

Predictor	Normal	Alternative	Normal mutated	Alternative mutated
	CTT/gtaagt	taa/gtaagc	CTT/gtaaag	aaa/gtaagc
Splice site score calculation (Max = 12.6)¹	6.9	6.2	2	7.6
Analyzer splice tool (Max = 100)²	78.94	78.17	61.92	81.99
Splice site prediction by neuronal network (Max = 1) [29]	0.95	0	0	0.97
MaxEntScan (Max = 12) [30]	8	5.66	-3	7.31
Human splicing finder (Max = 100) [31]	84.38	85.27	70.28	86.29

1. http://rulai.cshl.edu/new_alt_exon_db2/HTML/score.html

2. <http://ibis.tau.ac.il/ssat/SpliceSiteFrame.htm>