



Supplementary Figure S11: Analysis of alternative C-terminal protein sequences in the GENCODE annotated mouse proteome.

(A) Frequencies of primarily splice-driven or frameshift-driven mechanisms underlying C-terminal protein variation. (B) Distribution of different splice-driven altered C-terminal categories and change in the altered region length normalized by the APPRIS-ref isoform length and the change in reference-relative lengths of their altered C-terminal regions. (C) Distribution of the frequencies for various splice-driven patterns. (D) Distribution of the normalized altered region lengths for various splice-driven patterns (shown in (C)).