Skippy Reference: variants that create ectopic splice sites (hg18)

SNP No.	Chr	Position	Gene	Potential Coding effect	Splice site activation	Variant (DNA)	max(Δss) Reference	
1	3	15651990	**BTD**	Missense	3’	G->A	7.96	(1)
2	4	155708309	**FGB**	Missense	3’	T->A	8.37	(2)
3	6	31432182	**HLA-B**	Missense	3’	G->C	9.95	(3)
4	11	107511124	**ACAT1**	Missense	3’	C->T	2.11	(4)
5	15	41282727	**EPB42**	Missense	3’	G->T	13.31	(5)
6	19	17860206	**SLC5A5**	Missense	3’	C->G	8.06	(6)
7	X	32238252	**DMD**	Nonsense	3’	T->G	8.59	(7)
8	11	67567654	**TCIRG1**	Synonymous	3’	G->A	7.95	(8)
9	19	11084983	**LDLR**	Synonymous	3’	C->A	8.04	(9)
10	X	153838609	**F8**	Synonymous	3’	G->T	2.51	(10)
11	1	154375026	**LMNA**	Missense	5’	G->A	1.53	(11)
12	4	89215079	**PKD2**	Nonsense	5’	C->T	0	(12)
13	5	70980772	**MCCB**	Missense	5’	A->G	8.18	(13)
14	7	91702781	**KRIT1**	Missense	5’	C->G	1.81	(14)
15	7	91703738	**KRIT1**	Missense	5’	A->G	8.18	(14)
16	10	89682830	**PTEN**	Missense	5’	C->G	8.27	(15)
17	11	107708775	**ATM**	Missense	5’	C->T	7.75	(16)
18	16	67413594	**CDH1**	Missense	5’	C->T	7.75	(17)
19	17	26551666	**NF1**	Missense	5’	C->T	7.75	(18)
20	17	26580523	**NF1**	Missense	5’	G->A	3.81	(19)
21	17	26565668	**NF1**	Nonsense	5’	A->G	8.18	(19)
22	17	26583297	**NF1**	Missense	5’	T->A	5.33	(18)
23	17	38512021	**BRCA1**	Missense	5’	T->G	8.6	(20)
24	17	45619024	**COL1A1**	Missense	5’	A->G	8.18	(21)
25	X	100516810	**BTK**	Nonsense	5’	A->G	8.18	(22)
26	X	133436951	**HPRT1**	Missense	5’	G->T	7.65	(23)
27	X	135560190	**CD40L**	Nonsense	5’	G->T	4.23	(24)
28	X	148392770	**IDS**	Missense	5’	C->G	0	(25)
29	X	153235823	**FLNA**	Missense	5’	C->T	0	(26)
30	1	154375028	**LMNA**	Synonymous	5’	C->T	0.49	(27)
31	1	153474560	**GBA**	Synonymous	5’	A->G	8.18	(28)
32	1	156912581	**SPTA1**	Synonymous	5’	A->G	8.18	(29)
33	2	219382723	**CYP27A1**	Synonymous	5’	G->T	7.65	(30)
34	3	48593334	**COL7A1**	Synonymous	5’	C->T	7.75	(31)
35	3	143757971	**ATR**	Synonymous	5’	A->G	0	(32)
36	3	185449258	**ALG3**	Synonymous	5’	C->T	0.75	(33)
37	5	42747170	**GHR**	Synonymous	5’	C->T	7.76	(34)
38	5	42735837	**GHR**	Synonymous	5’	A->G	4.8	(35)
39	7	65070189	**GUSB**	Synonymous	5’	C->T	7.76	(36)
40	11	5204749	**HBB**	Synonymous	5’	G->A	2.24	(37)
41	11	5204753	**HBB**	Synonymous	5’	T->A	2.28	(38)
42	12	108497085	**MVK**	Synonymous	5’	C->T	7.75	(39)
43	12	46535733	**VDR**	Synonymous	5’	C->G	3.5	(40)
44	15	70428488	**HEXA**	Synonymous	5’	T->A	4.52	(41)
45	16	162980	**HBA2**	Synonymous	5’	C->T	7.76	(42)
46	16	66027759	**HSD11B2**	Synonymous	5’	C->G	2.51	(43)
47	17	42731797	**ITGB3**	Synonymous	5’	C->T	7.76	(44)
48	17	26580468	**NF1**	Synonymous	5’	G->A	3.51	(19)
49	17	26586877	**NF1**	Synonymous	5’	C->T	7.75	(19)
50	19	17808957	**JAK3**	Synonymous	5’	C->T	7.75	(45)
51	19	806755	**ELA2**	Synonymous	5’	C->A	7.03	(46)
52	X	100500293	**BTK**	Synonymous	5’	A->G	0	(47)
53	X	153301140	**TAZ**	Synonymous	5’	G->A	0	(48)
54	X	148376419	**IDS**	Synonymous	5’	C->T	7.75	(49)

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