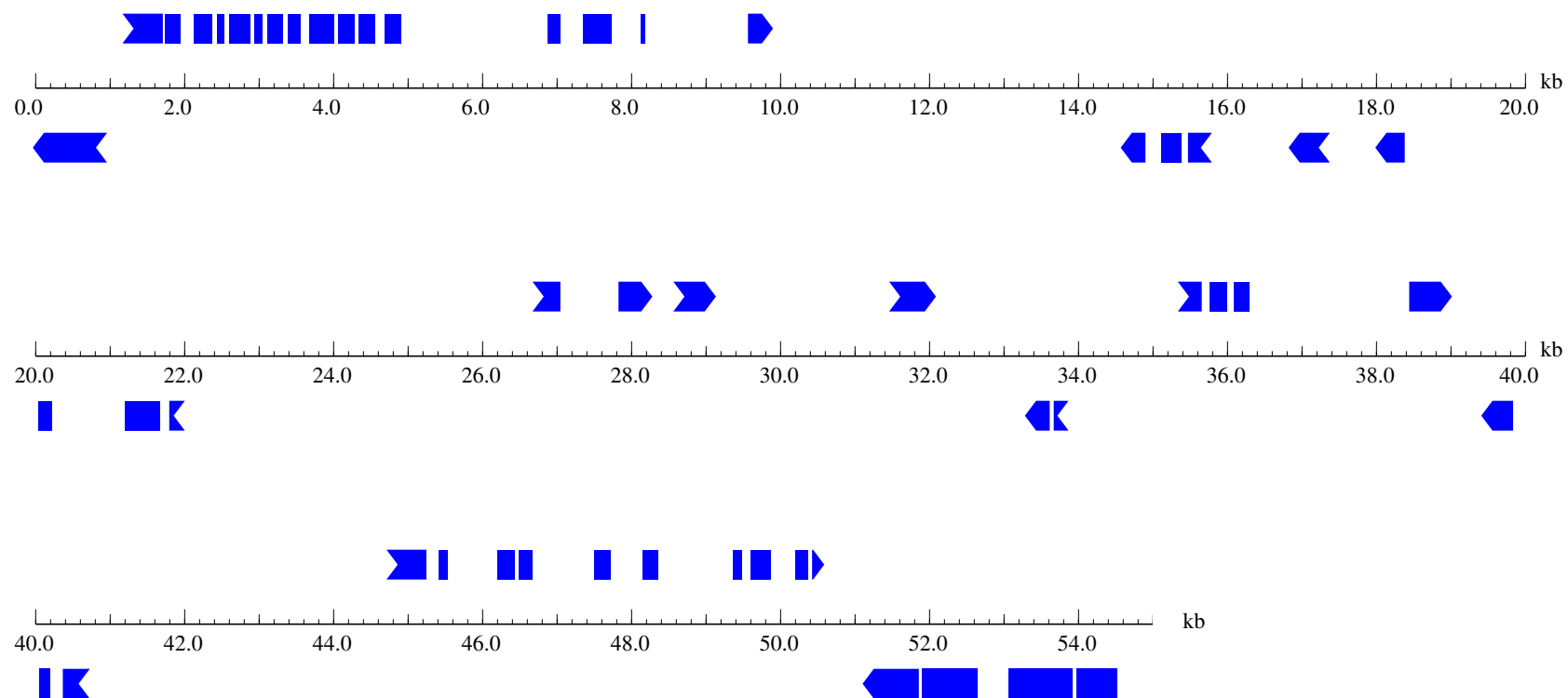




GENSCAN predicted genes in sequence contig52





Key:


 Initial
exon

 Internal
exon

 Terminal
exon

 Single-exon
gene

 Optimal exon

 Suboptimal exon