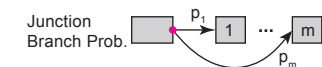
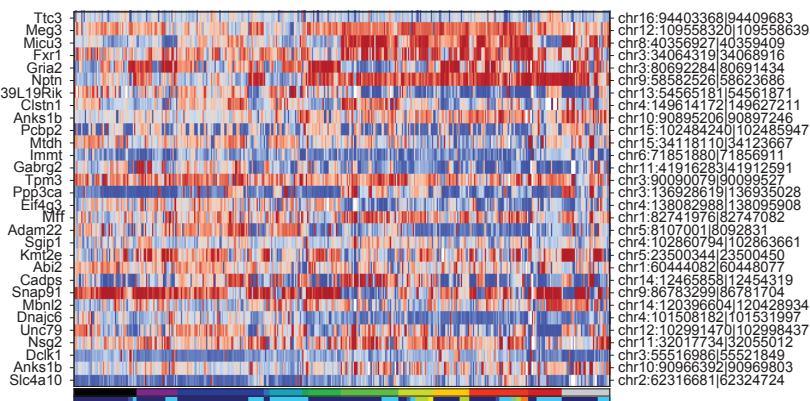
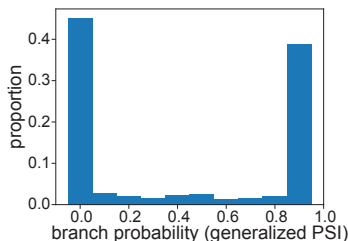


A Branch probabilities

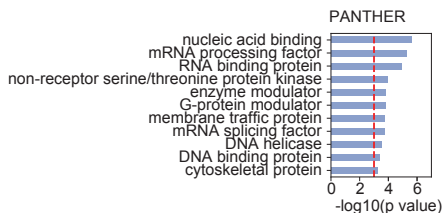
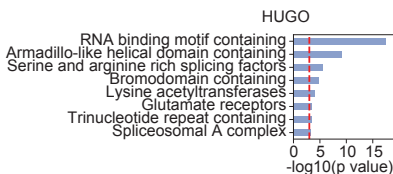
B Genes with top 30 splice DEF



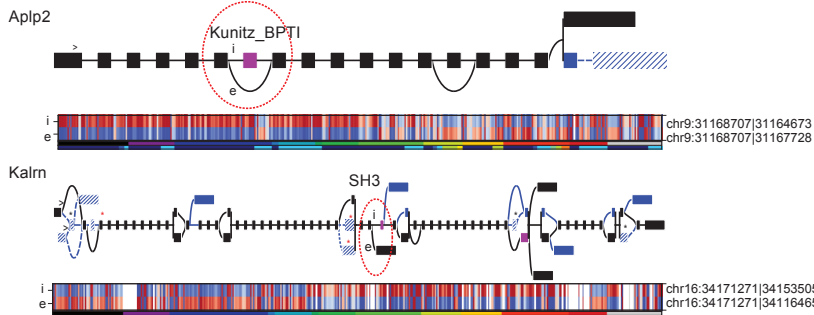
Distribution of branch probabilities



C Enriched families in top 1000 splice DEF



D Examples of alternative splicing leading to inclusion/exclusion of Pfam motifs



E Proportion of alternative sites changing Pfam or NMD

