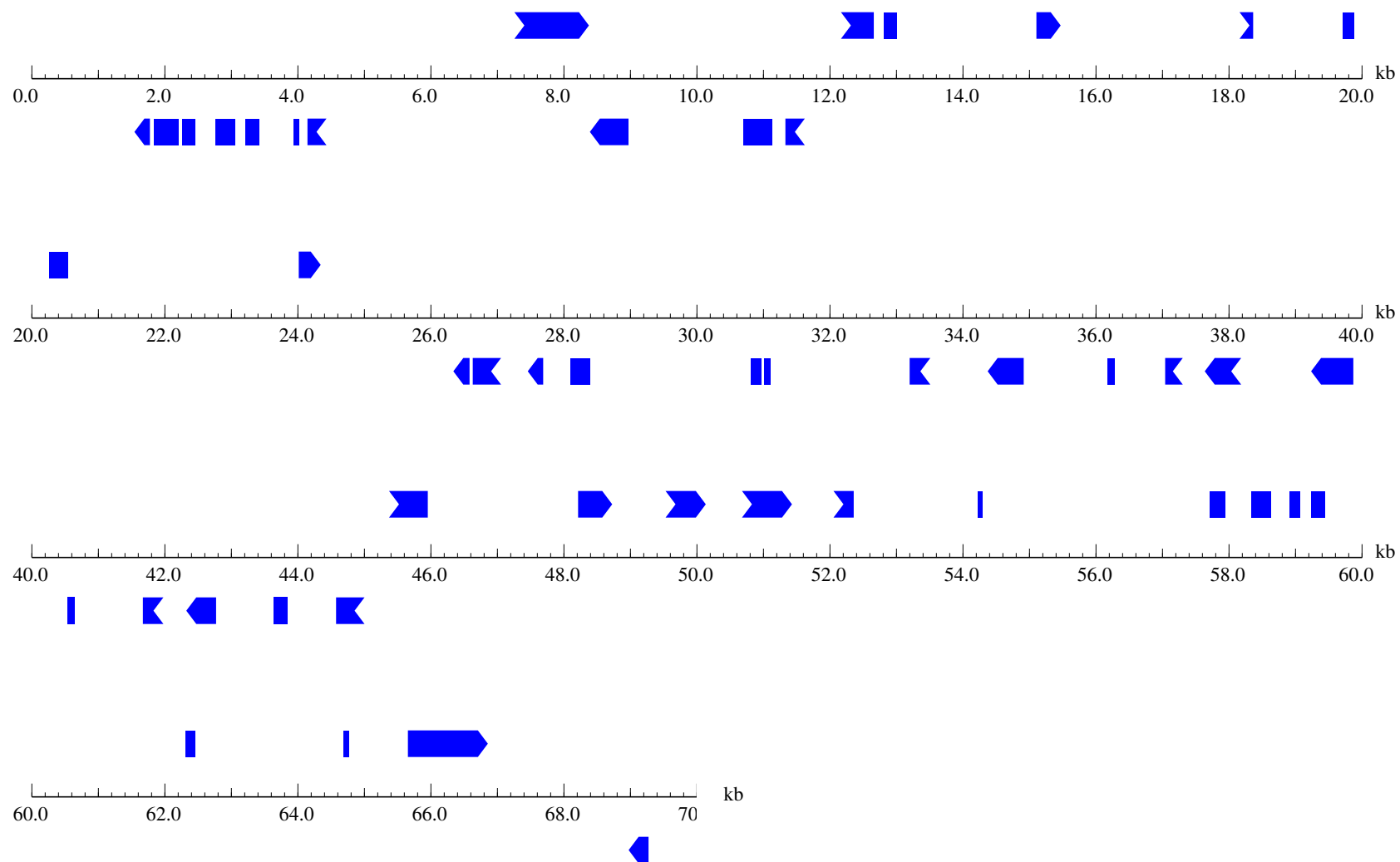


# GENSCAN predicted genes in sequence contig32



**Key:**



Initial  
exon



Internal  
exon



Terminal  
exon



Single-exon  
gene



Optimal exon



Suboptimal exon