

GeneMark Prediction Report

General Statistics

Gene count: 14836

Single-exon genes: 2370

Multi-exon genes: 12466

Introns per gene: 2.92

Introns per multi-exon gene: 3.47

Genes fully supported by external evidence: 5730 (38.62%)

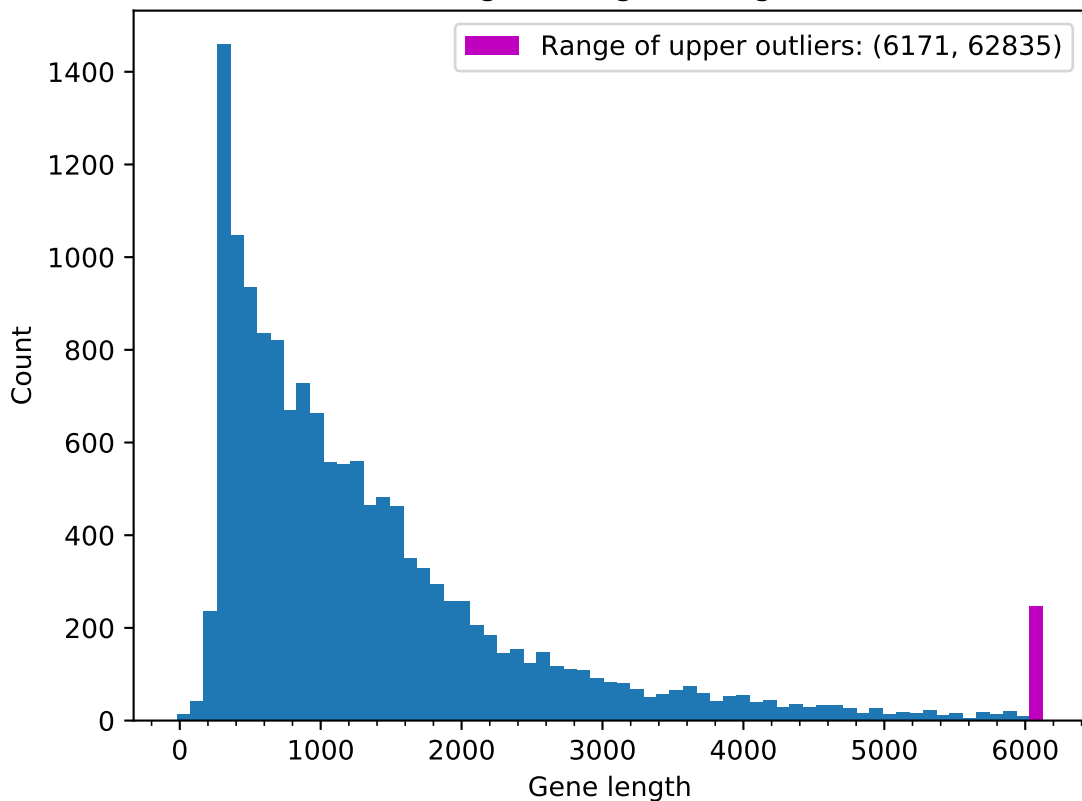
Genes partially supported by external evidence: 10783 (72.68%)

Genes unsupported by any external evidence: 4053 (27.32%)

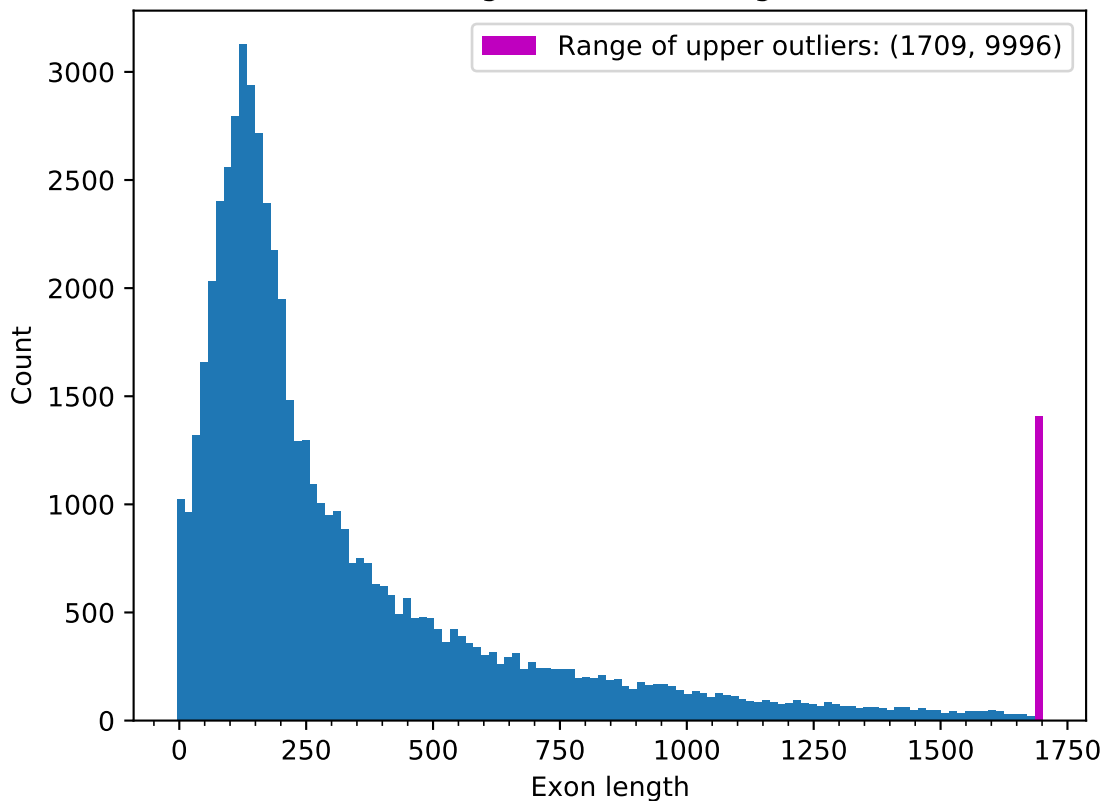
Complete genes: 14597 (98.39%)

Partial genes: 239 (1.61%)

