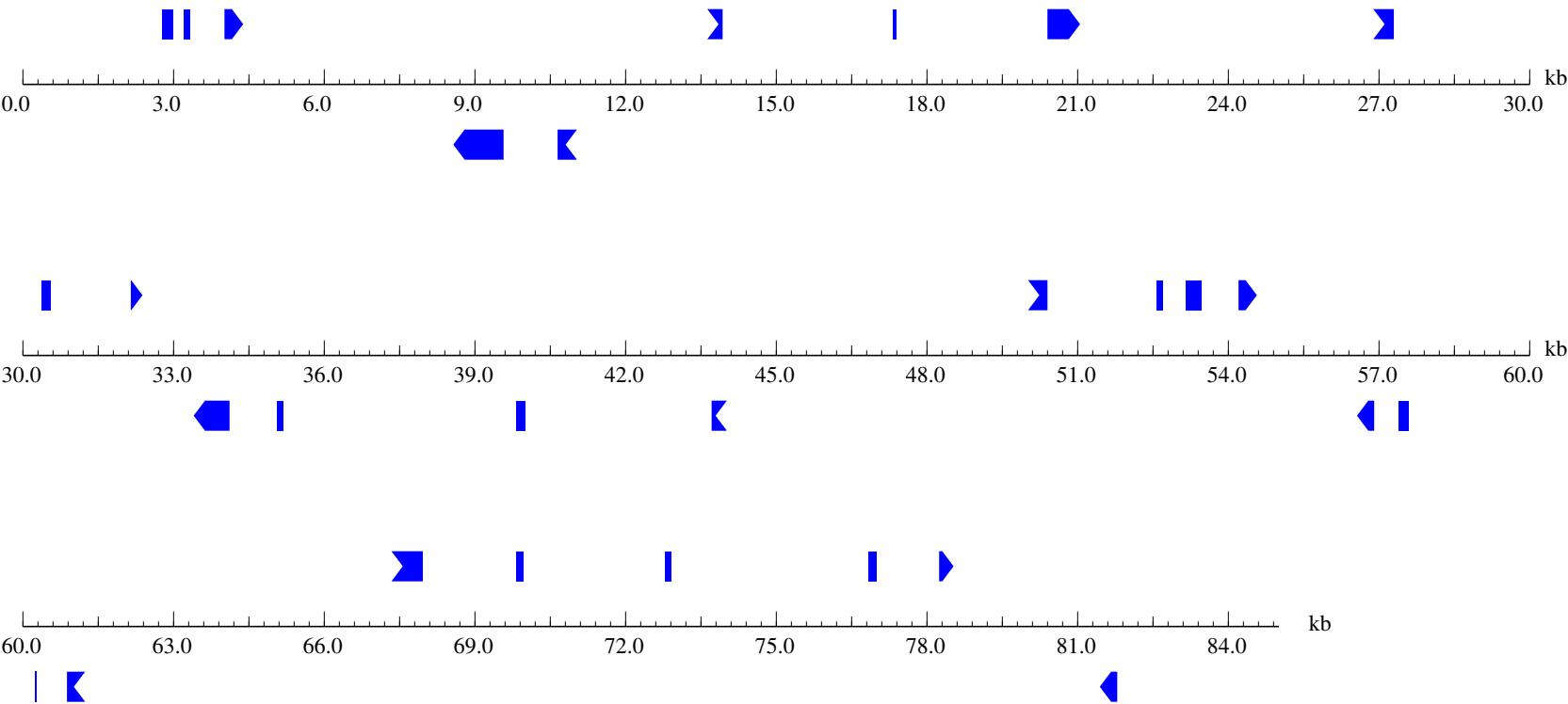


GENSCAN predicted genes in sequence contig32



Key:

	Initial exon		Internal exon		Terminal exon		Single-exon gene		Optimal exon
									Suboptimal exon