	0	0
SINE	Alu	B1_Mus2	chr13	67715461 67715607 -	0	0	0
LTR	MaLR	MTB	chr13	69359070 69359466 -	0	0	0
SINE	B2	B2_Mm2	chr13	69623586 69623773 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	69943316 69943714 +	0	0	0
LTR	MaLR	MTB_Mm	chr13	70141061 70141441 -	0.332194	0	0

SINE	Alu	B1_Mus1	chr13	70708555 70708702 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	70708703 70708746 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	71123828 71124222 +	0	0	0
SINE	Alu	B1_Mm	chr13	71171973 71172119 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	71172120 71172143 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	71413475 71413503 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	71635564 71635592 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	71806124 71806160 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	50747265 50747300 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	72431764 72431804 +	0	0	0
SINE	Alu	B1_Mm	chr13	72592812 72592947 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	72592950 72592976 +	0	0.380568	0
LTR	MaLR	MTA_Mm	chr13	72914149 72914544 -	0	0	0
SINE	Alu	B1_Mus2	chr13	73053550 73053695 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	73168501 73168537 +	0	0	0
SINE	Alu	B1_Mus2	chr13	73561803 73561949 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	74037466 74037504 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	74773048 74773442 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	75002148 75002180 +	0.332194	0	0
SINE	B2	B2_Mm1a	chr13	75498002 75498193 +	0	0	0
LINE	L1	L1_Mus4	chr13	75628623 75634697 +	0	0	0
SINE	Alu	B1_Mm	chr10	51196566 51196700 -	0	0	0
SINE	Alu	B1_Mm	chr13	76335597 76335743 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	76335744 76335768 +	0	0	0
LINE	L1	Lx2	chr13	78377855 78383139 -	0	0	0
LINE	L1	L1Md_A	chr13	78437960 78442838 +	0	0	0
LINE	L1	L1_Mus2	chr13	78558013 78563950 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	78611711 78611760 +	0	0	0
SINE	Alu	B1_Mur2	chr13	79398939 79399079 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	79631496 79631538 +	0	0.380568	0
SINE	Alu	B1_Mm	chr13	79693013 79693159 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	79693160 79693183 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	79922006 79922032 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	79995709 79996103 +	0	0	0
Simple_repeat	Simple_repeat	(CAAAAA)n	chr13	80532706 80532810 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	82293486 82293877 +	0	0	0
SINE	B2	B2_Mm1t	chr13	83055656 83055847 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	83055848 83055882 +	0.664388	0.380568	0
SINE	Alu	B1_Mm	chr13	83070981 83071127 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	83372778 83372806 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	83518060 83518104 +	0	0	0
LTR	ERVK	RMER19B	chr13	83840303 83841215 -	0.591053	0	0
LINE	L1	L1Md_T	chr13	84623034 84630285 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	85232746 85232780 +	0	0	0
SINE	Alu	B1_Mus1	chr13	85289023 85289167 -	0	0	0
SINE	Alu	B1_Mur4	chr13	85718064 85718210 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	85718211 85718238 +	0	0	0
SINE	Alu	B1_Mus2	chr13	85892462 85892600 -	0	0	0
SINE	Alu	B1_Mm	chr13	86103579 86103724 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	86103725 86103770 +	0	0	0
SINE	Alu	B1_Mus2	chr13	86313869 86314013 -	0	0	0
LTR	MaLR	ORR1B2-int	chr13	86445516 86446832 +	0	0	0
SINE	Alu	B1_Mm	chr13	86905562 86905708 -	0	0	0
SINE	Alu	B1_Mm	chr13	87628756 87628896 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	87628897 87628934 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	87896976 87897022 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	87998283 87998677 -	0	0	0
SINE	Alu	B1_Mus2	chr13	88224379 88224525 +	0	0	0
SINE	Alu	B1_Mus2	chr13	88351362 88351508 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	88351509 88351589 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	88363431 88363457 +	0	0	0
