

A.thaliana					
Donor Splice Sites			Acceptor Splice Sites		
pos	oligomer	contribution	pos	oligomer	contribution
5	G	151.56	-3	G	-116.54
0	G	110.63	-3	C	99.41
3	A	96.96	-5	T	77.45
2	T	93.08	-8	T	75.91
2	C	-93.08	-9	T	74.65
-1	A	75.91	-12	T	74.05
5	GT	156.03	-9	AG	-132.66
2	TA	137.40	-8	AG	-128.56
4	AG	130.96	-3	GA	-120.51
-1	AG	119.31	-10	AG	-120.33
0	GG	114.40	-11	AG	-114.00
1	GT	96.25	-3	CA	102.80
0	GGT	135.92	-3	GAG	-101.65
1	GTA	119.54	-3	CAG	86.71
-1	AGG	104.42	-10	CAG	-76.63
4	AGT	97.90	-9	CAG	-69.93
2	TAA	75.79	-11	CAG	-67.34
-2	CAG	62.07	-10	TAG	-62.96
0	GGTA	93.28	-3	TAGG	51.43
-1	AGGT	92.86	-4	AGAG	-50.37
1	GTAA	62.50	-3	CAGG	49.98
-2	CAGG	51.34	-4	GCAG	43.11
2	TGAG	47.51	-26	CTAA	41.59
2	TAAG	42.76	-4	ACAG	41.49
-2	CAGGT	38.23	-5	TGCAG	26.50
1	GTGAG	37.66	-4	AGAGG	-24.45
-1	AGGTT	37.06	-4	ATAGG	23.57
1	GTAAG	33.49	-5	TGTAG	21.80
-2	AAGGT	30.45	-6	TTTCA	20.65
0	GGTAA	29.85	-26	CTAAC	19.93
0	TGTAAG	25.16	-6	TTTCAG	14.68
-2	CAGGTT	24.36	-5	TGTAGG	11.68
1	GTGAGT	21.61	-6	TGGCAG	11.12
0	GGTAAG	-20.27	-6	AGGCAG	-10.37
-1	AGGTTT	19.27	-5	TATAGG	9.64
0	GGTAAA	17.02	-5	TGCAGT	9.51
-1	AGGTAAG	-19.54	-6	TTTCAGG	7.08
0	GGTAAGT	-12.72	-7	CAGGCAG	-4.62
-2	CAGGTTT	11.43	-7	TTGGCAG	4.53
0	TGTAAGT	10.27	-6	TGGCAGG	4.52
0	TGTGAGT	9.79	-7	TTTTTCAG	4.30
-1	AGGTGAG	-9.72	-5	TGCAGTG	4.22
-1	AGGTAAGT	-5.73	-6	TTTCAGGG	1.57
-2	AAGGTAAG	-4.62	-3	TAGGGTTT	-1.50
-1	AGGTGAGT	-4.56	-7	ATTCAGG	1.45
-2	CAGGTAAG	-3.69	-8	TGTAACAG	1.33
1	GTATCCTT	3.15	-7	TTTTTCAGG	1.32
-1	AGGTAAGC	-2.96	-8	TTTGGCAG	1.31