

GENSCAN predicted genes in sequence fosmid34

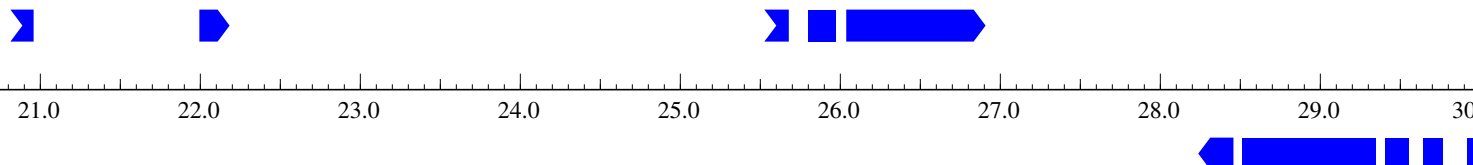
0.0 1.0 2.0 3.0 4.0 5.0 6.0 7.0 8.0 9.0 10.0 kb



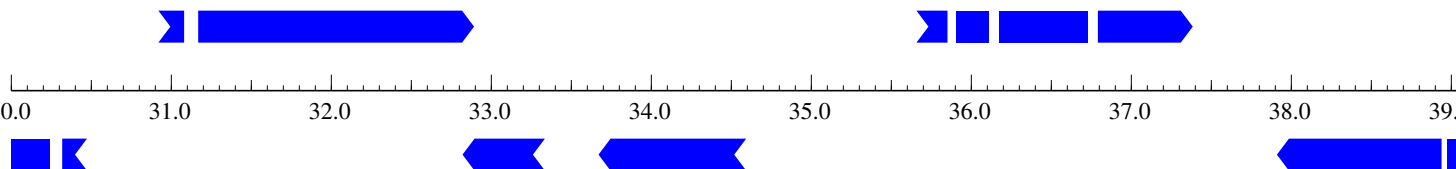
10.0 11.0 12.0 13.0 14.0 15.0 16.0 17.0 18.0 19.0 20.0 kb



20.0 21.0 22.0 23.0 24.0 25.0 26.0 27.0 28.0 29.0 30.0 kb



30.0 31.0 32.0 33.0 34.0 35.0 36.0 37.0 38.0 39.0 40.0 kb



Key:

Initial
exon

Internal
exon

Terminal
exon

Single-exon
gene

Optimal exon
Suboptimal exon