SINE	Alu	B1_Mm	chr13	88431079 88431225 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	88784776 88785170 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	52250094 52250134 +	0.664388	0	1.02787
Simple_repeat	Simple_repeat	(A)n	chr13	90868424 90868458 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	90940306 90940704 -	0	0	0
SINE	Alu	B1_Mus1	chr13	91535293 91535442 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr13	91535443	91535469 +	0	0	0
SINE	B2	B2_Mm2	chr13	91934904	91935096 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	92904032	92904072 +	0	0	0
SINE	B2	B2_Mm1a	chr13	93179717	93179908 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	93179909	93179942 +	0.664388	0	0.342623
SINE	Alu	B1_Mus2	chr13	94007260	94007405 -	0	0	0
SINE	Alu	B1_Mus2	chr13	94320870	94321015 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr13	94321016	94321049 +	0	0	0
SINE	Alu	B1_Mus1	chr13	94364208	94364354 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	94364355	94364390 +	0	0.380568	0
SINE	Alu	B1_Mus1	chr13	95111699	95111844 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	95111845	95111883 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	96686650	96686688 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	96976470	96976865 -	0	0	0
LTR	MaLR	MTA_Mm	chr13	97479721	97480106 -	0	0	0
SINE	Alu	B1_Mm	chr13	97520114	97520251 -	0	0	0
LTR	MaLR	MTA_Mm	chr13	97553052	97553433 -	0	0	0
SINE	Alu	B1_Mm	chr13	98144829	98144974 -	0	0	0
SINE	Alu	B1_Mus2	chr13	99046571	99046717 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	99046718	99046740 +	0	0	0
LTR	MaLR	MTA_Mm	chr10	53174693	53175069 +	0	0	0
SINE	Alu	B1_Mus1	chr13	99116714	99116869 +	0	0	0
SINE	Alu	B1_Mus1	chr10	53178343	53178488 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	53178489	53178518 +	0	0	0
SINE	Alu	B1_Mus2	chr13	99598967	99599114 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	99599115	99599143 +	0	0	0
SINE	Alu	B1_Mus2	chr13	99699179	99699325 -	0	0	0
SINE	Alu	B1_Mm	chr13	99750249	99750394 +	0	0	0
SINE	Alu	B1_Mus2	chr13	10116095	10116109 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	10123716	10123719 +	0	0	0
SINE	Alu	B1_Mus2	chr13	10161799	10161813 -	0	0	0
LTR	MaLR	MTA_Mm	chr13	10163368	10163407 -	0	0	0.342623
LINE	L1	L1Md_F2	chr10	53549275	53555455 -	0	0	0
LTR	MaLR	MTA_Mm	chr13	10211032	10211071 +	0	0	0
SINE	Alu	B1_Mus1	chr13	10423011	10423021 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	10423021	10423025 +	0	0	0
SINE	Alu	B1_Mm	chr13	10423176	10423191 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	10423191	10423196 +	0	0	0
SINE	Alu	B1_Mus2	chr13	10480008	10480023 -	0	0	0
SINE	Alu	B1_Mus1	chr10	7822651	7822780 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	10608937	10608942 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	10651802	10651805 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	10727185	10727224 -	0	0	0
LTR	MaLR	MTA_Mm	chr13	10729853	10729892 -	0	0	0
SINE	Alu	B1_Mm	chr13	10787496	10787510 -	0	0	0
SINE	Alu	B1_Mus1	chr13	10827701	10827716 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	10827716	10827723 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	10829685	10829690 +	0	0	0
LINE	L1	L1Md_T	chr13	10929342	10930007 -	0	0	0
SINE	Alu	PB1D7	chr10	54360832	54360953 -	0	0.708814	0
SINE	Alu	B1_Mus2	chr10	54470339	54470487 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	11109186	11109190 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	11132979	11132983 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr10	54558813	54558842 +	0	0	0
LTR	MaLR	MTB	chr13	11160828	11160868 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	11215978	11215980 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	11219905	11219945 +	0	0	0
SINE	Alu	B1_Mm	chr13	11228450	11228464 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	11228464	11228467 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	11230415	11230454 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	11230564	11230604 +	1.18211	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	11231660	11231666 +	0	0	0
