

D.rerio					
Donor Splice Sites			Acceptor Splice Sites		
pos	oligomer	contribution	pos	oligomer	contribution
5	G	140.63	-3	G	-105.25
0	G	87.69	-11	T	90.23
2	T	78.05	-12	T	86.63
2	C	-78.05	-7	G	-85.46
3	A	70.16	-9	T	85.29
-1	A	69.50	-10	T	84.95
5	GT	147.00	-8	AG	-143.14
4	AG	122.57	-9	AG	-135.94
2	TA	111.33	-10	AG	-126.06
0	GG	90.69	-11	AG	-122.94
-1	AG	87.13	-7	AG	-110.51
1	GT	80.71	-12	AG	-110.14
0	GGT	110.50	-3	GAG	-91.80
1	GTA	98.23	-9	CAG	-77.43
4	AGT	90.91	-10	CAG	-75.07
-1	AGG	73.64	-8	CAG	-70.07
3	GAG	61.86	-11	CAG	-64.96
5	GTA	61.77	-12	CAG	-64.50
0	GGTA	85.86	-4	AGAG	-50.04
-1	AGGT	65.08	-5	CAGA	-37.10
2	TGAG	56.06	-10	TCAG	-36.29
1	GTAA	49.67	-4	GGAG	-35.81
2	TAAG	47.13	-3	CAGG	35.08
3	GAGT	43.38	-11	TCAG	-34.63
1	GTGAG	44.92	-5	CAGAG	-28.76
1	GTAAG	37.02	-4	AGAGG	-21.30
2	TGAGT	35.33	-6	TTTCA	19.88
0	GGTAA	33.05	-5	TAGAG	-19.35
3	ATCCT	31.99	-10	TGTGT	18.89
4	TCCTT	30.78	-3	TAGGA	18.68
3	ATCCTT	25.72	-6	TTTCAG	14.12
1	GTGAGT	25.66	-3	TAGGAG	12.16
2	TATCCT	23.77	-6	TCAGAG	-11.90
0	GGTGAG	22.57	-5	CAGAGG	-11.87
0	GGTCAG	17.80	-5	AGCAGG	-11.29
0	CGTAAG	17.55	-10	TGTGTT	9.82
-1	AGGTAAG	-18.55	-5	CGCAGGT	5.58
2	TATCCTT	15.99	-6	CAGCAGG	-5.40
1	GTATCCT	14.44	-7	CAGGCAG	-5.24
0	GGTAAGT	-13.03	-6	TTTCAGA	5.20
-1	TGGTGAG	12.16	-7	TTCACAG	5.18
0	CGTAAGT	12.03	-5	AGCAGGT	-4.77
-1	AGGTAAGT	-8.14	-9	TGTGTGCA	1.77
-1	AGGTGAGT	-7.96	-6	TTTCAGAG	1.72
1	GTATCCTT	6.62	-8	TGTTTCAG	1.72
-2	CAGGTAAG	-5.94	-5	TTTAGGTT	-1.68
0	TGTATCCT	4.68	-6	CAGCAGGT	-1.60
-2	AAGGTAAG	-4.52	-8	CTTCACAG	1.44