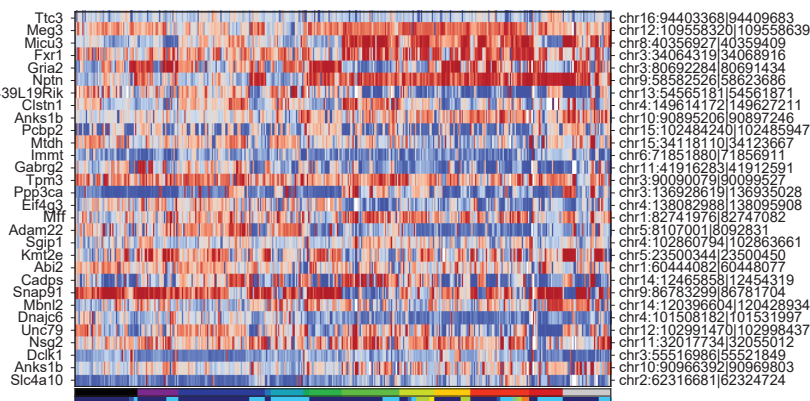
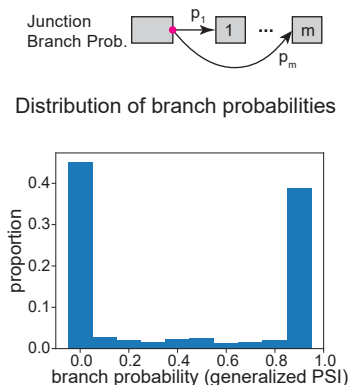
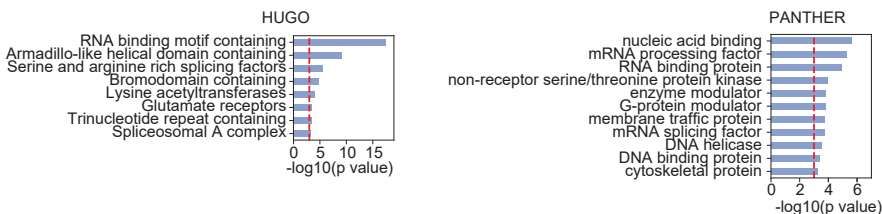


A Branch probabilities

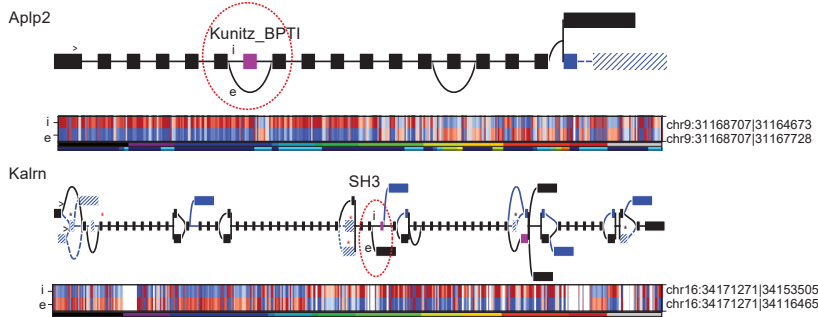
B Branch probabilities with top 30 splice DEF



C Enriched families splice DEF top 1000



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



E Proportion of alternative sites w/ Pfam or NMD choice