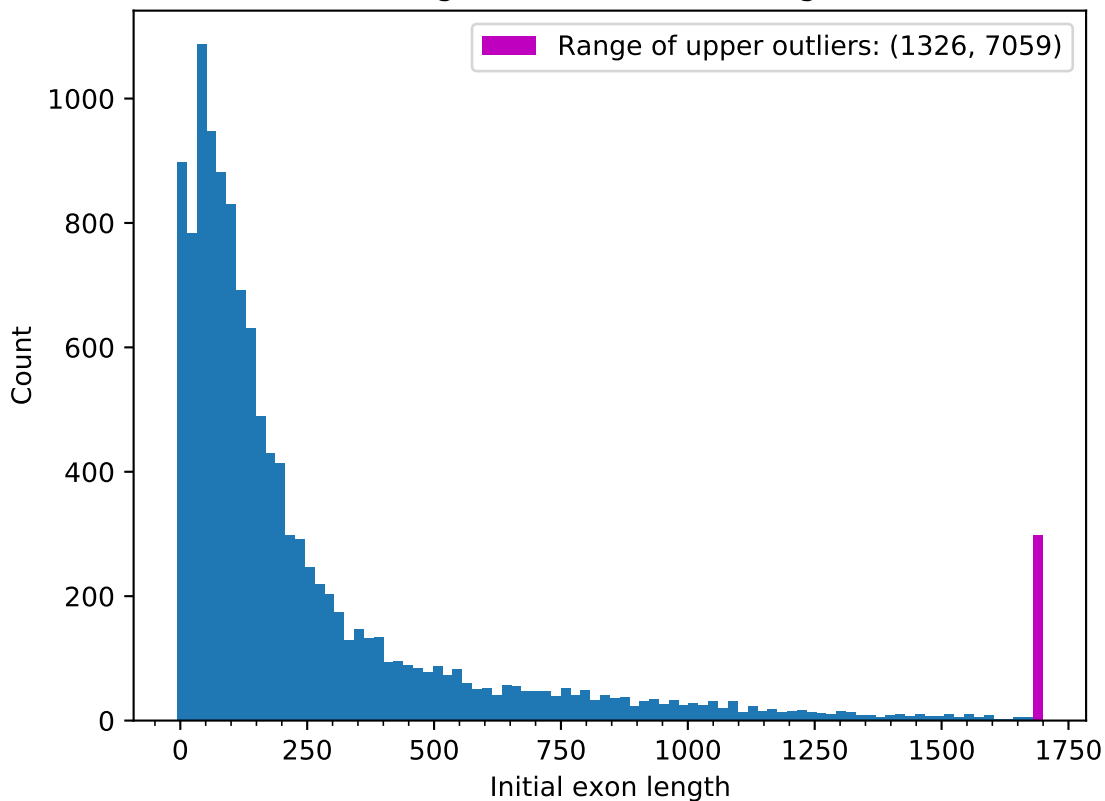
Histogram of gene lengths



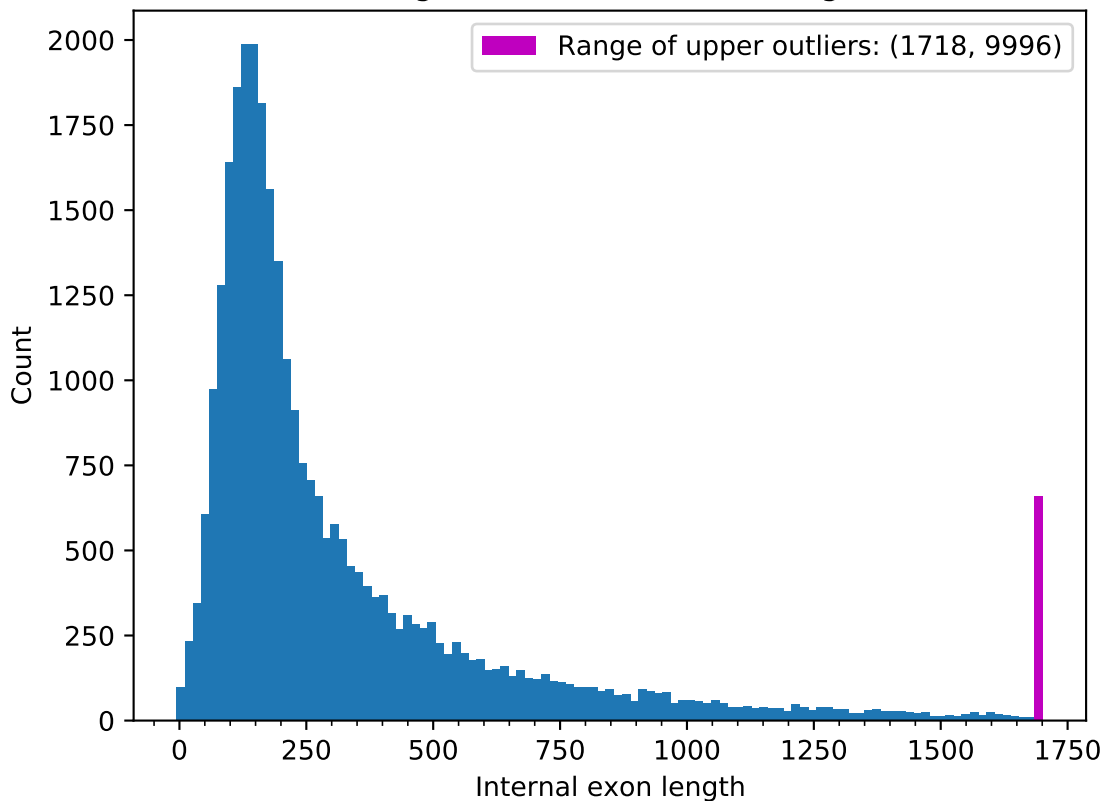
Histogram of exon lengths



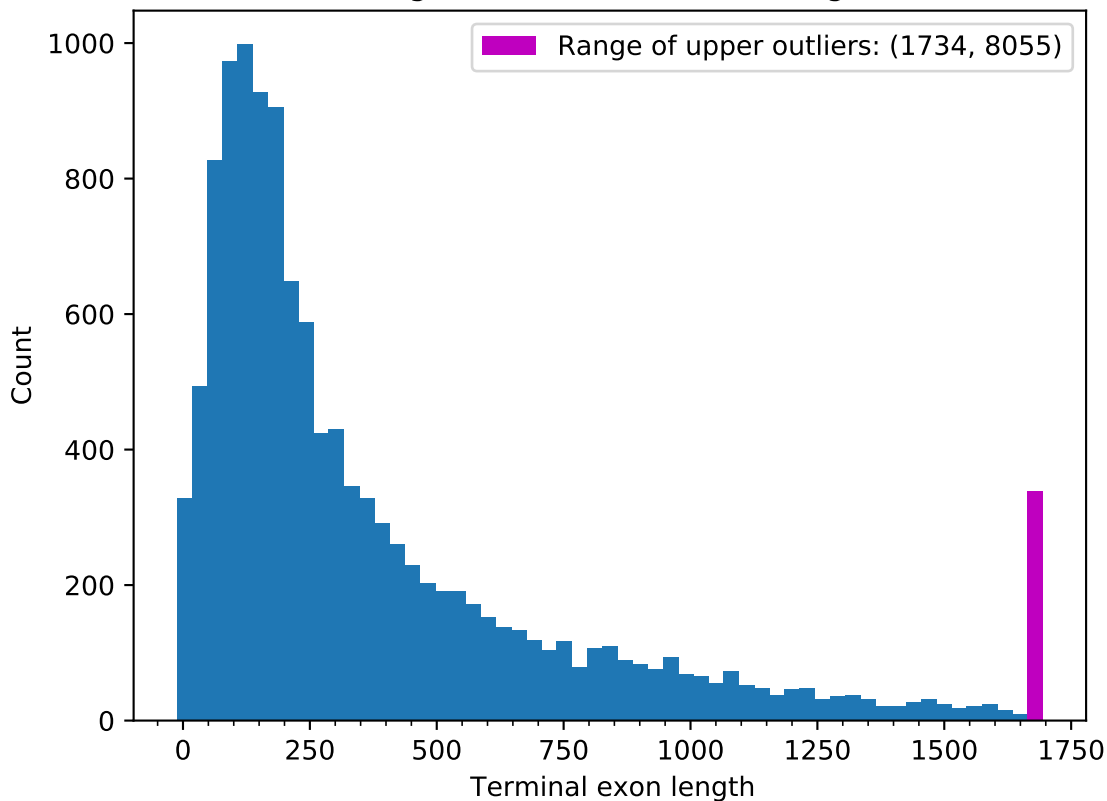
