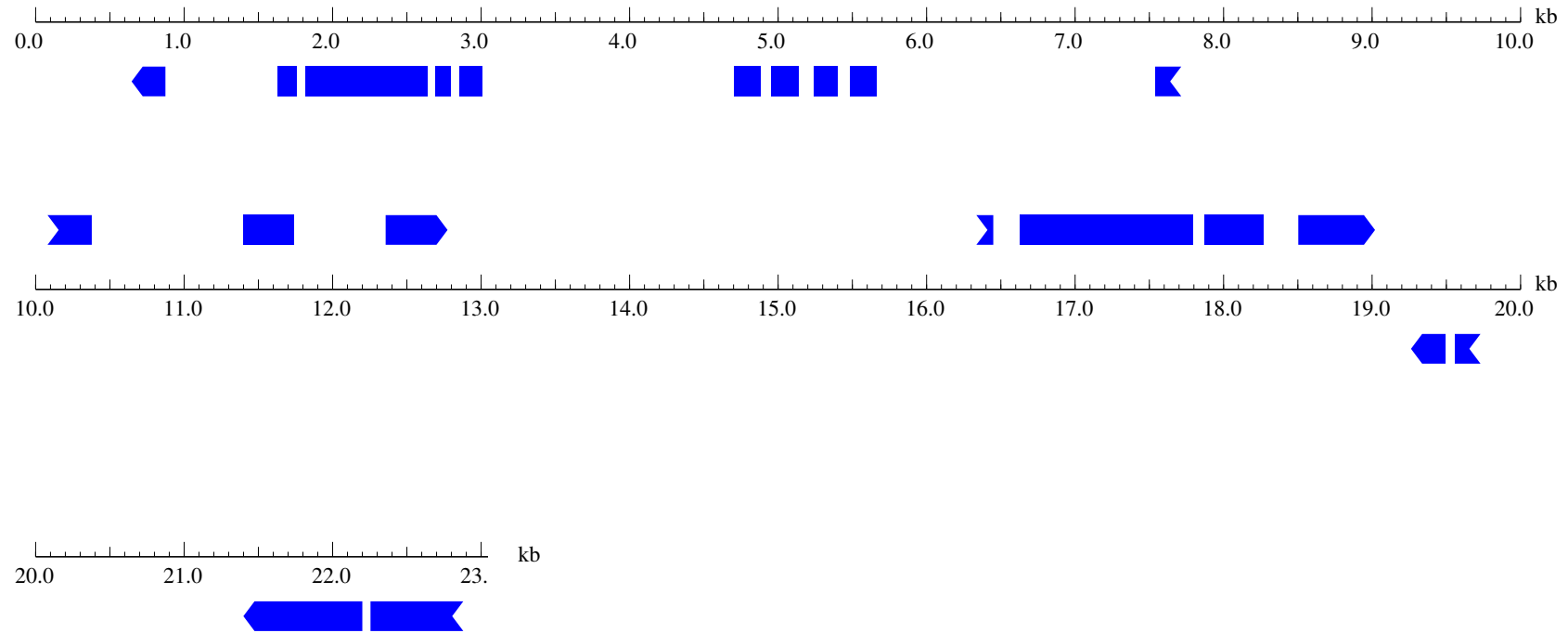


# GENSCAN predicted genes in sequence contig1



**Key:**



Initial  
exon



Internal  
exon



Terminal  
exon



Single-exon  
gene



Optimal exon  
Suboptimal exon