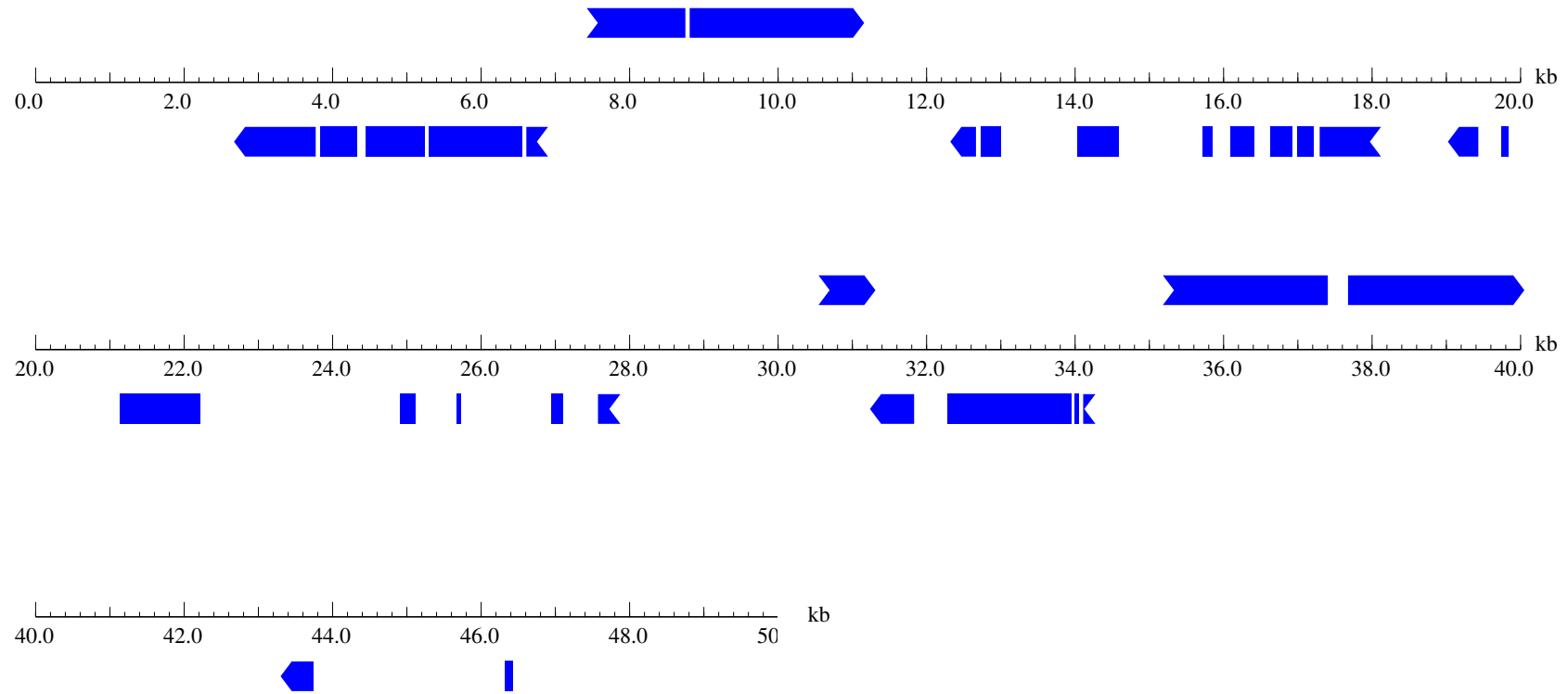


GENSCAN predicted genes in sequence contig32



Key:

Initial
exon

Internal
exon

Terminal
exon

Single-exon
gene

Optimal exon

Suboptimal exon