SINE	Alu	B1_Mm	chr13	11232909	11232924 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr13	11232924	11232926 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr13	11252844	11252858 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr13	11252858	11252860 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	11284881	11284885 +	0	0	0

Simple_repeat	Simple_repeat	(A)n	chr13	11289059 11289062 +	0	0	0
LTR	MaLR	MTB	chr13	11307347 11307387 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	11321759 11321764 +	0	0.380568	0
LTR	MaLR	MTA_Mm	chr13	11414840 11414880 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	11423443 11423483 +	0	0	0
LTR	MaLR	MTA_Mm	chr13	11430184 11430223 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	11457805 11457807 +	0	0	0
SINE	B2	B3	chr13	11523869 11523888 -	0	0	0
LTR	MaLR	MTB	chr13	11526604 11526644 +	0	0	0
LTR	MaLR	MTA_Mm	chr10	55147670 55148064 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr13	11679166 11679170 +	0.591053	0.380568	0.957947
LTR	MaLR	MTA_Mm	chr13	11759153 11759193 -	0	0	0
LTR	MaLR	MTA_Mm	chr13	11925603 11925642 -	0	0	0
LINE	L1	L1Md_F2	chr13	11978540 11979067 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	3019467 3019492 +	0	0	0
SINE	Alu	B1_Mus1	chr14	4271863 4272009 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	5223751 5223776 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	5360784 5360819 +	0	0	0.342623
SINE	Alu	B1_Mus1	chr14	5402442 5402589 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr14	5402590 5402639 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	6408820 6408845 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	6545857 6545890 +	0	0	0.342623
SINE	Alu	B1_Mus1	chr14	6587705 6587852 +	0	0	0
Low_complexity	Low_complexity	A-rich	chr14	6587853 6587903 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr14	7165543 7165613 +	0	0	0
SINE	B2	B2_Mm1a	chr14	8516825 8517021 +	0	0	0
SINE	Alu	B1_Mm	chr10	56003776 56003921 +	0	0	0
SINE	Alu	B1_Mus2	chr14	8594154 8594300 -	0	0	0
LTR	MaLR	MTA_Mm	chr14	8599962 8600343 -	0	0	0
LTR	MaLR	MTA_Mm	chr14	8646969 8647364 +	0	0	0
SINE	Alu	B1_Mm	chr14	9021558 9021703 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	9021704 9021758 +	0	0	0
LTR	MaLR	MTA_Mm	chr14	9850607 9851000 +	0	0	0
LTR	MaLR	MTA_Mm	chr14	9852100 9852493 +	0	0	0.342623
SINE	Alu	B1_Mus2	chr10	56113910 56114056 -	0	0	0
SINE	Alu	B1_Mus2	chr10	56145755 56145908 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	56145909 56145936 +	0	0	0
SINE	Alu	B1_Mur2	chr10	56193647 56193784 -	0	0	0
SINE	Alu	B1_Mm	chr14	12125289 12125435 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	12125436 12125461 +	0	0	0
SINE	Alu	B1_Mm	chr10	56267522 56267668 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	56267669 56267694 +	0	0	0
LTR	MaLR	MTA_Mm	chr14	13143360 13143751 -	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	56300529 56300585 +	0	0	0
LINE	L1	L1Md_F2	chr10	56338075 56344228 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	14485581 14485607 +	0	0	0
SINE	Alu	B1_Mus1	chr14	14732433 14732576 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	14732577 14732609 +	0	0	0
SINE	Alu	B1_Mm	chr14	14827144 14827289 -	0	0	0
LINE	L1	L1_Mus1	chr14	15087175 15091630 -	0	0.380568	0
LINE	L1	L1_Mus2	chr14	15188554 15193433 +	0	0	0
LINE	L1	L1_Mus2	chr14	15417219 15423484 -	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr14	15508711 15508739 +	0	0	0.615324
LTR	MaLR	MTA_Mm	chr14	15647598 15647992 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	15897940 15897986 +	0	0	0
SINE	Alu	B1_Mus1	chr14	16030748 16030895 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	16030896 16030943 +	0	0	0
LINE	L1	L1_Mus3	chr14	16108848 16113632 +	0	0	0
LINE	L1	L1_Mus1	chr14	17183355 17187733 -	0	0	0
LTR	MaLR	MTA_Mm	chr14	17969965 17970351 -	0.591053	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr14	18221184 18221221 +	0	0	0
LTR	MaLR	MTA_Mm	chr14	18415648 18416040 -	0.591053	0	0.615324
LTR	MaLR	MTA_Mm	chr14	18486297 18486696 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr10	56759332 56759366 +	0.591053	0	0
