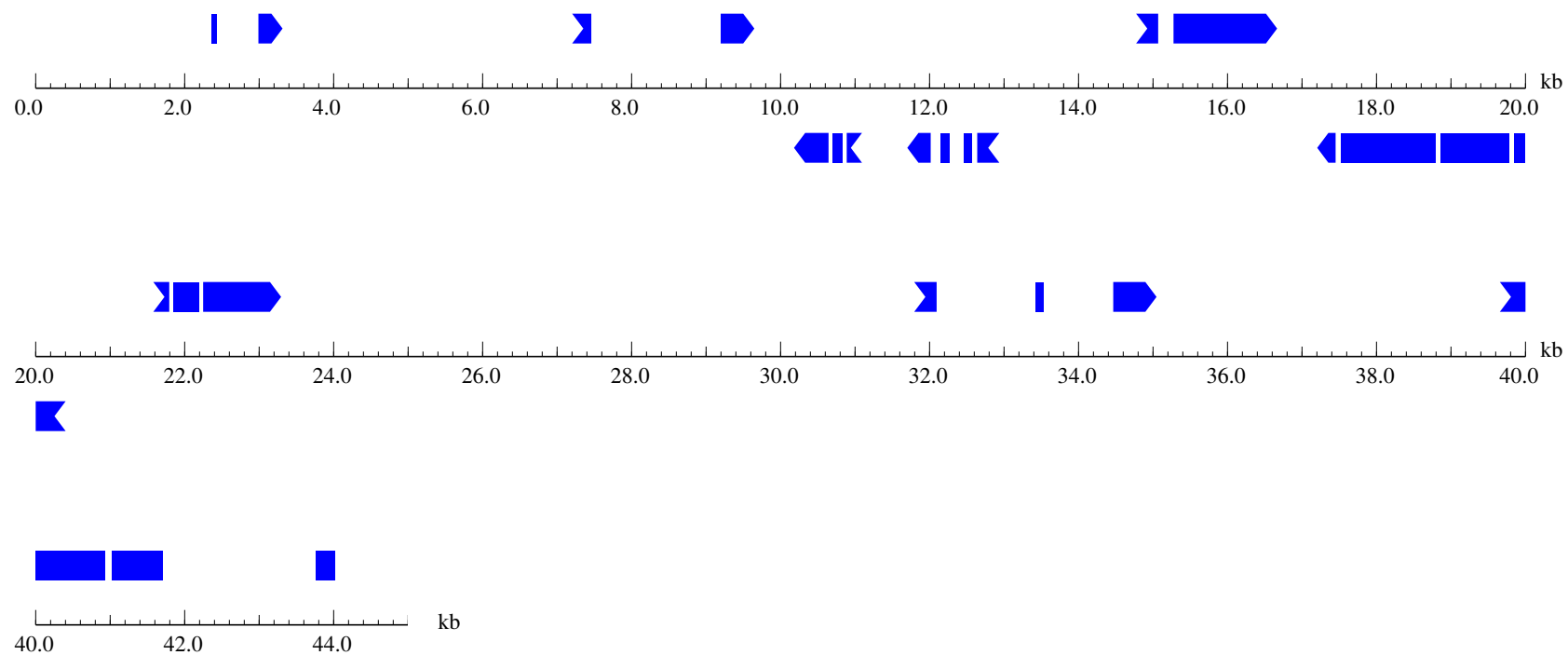


GENSCAN predicted genes in sequence contig1



Key:

Initial exon

Internal exon

Terminal exon

Single-exon gene

Optimal exon

Suboptimal exon