



Figure S4: 3' end processing factors affect the splicing of both terminal and not terminal introns.

Splicing sensitive microarrays of 3' end processing factors *ppn1* (A) and *ssu72* (B). Only transcripts that contained multiple introns were considered for this analysis. Introns were called 'terminal' when they were the last intron annotated within a transcript, the remainder were considered 'not terminal'. Each heatmap was organized by hierarchical clustering. Global splicing defects are observed for each of the mutants, regardless of relative intron location.