SINE	Alu	B1_Mus2	chr14	19097662 19097800 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	19097801 19097848 +	0	0	0
SINE	Alu	B1_Mm	chr14	20023156 20023292 +	0	0	0



Simple_repeat	Simple_repeat	(A)n	chr14	20023293 20023320 +	0	0	0
LTR	MaLR	MTA_Mm	chr14	20053001 20053395 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	20305130 20305184 +	0	0	0
SINE	Alu	B1_Mus1	chr14	20685450 20685603 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	20685604 20685627 +	0	0	0
SINE	Alu	B1_Mus1	chr14	20844841 20844962 +	0	0	0
Simple_repeat	Simple_repeat	(GAAA)n	chr14	20844963 20845013 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	20931794 20931824 +	0	0	0
SINE	Alu	B1_Mm	chr14	20955724 20955860 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	20955861 20955883 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	21300380 21300409 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	21450957 21451005 +	0	0	0
SINE	Alu	B1_Mus1	chr14	21688914 21689060 -	0.591053	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	21713466 21713492 +	0	0	0
SINE	B2	B2_Mm1a	chr14	21715851 21716043 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	21716044 21716069 +	1.38382	0.708814	0.685246
LTR	MaLR	MTA_Mm	chr14	23995720 23996113 +	0	0	0
SINE	Alu	B1_Mm	chr14	26952216 26952361 -	0	0	0
SINE	Alu	B1_Mus2	chr14	26952950 26953096 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	26953097 26953120 +	0	0.380568	0
SINE	Alu	B1_Mm	chr14	27091865 27092010 -	0	0	0
SINE	Alu	B1_Mus2	chr14	27092599 27092745 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	27092746 27092769 +	0	0	0
SINE	Alu	B1_Mus1	chr10	57774628 57774775 -	0	0	0
SINE	Alu	B1_Mus2	chr14	28223983 28224129 +	0	0.708814	0
Simple_repeat	Simple_repeat	(A)n	chr14	28224130 28224158 +	0	0.708814	0
LTR	MaLR	MTA_Mm	chr14	28363619 28364014 -	0.591053	0	0
SINE	Alu	B1_Mus2	chr14	28401938 28402082 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	29610337 29610441 +	0	0	0
SINE	Alu	B1_Mus2	chr14	30745642 30745783 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	30745784 30745828 +	0	0	0
SINE	Alu	B1_Mus1	chr14	30775571 30775716 -	0	0	0
SINE	Alu	B1_Mus2	chr14	31936139 31936284 -	0	0	0
LTR	MaLR	MTB	chr10	58181127 58181525 +	0	0	0
LTR	MaLR	MTA_Mm	chr14	32143313 32143708 -	0.332194	0	0
LTR	MaLR	MTA_Mm	chr14	32398327 32398725 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	32496883 32496912 +	0	0	0
SINE	Alu	B1_Mus1	chr14	32648480 32648623 -	0	0	0
SINE	Alu	B1_Mus2	chr14	32935988 32936134 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	32936135 32936157 +	0	0	0
SINE	Alu	B1_Mm	chr14	33555429 33555576 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	33555577 33555670 +	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	34184460 34184495 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	34296510 34296538 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr14	34563721 34563759 +	0	0	0
LTR	MaLR	MTB	chr14	34671328 34671725 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	34765411 34765445 +	0	0	0
SINE	Alu	B1_Mus2	chr14	35484377 35484526 +	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	35484527 35484570 +	0	0	0
LINE	L1	L1_Mus2	chr14	37040240 37040608 -	0	0	0
LINE	L1	L1Md_F2	chr14	37052070 37058253 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	37100296 37100324 +	0	0	0
SINE	Alu	B1_Mus2	chr14	37389033 37389179 -	0	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	38074507 38074555 +	0	0	0
LTR	MaLR	MTA_Mm	chr14	38271599 38271994 -	0.332194	0	0
Simple_repeat	Simple_repeat	(A)n	chr14	38300956 38300999 +	0	0	0.342623
Simple_repeat	Simple_repeat	(A)n	chr14	38323203 38323240 +	0	0	0
SINE	Alu	B1_Mus2	chr14	38938544 38938659 -	0	0	0
LINE	L1	L1Md_F2	chr14	40529522 40535385 -	0	0	0
SINE	Alu	B1_Mus2	chr14	40783900 40784045 -	0	0	0
SINE	Alu	B1_Mm	chr14	41186299 41186450 +	0	0	0
SINE	Alu	B1_Mus2	chr14	42329045 42329190 -	0	0	0

Supplementary Table S4: Predicted repeat targets based on piRNA cleavage signature in addition to their class; family; subfamily; genomic coordinates and normalized CAGE expression throughout postnatal development in brain