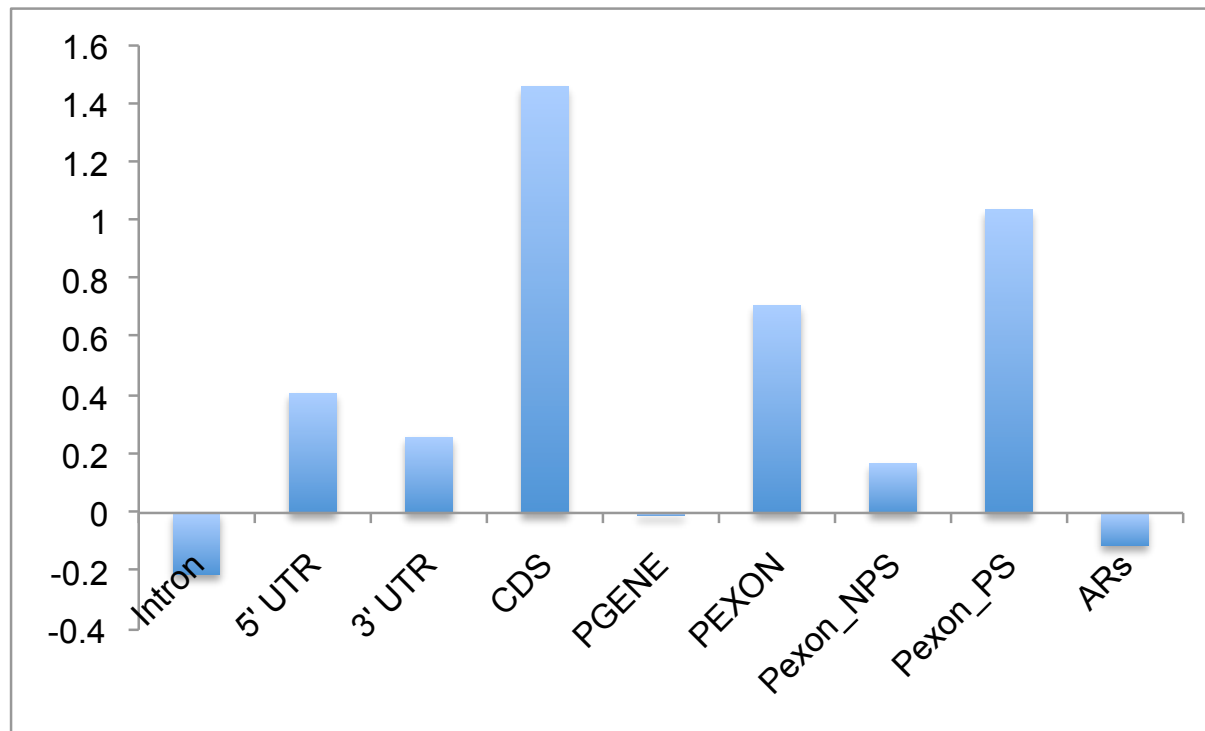
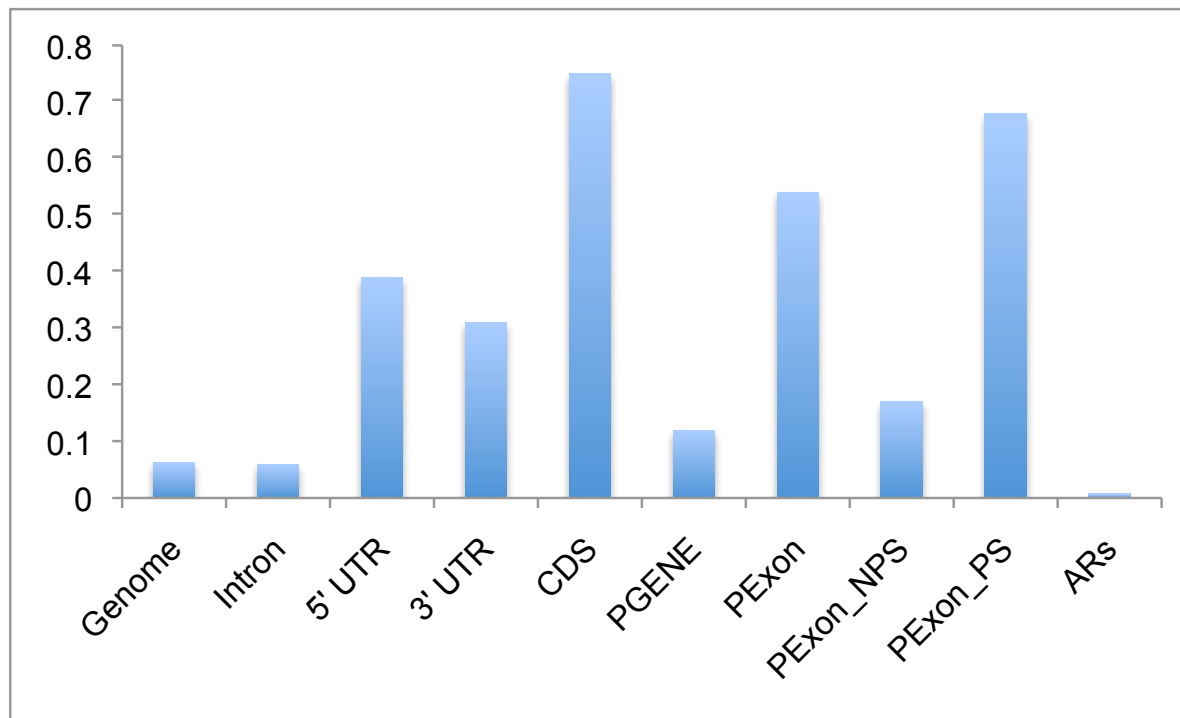


Average RS Scores in Different Regions

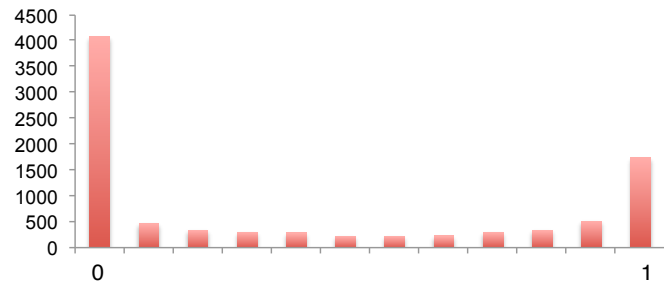


Percentage of Genomic Region Under Constraint



Distribution of Constraint Region Percentage for Pseudo-exons

Exon distributions of non-processed Pgenes



Exon distribution of processed Pgenes

