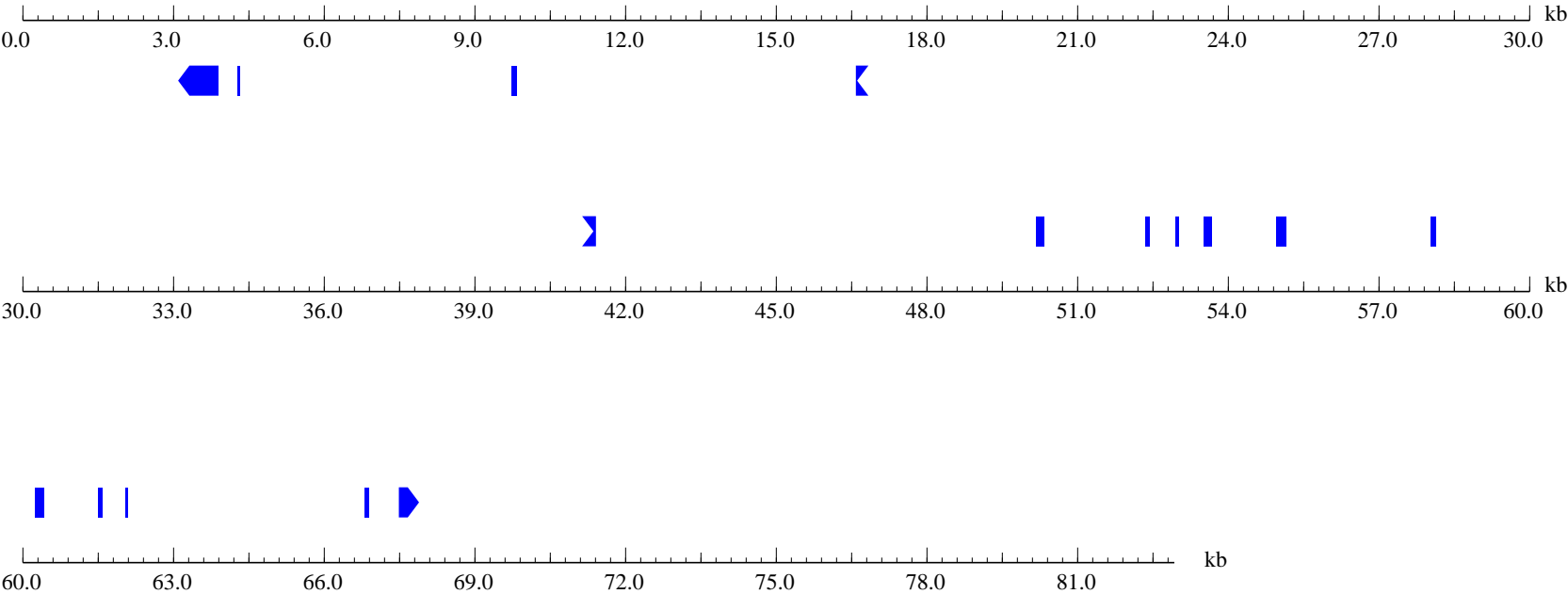


GENSCAN predicted genes in sequence Pan\_troglodytes\_annotation\_chunk\_5



**Key:**

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