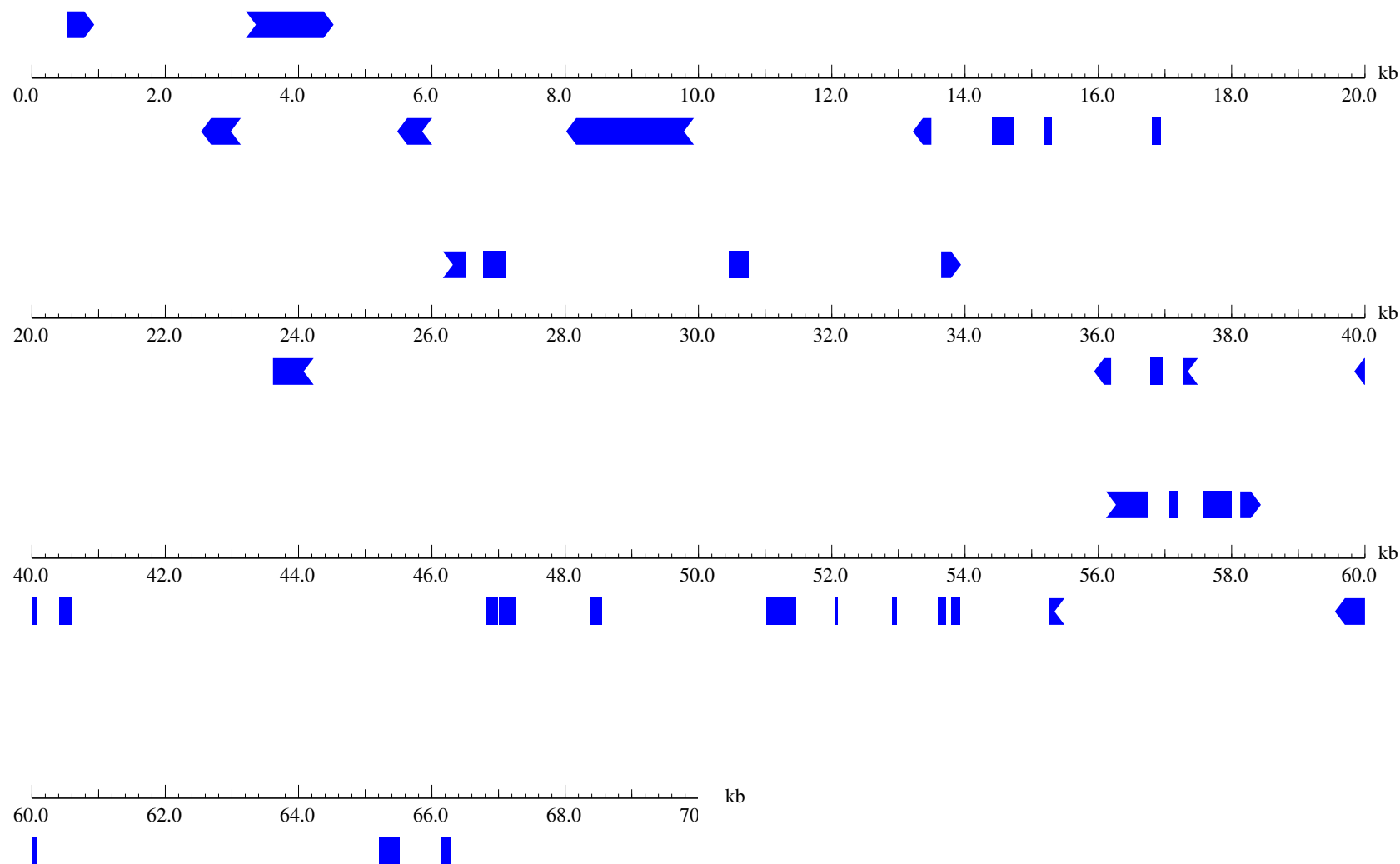





# GENSCAN predicted genes in sequence contig28




**Key:**


 Initial exon

 Internal exon

 Terminal exon

 Single-exon gene

 Optimal exon

 Suboptimal exon