

4	1.3E+08 .	T	G	base_qual	SNV	243	8	251	0.032	205	8	213	0.038	link	-	Elf3l	c.529-2A>-	ENSMUST splice_acceptor_variant+intron_variant
5	88642927 .	A	C	base_qual	SNV	242	2	244	0.008	227	3	230	0.013	link	-	Rufy3	c.1488-2A:-	ENSMUST splice_acceptor_variant+intron_variant
5	1.45E+08 .	T	G	base_qual	SNV	188	10	198	0.051	224	10	234	0.043	link	-	Trrap	c.261+2T>-	ENSMUST splice_donor_variant+intron_variant
5	1.47E+08 .	A	C	base_qual	SNV	241	12	253	0.047	210	24	234	0.103	link	-	Cdx2	c.538+2T>-	ENSMUST splice_donor_variant+intron_variant
6	69992764 .	T	G	base_qual	SNV	174	3	177	0.017	179	7	186	0.038	link	-	Igkv18-36	c.50-5A>C-	ENSMUST splice_region_variant+intron_variant
6	1.47E+08 .	T	G	base_qual	SNV	193	5	198	0.025	181	8	189	0.042	link	-	Asun	c.504-8A>-	ENSMUST splice_region_variant+intron_variant
8	84225814 .	A	C	base_qual	SNV	127	6	133	0.045	114	3	117	0.026	link	-	Zswim4	c.1516+2T-	ENSMUST splice_donor_variant+intron_variant
9	20350524 .	G	A	base_qual	SNV	173	9	182	0.049	126	13	139	0.094	link	-	Zfp560	c.322+8C>-	ENSMUST splice_region_variant+intron_variant
9	37646292 .	A	C	base_qual	SNV	144	11	155	0.071	146	11	157	0.07	link	-	Siae	c.1399-5A:-	ENSMUST splice_region_variant+intron_variant
9	67031450 .	A	C	base_qual	SNV	148	2	150	0.013	173	3	176	0.017	link	-	Tpm1	c.828+2T>-	ENSMUST splice_donor_variant+intron_variant
X	37157086 .	A	C	base_qual	SNV	71	2	73	0.027	79	9	88	0.102	link	-	Akap14	c.1464+2T-	ENSMUST splice_donor_variant+intron_variant
X	57275980 .	T	G	base_qual	SNV	93	1	94	0.011	94	4	98	0.041	link	-	Arhgef6	c.897-7A>-	ENSMUST splice_region_variant+intron_variant
X	1.15E+08 .	A	C	base_qual	SNV	90	8	98	0.082	91	3	94	0.032	link	-	Klhl4	c.1547-2A:-	ENSMUST splice_acceptor_variant+intron_variant
X	1.58E+08 .	C	T	PASS	SNV	133	0	133	0	130	9	139	0.065	link	-	Cnksr2	c.1091+1G-	ENSMUST splice_donor_variant+intron_variant