Histogram of initial exon lengths



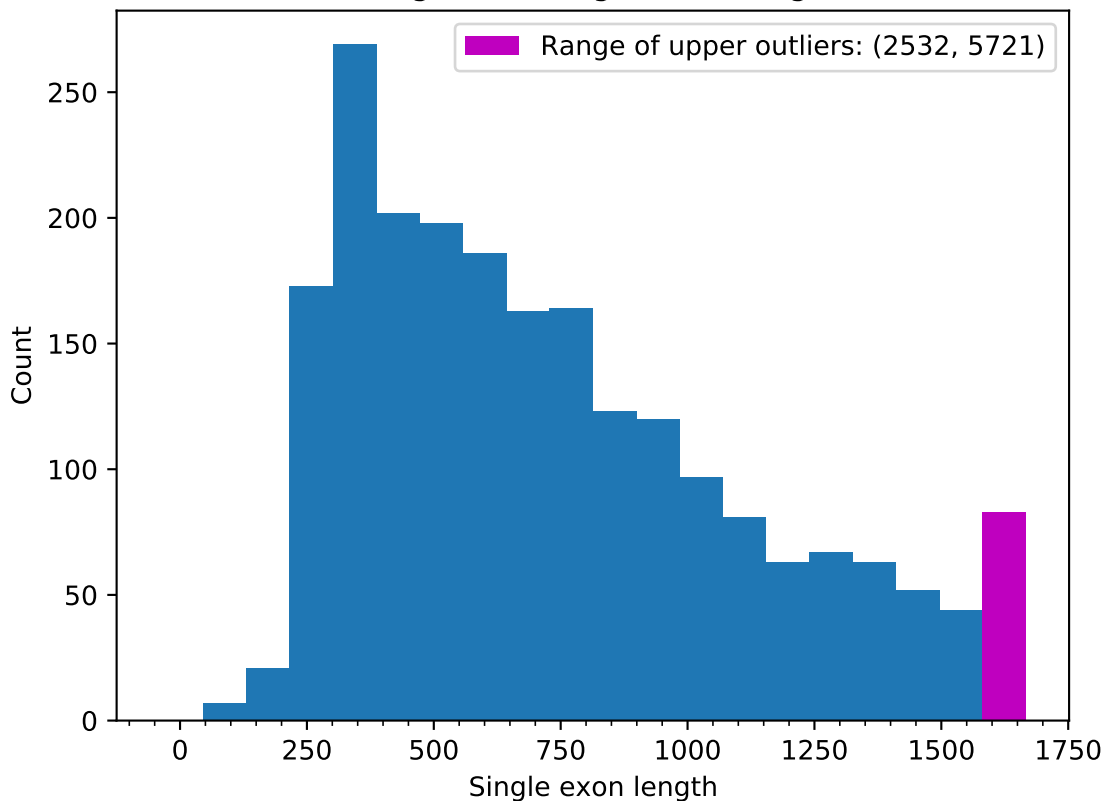
Histogram of internal exon lengths



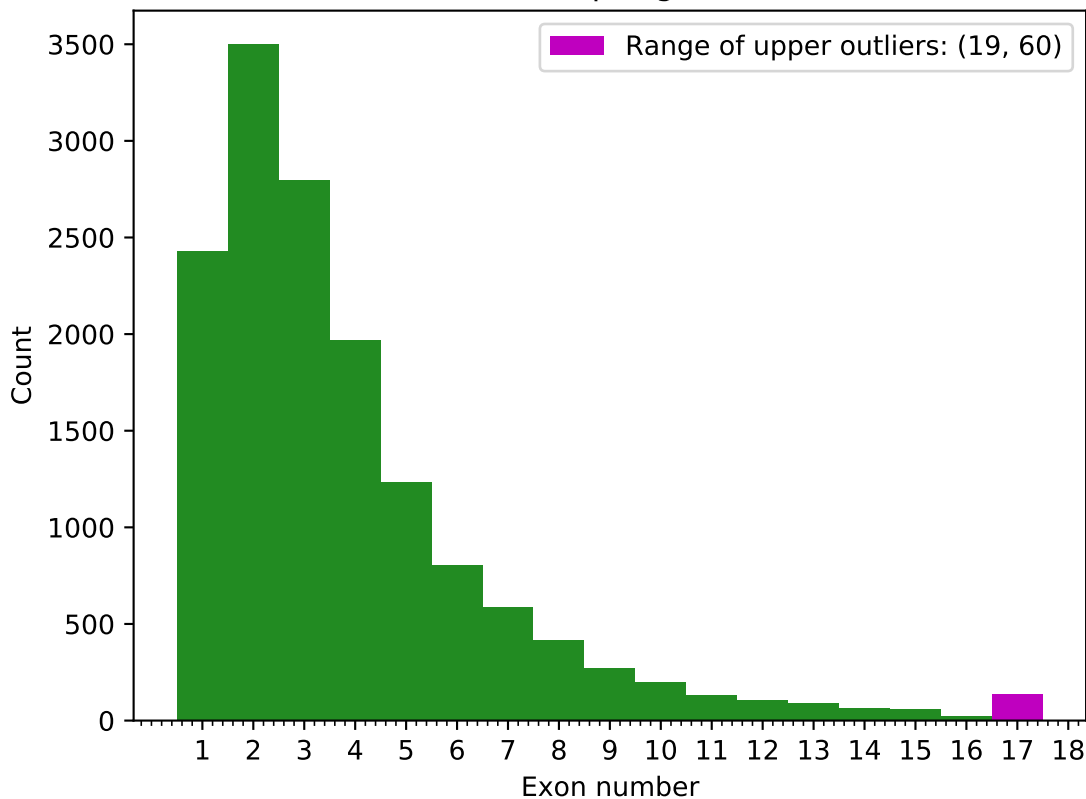
Histogram of terminal exon lengths



