



Supplementary Figure 8 - Splice site and polypyrimidine tract scores of each branchpoint motif model. (A) MaxentScan score distribution of 3'ss and 5'ss for each motif model. P-values determined using t-test. (B) Polypyrimidine tract score distribution for each motif model. Polypyrimidine tract scores determined using algorithm from (Hum Mol Genet. 2002 Feb 15;11(4):451-64). P-values determined using wilcoxon test.