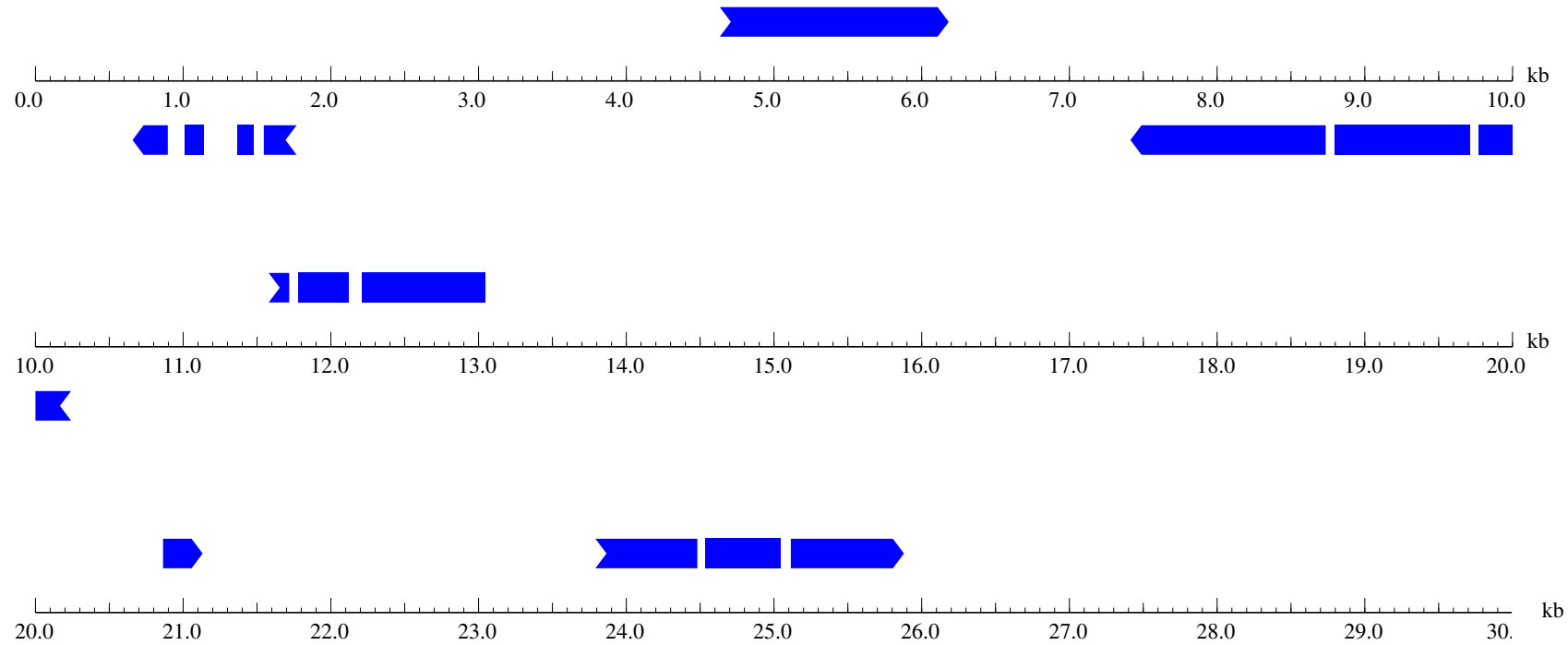


## GENSCAN predicted genes in sequence contig58



**Key:**



Initial  
exon



Internal  
exon



Terminal  
exon



Single-exon  
gene



Optimal exon  
Suboptimal exon