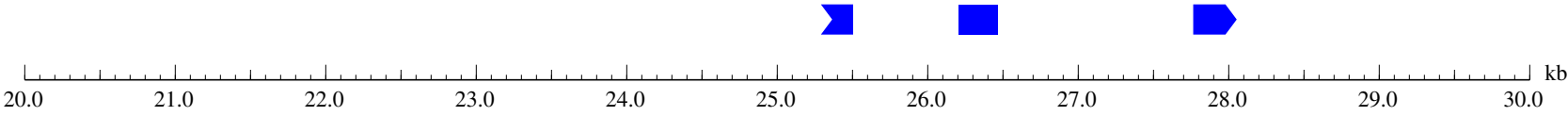
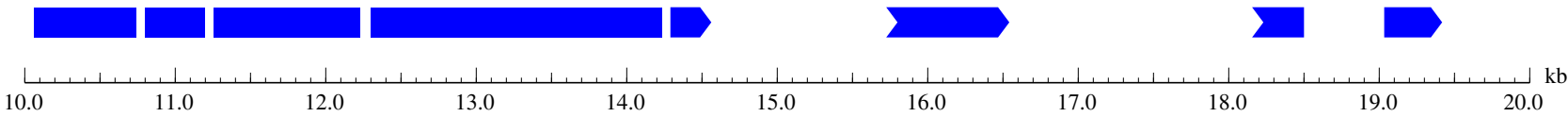
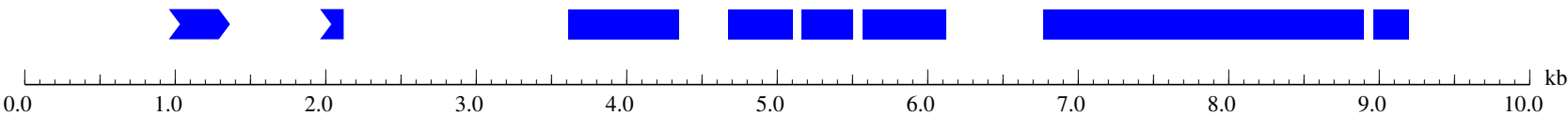


GENSCAN predicted genes in sequence contig34



30.0 kb

