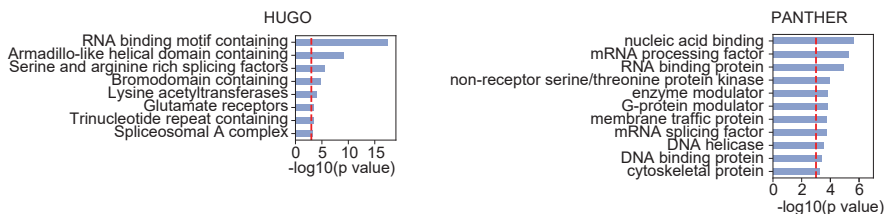
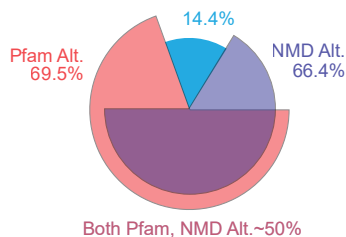


**C** Enriched families splice DEF top 1000



**D** Proportion of alternative sites w/ Pfam or NMD choice



**E** Examples of alternative splicing leading to inclusion/exclusion of Pfam motifs

