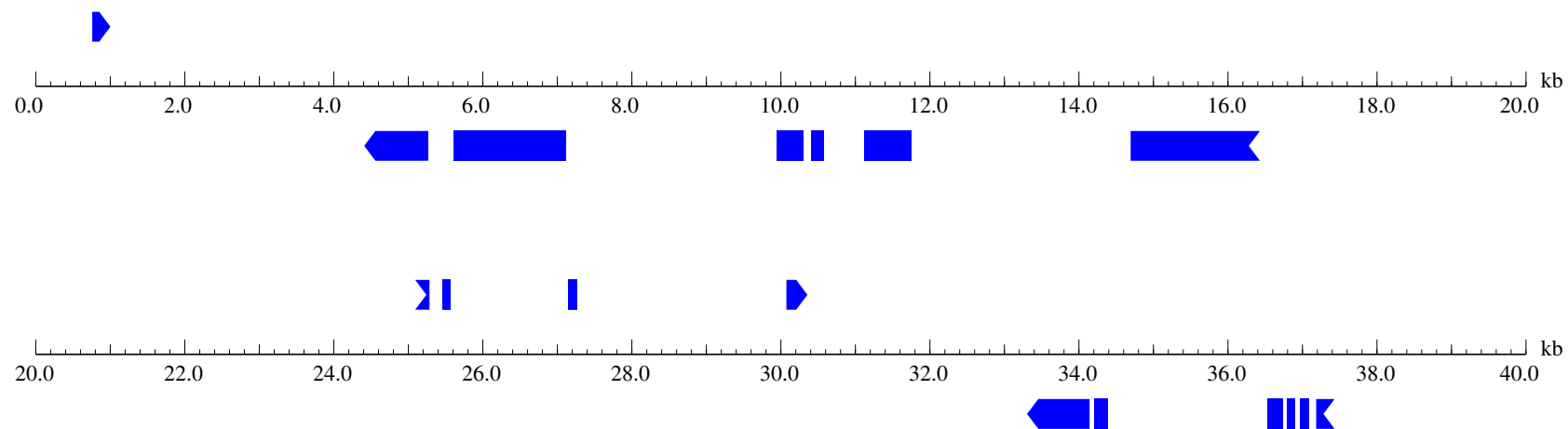


## GENSCAN predicted genes in sequence contig33



40 kb

**Key:**



Initial  
exon



Internal  
exon



Terminal  
exon



Single-exon  
gene



Optimal exon  
Suboptimal exon