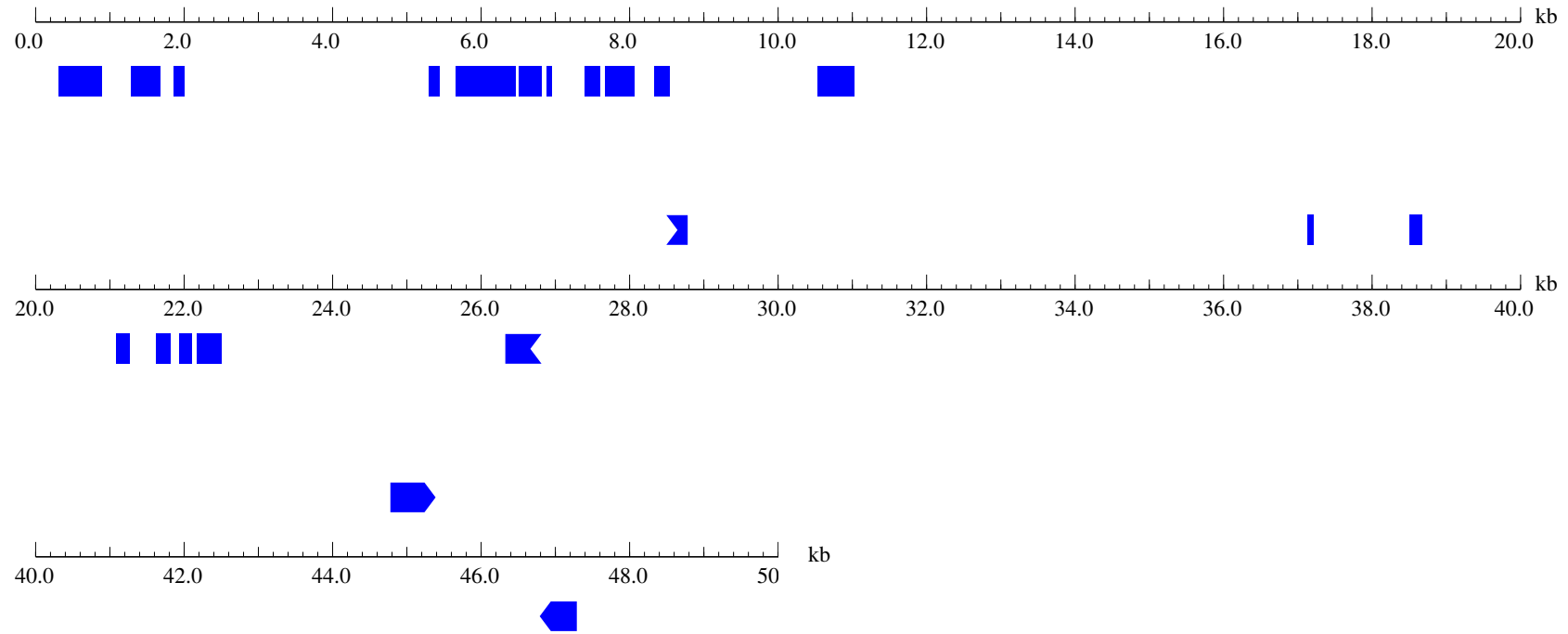


## GENSCAN predicted genes in sequence contig29



**Key:**

Initial  
exon

Internal  
exon

Terminal  
exon

Single-exon  
gene

Optimal exon

Suboptimal exon