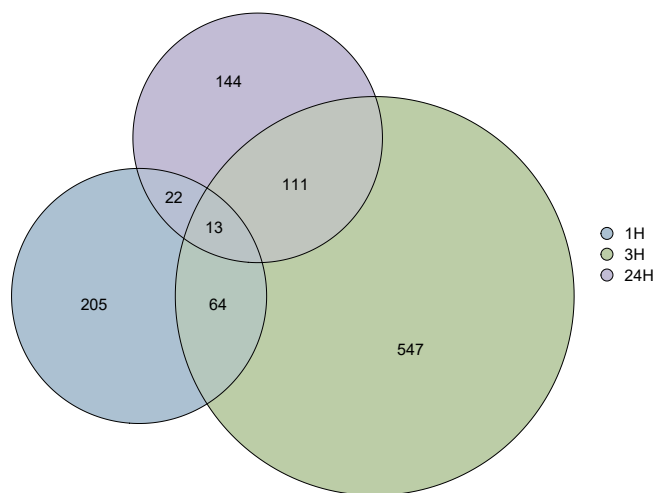


Supplemental Figure 3

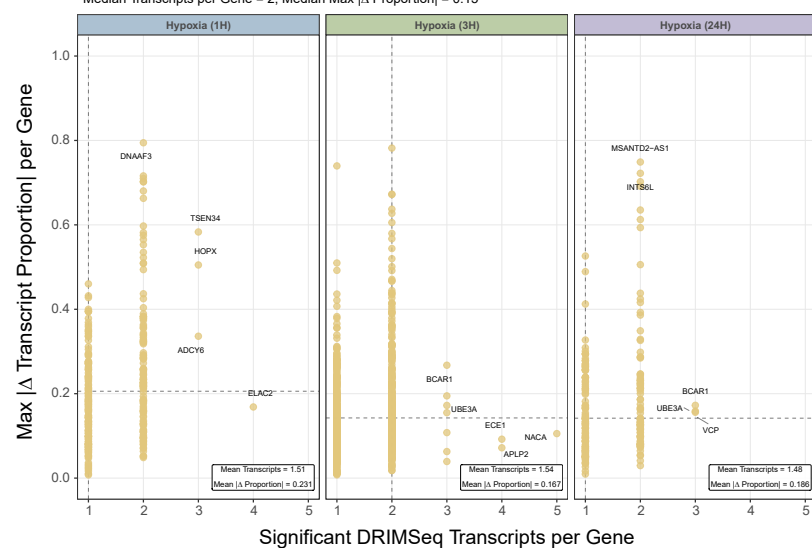
A

DRIMSeq: Gene Overlap Across Hypoxia Timepoints

**B**

DRIMSeq: Transcript Multiplicity and Magnitude

Median Transcripts per Gene = 2; Median Max $|\Delta|$ Proportion = 0.15

**C**

DRIMSeq: Transcript Usage Trajectories

Top 5 DTU genes per timepoint, selected by $|\Delta|$ Proportion