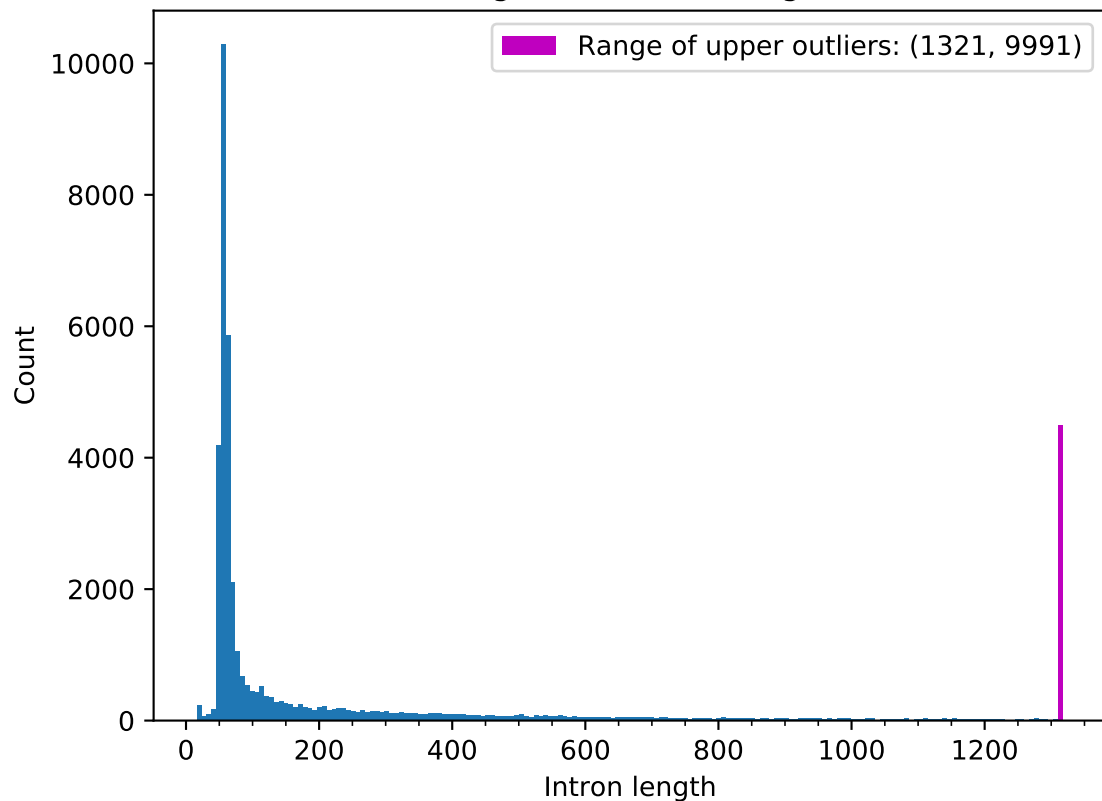
Histogram of single exon lengths



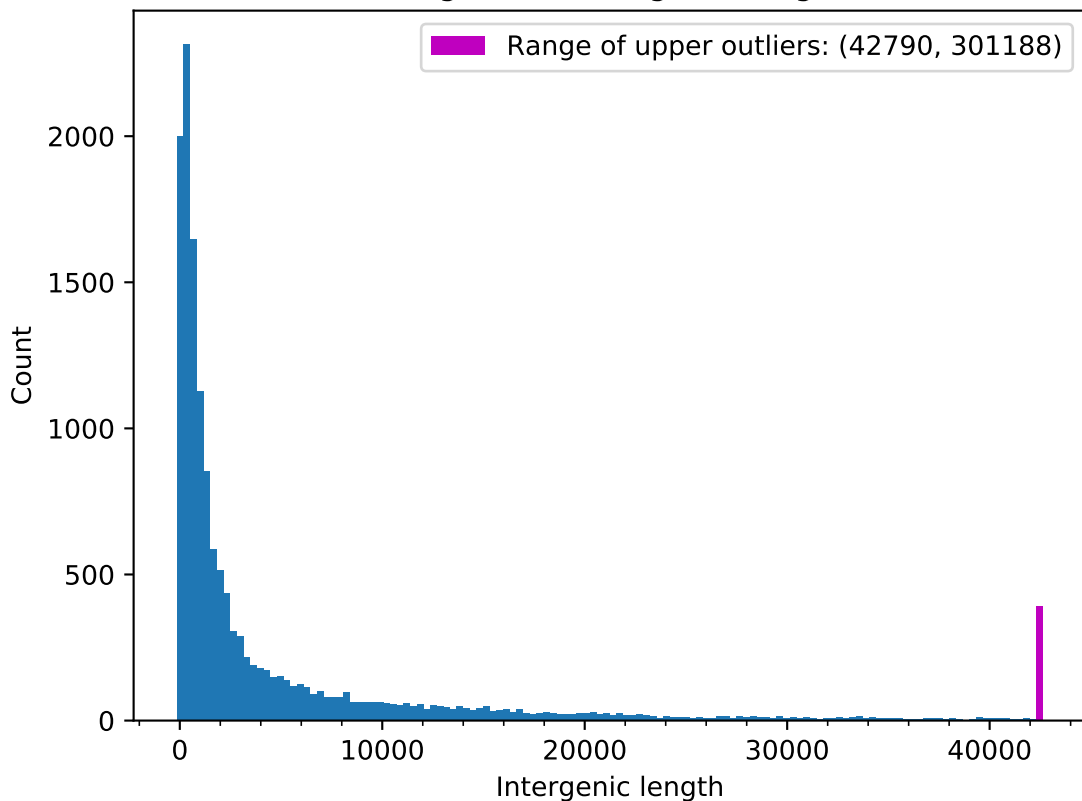
Exons per gene



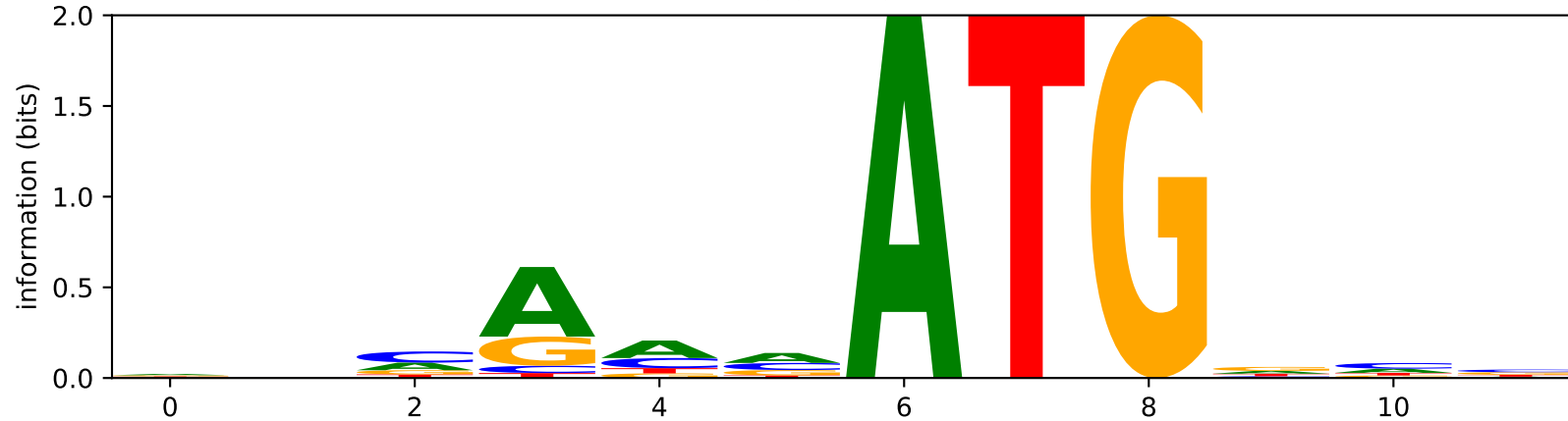
