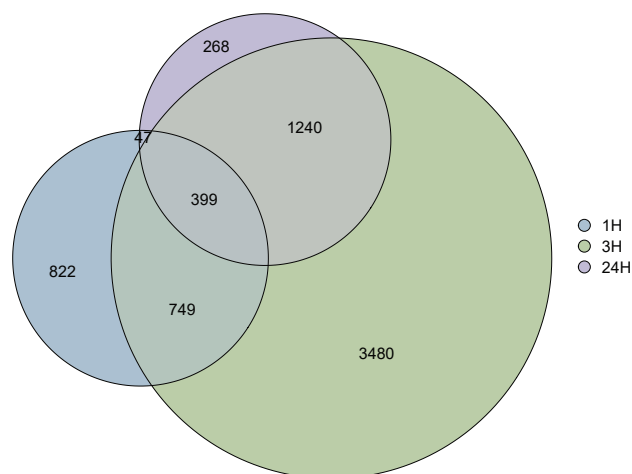


Supplemental Figure 2

A

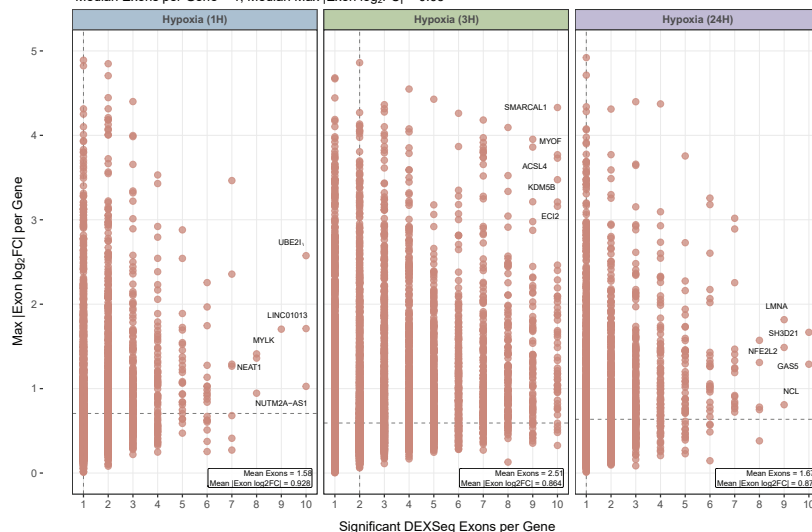
DEXSeq: Gene Overlap Across Hypoxia Timepoints



B

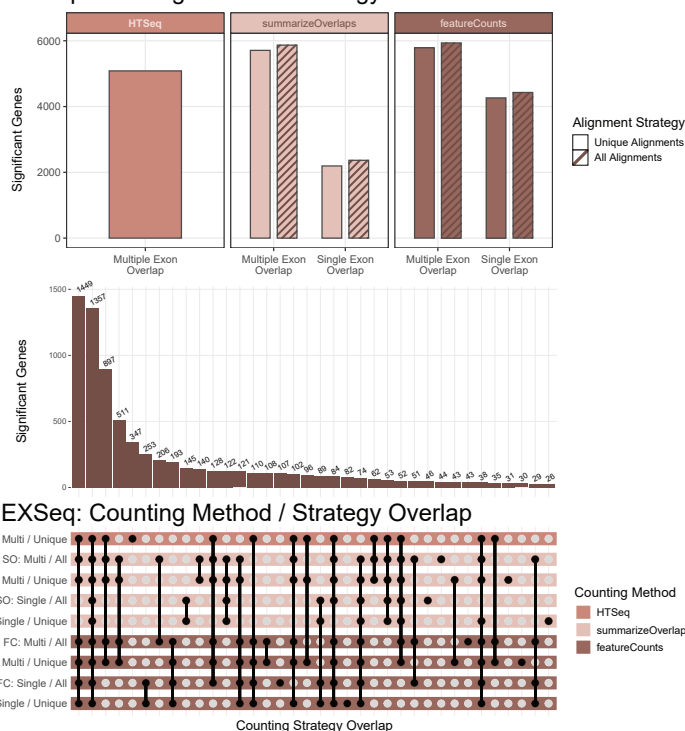
DEXSeq: Multiplicity and Magnitude

Median Exons per Gene = 1; Median Max |Exon log₂FC| = 0.63



C

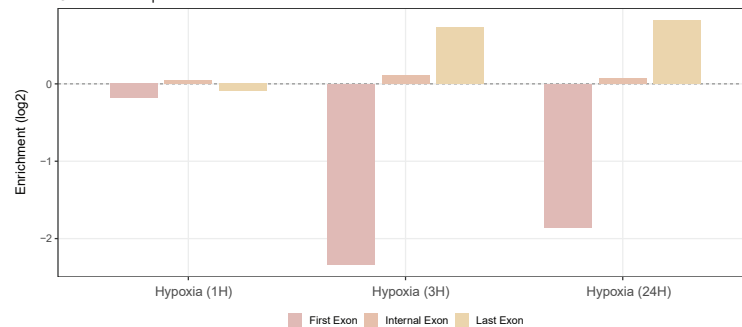
DEXSeq: Counting Method / Strategy Counts



D

DEXSeq: Exon Position Enrichment

Observed exon positions relative to the distribution of all tested exon bins



DEXSeq: Exon Direction Pattern Enrichment

Observed exon direction patterns relative to expectation based on global exon-level direction proportions for genes with >= 2 significant exon bins

