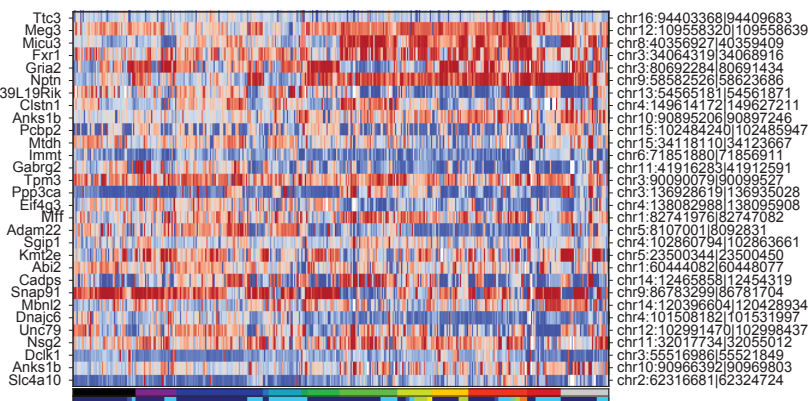
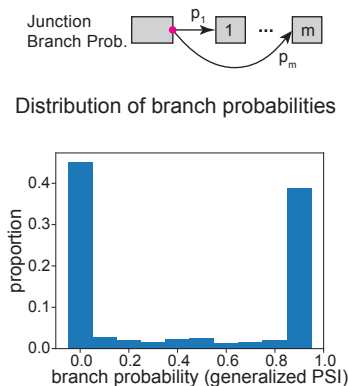
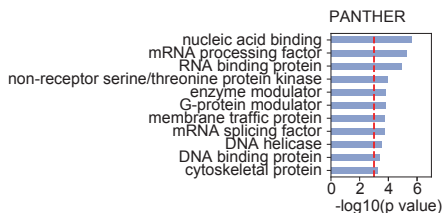
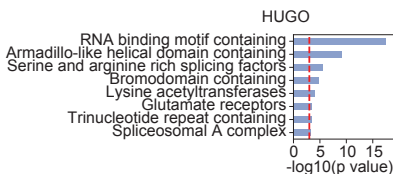


A Branch probabilities

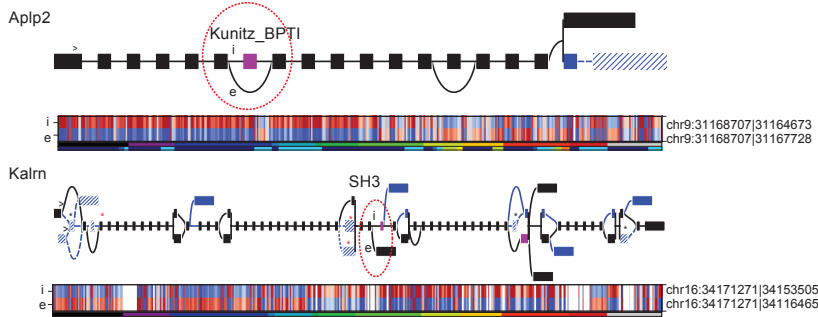
B Example sites with high differential splicing (top 30)



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