Histogram of intron lengths



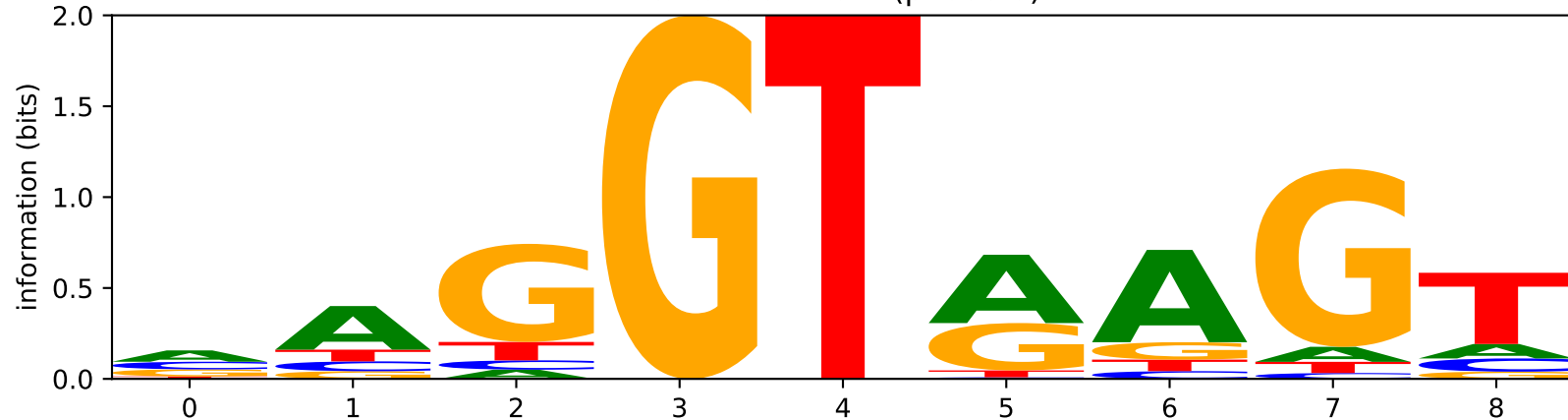
Histogram of intergenic lengths



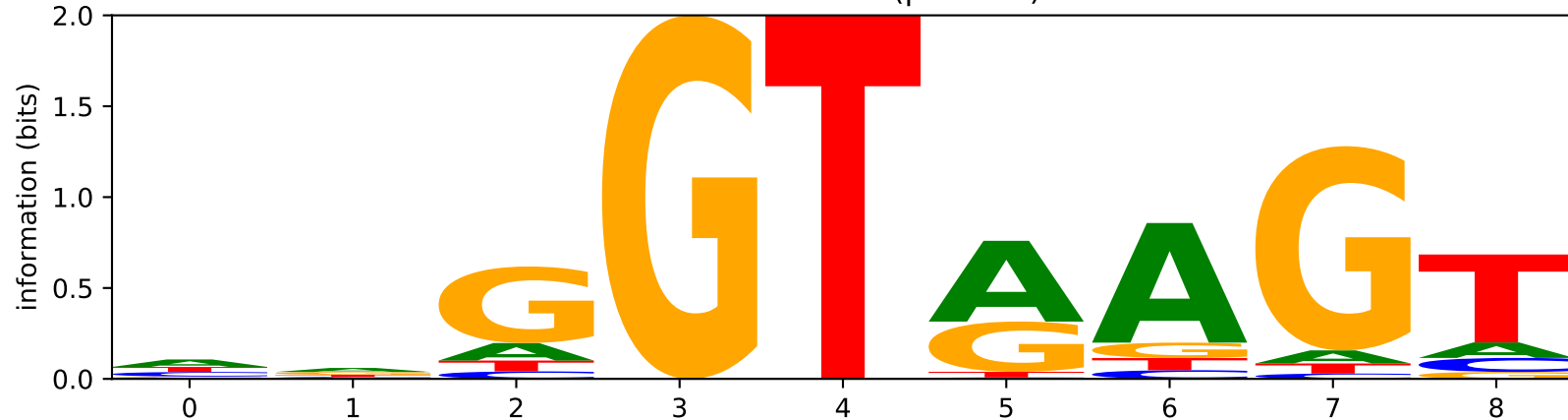
Start motif



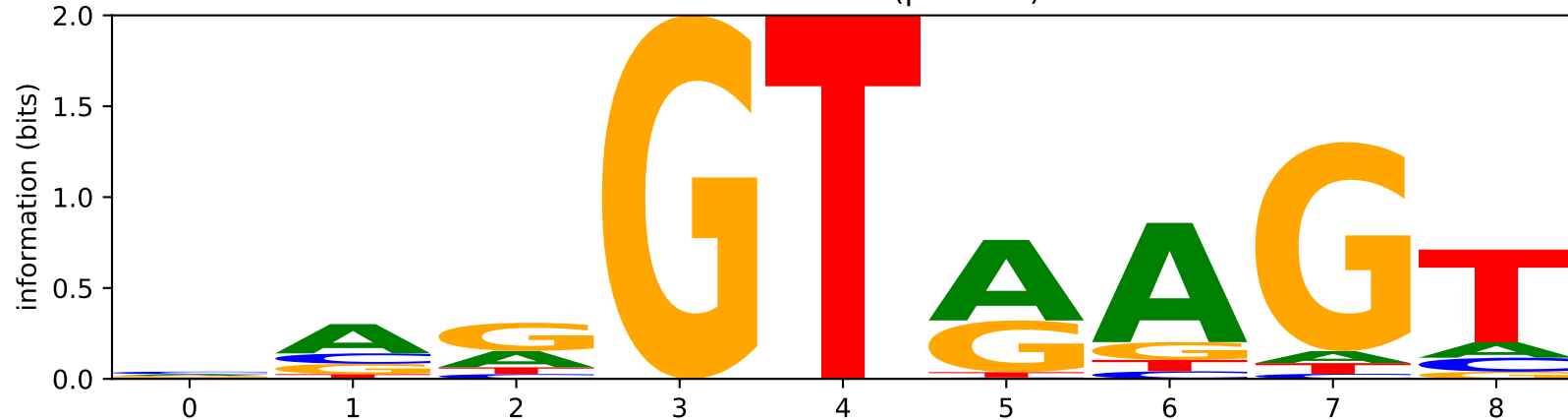
