

D.melanogaster					
Donor Splice Sites			Acceptor Splice Sites		
pos	oligomer	contribution	pos	oligomer	contribution
5	G	153.87	-3	G	-85.09
6	T	80.34	-7	G	-72.54
0	G	75.28	-5	G	-63.88
2	C	-69.41	-5	T	62.94
2	T	69.41	-8	G	-61.83
3	A	67.06	-10	G	-58.30
5	GT	169.42	-6	TT	90.15
4	AG	121.48	-3	GA	-87.99
2	TA	99.97	-9	AG	-84.01
0	GG	77.85	-8	AG	-82.13
6	TA	77.13	-10	AG	-71.38
-1	AG	73.79	-11	AG	-68.25
4	AGT	99.41	-3	GAG	-74.21
0	GGT	93.74	-3	CAG	49.56
1	GTA	87.53	-4	AGA	-46.62
5	GTA	80.54	-3	TAG	41.48
2	TAA	69.22	-7	ATT	39.96
3	AAG	66.85	-10	CAG	-39.93
0	GGTA	67.87	-4	AGAG	-38.08
2	TAAG	57.86	-4	ACAG	31.55
1	GTAA	57.84	-6	TTTC	29.80
2	TGAG	54.04	-58	GTAA	29.56
-1	AGGT	53.20	-25	CTAA	29.22
3	GAGT	51.95	-21	CTAA	28.89
2	TGAGT	46.96	-57	GTAAG	25.16
1	GTAAG	45.85	-6	TTTCA	24.17
1	GTGAG	43.43	-56	GTAAG	24.16
2	TAAGT	34.42	-58	GTAAG	23.40
0	GGTAA	34.23	-6	TTGCA	23.22
0	GGTGA	24.52	-59	GTAAG	21.30
1	GTGAGT	34.55	-6	TTTCAG	17.18
1	GTAAGT	25.40	-6	TTGCAG	16.50
0	GGTGAG	25.36	-57	GTAAGT	13.38
2	TGAGTA	21.24	-56	GTAAGT	11.82
0	GGTAAG	19.22	-59	GTGAGT	11.72
0	GGTAGG	13.14	-59	GTAAGT	11.66
1	GTGAGTA	13.37	-6	TTTCAGA	7.03
0	GGTGAGT	12.95	-6	TTGCAGA	6.37
-1	AGGTGAG	10.51	-7	ATTTTCAG	5.74
-1	AGGTGGG	7.80	-6	TTACAGA	5.42
0	CGTAAGT	7.52	-6	TTTCAGG	5.19
1	GTGAGTT	7.41	-7	CTTGCAG	5.16
-1	AGGTGGGT	3.08	-7	ATTTTCAGA	2.00
-1	AGGTGAGA	2.69	-7	CTTGCAGA	1.63
-2	CAGGTGAG	2.51	-6	TTTCAGAT	1.51
0	AGTGAGTA	2.45	-7	TTCGCAGA	1.49
1	GTGAGTAT	2.41	-12	TTTTTTTTT	-1.37
0	TGTGAGTA	2.32	-7	TTTCCAGG	1.30