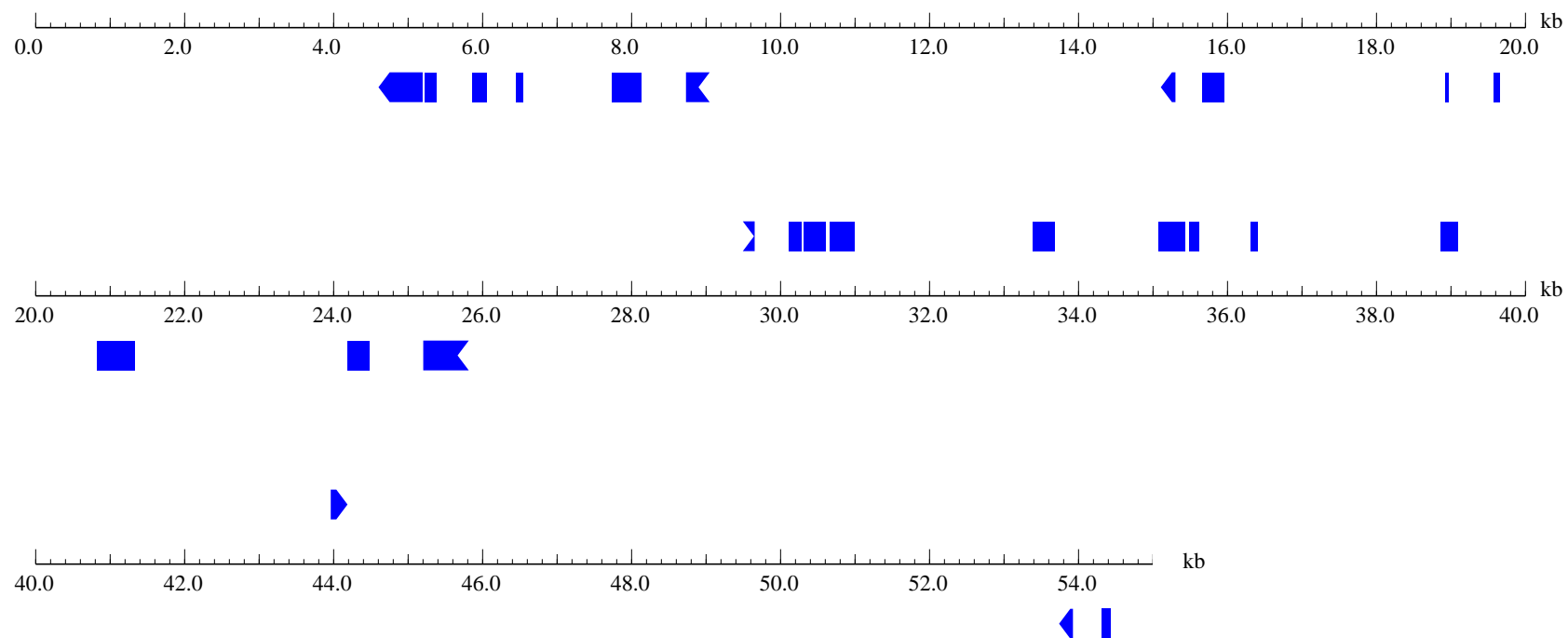


GENSCAN predicted genes in sequence contig32



Key:



Initial
exon



Internal
exon



Terminal
exon



Single-exon
gene



Optimal exon



Suboptimal exon