GT donor motif (phase 0)



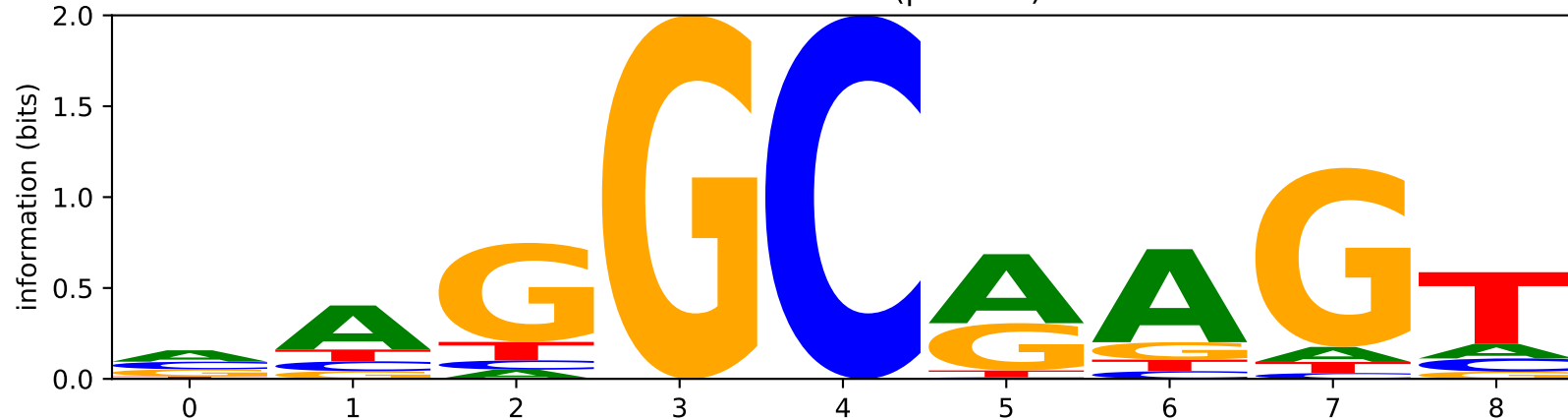
GT donor motif (phase 1)



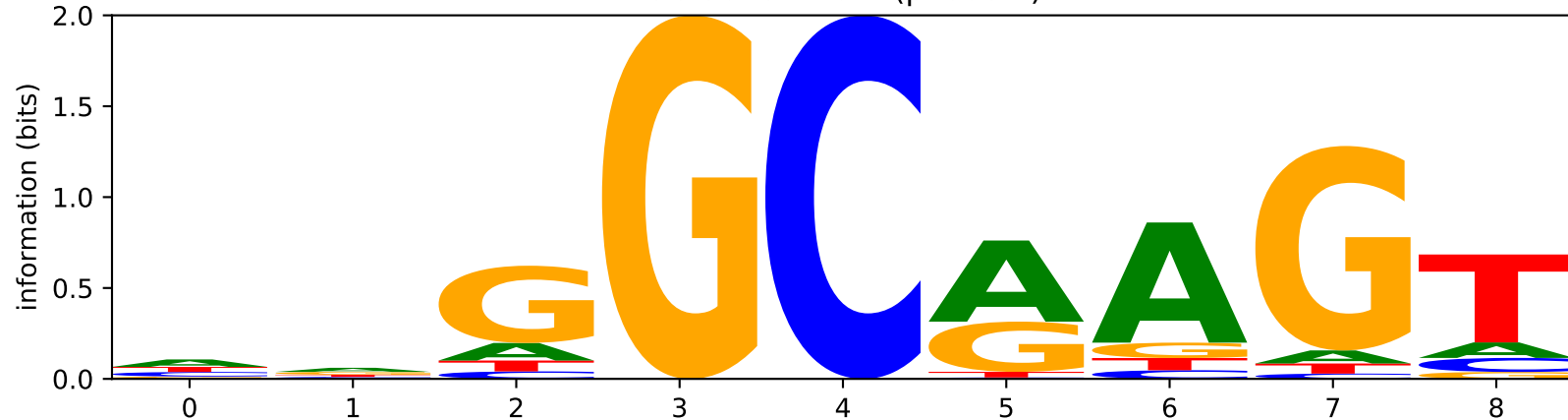
GT donor motif (phase 2)



