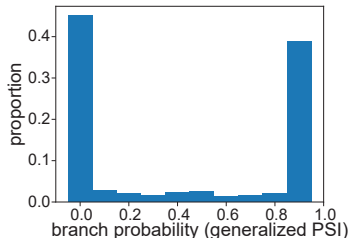
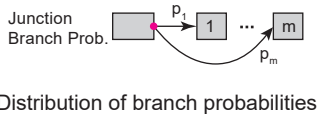


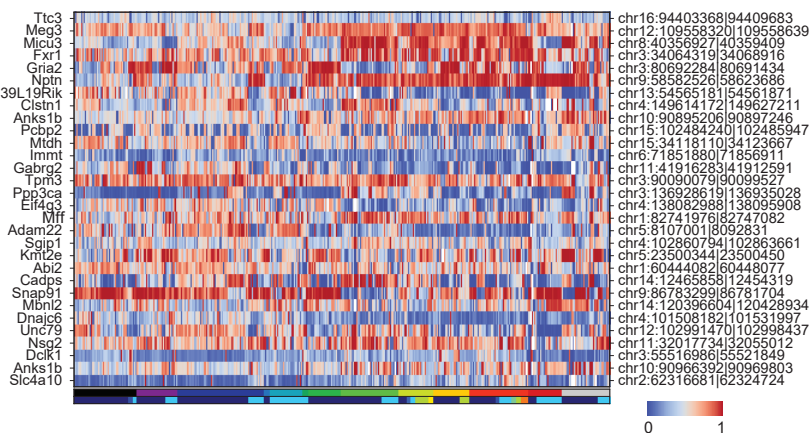
A

Branch probabilities



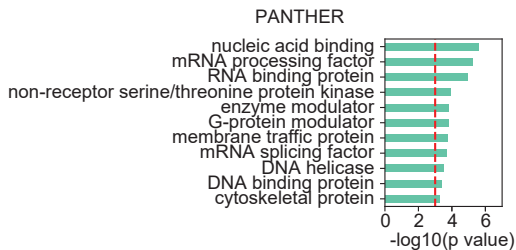
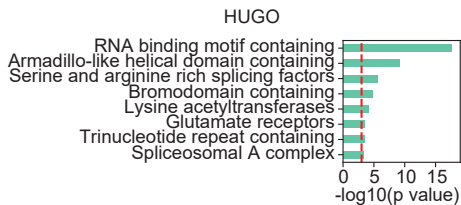
B

Example sites with high differential splicing (top 30)



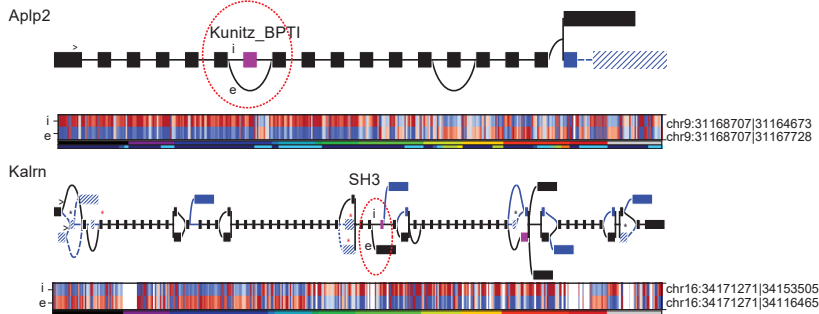
C

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