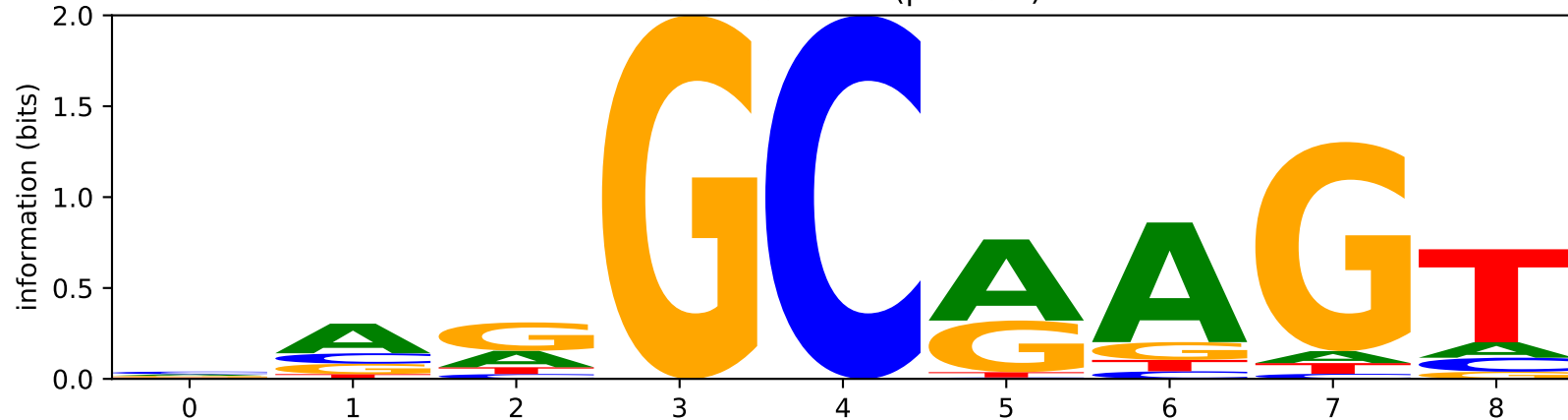
GC donor motif (phase 0)



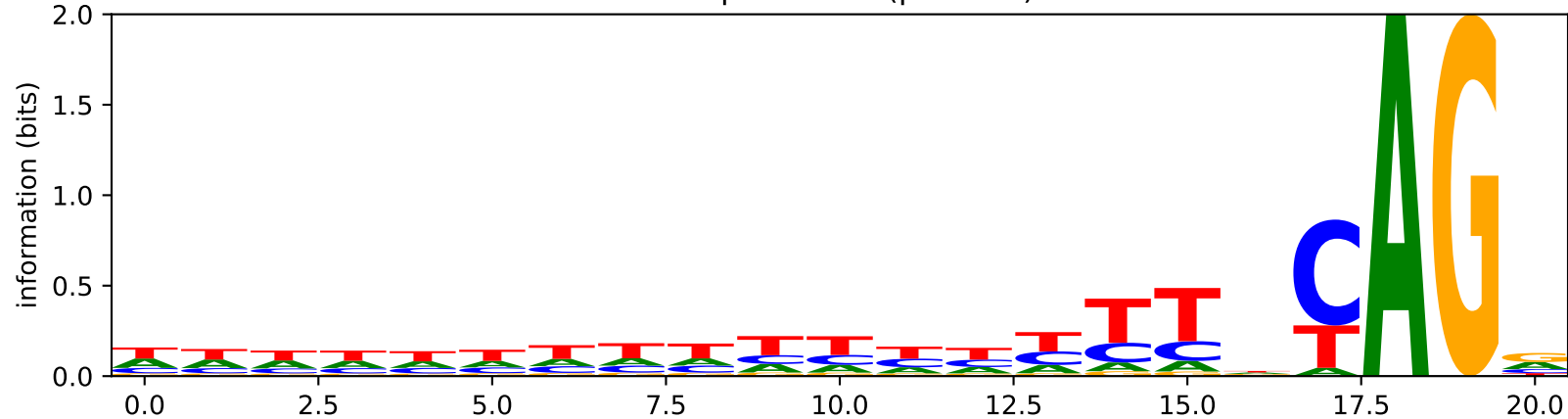
GC donor motif (phase 1)



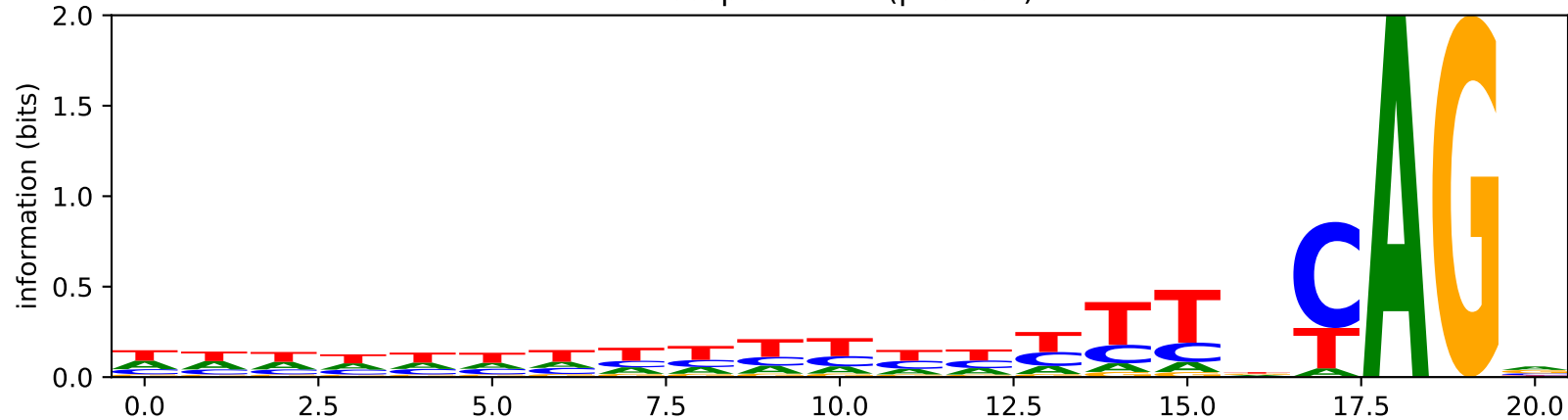
GC donor motif (phase 2)



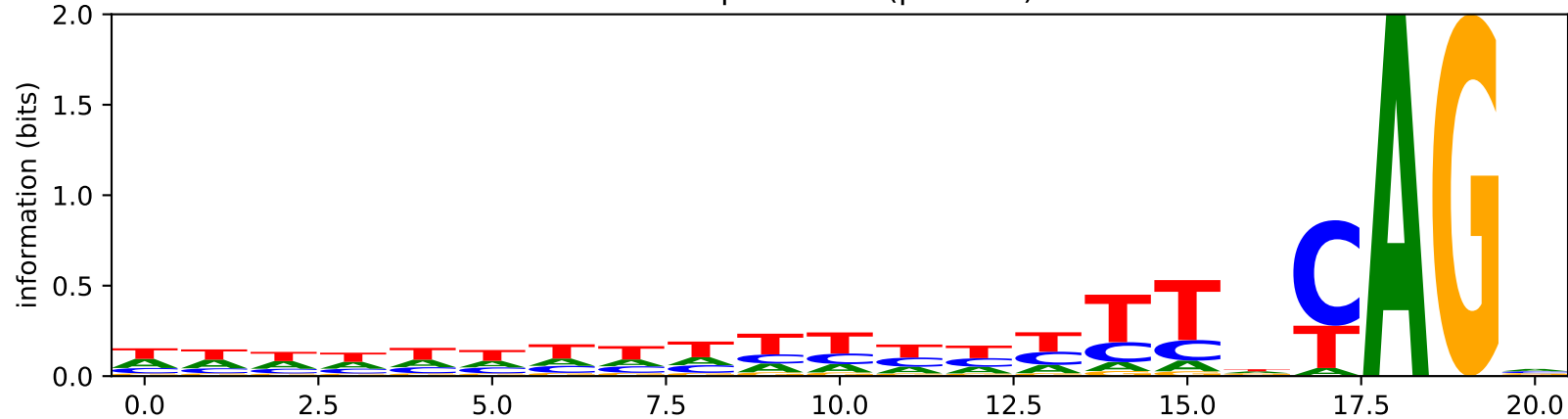
Acceptor motif (phase 0)



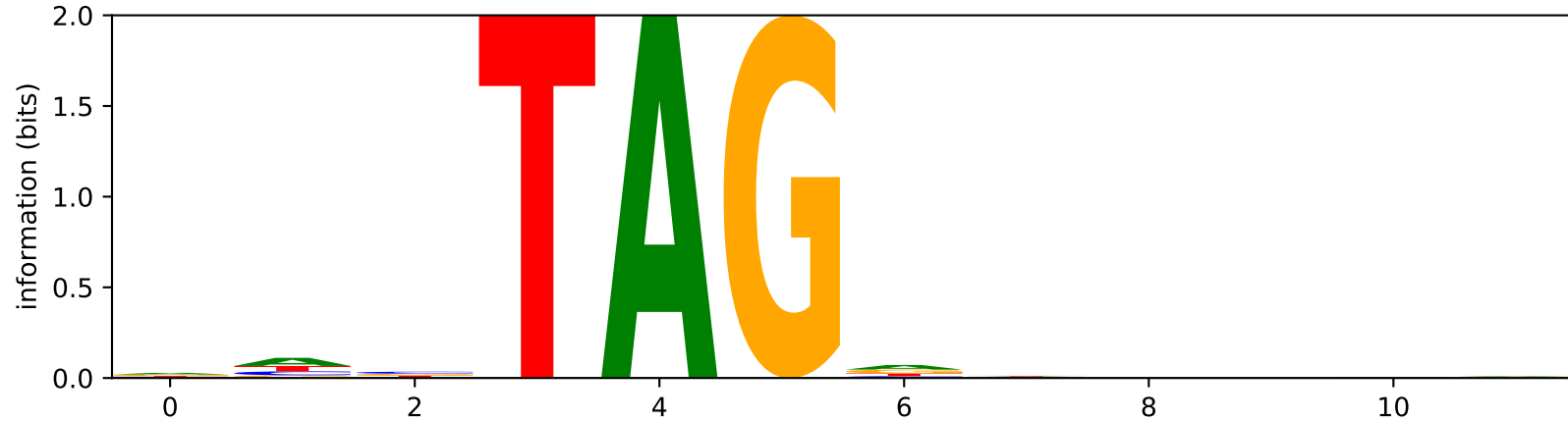
Acceptor motif (phase 1)



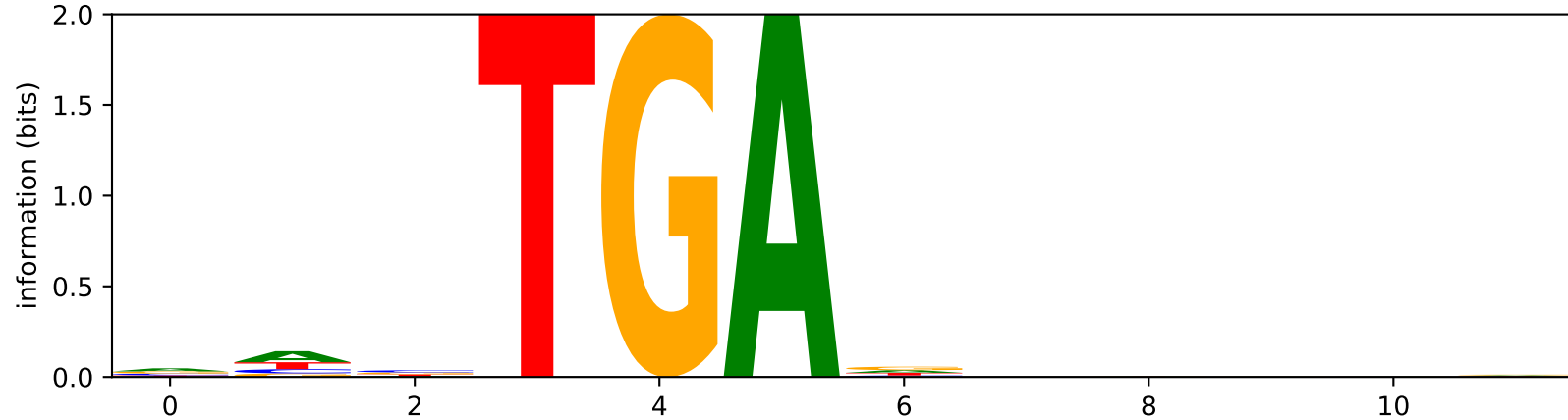
Acceptor motif (phase 2)



Amber motif



Umber motif



Ochre motif

