

GENSCAN predicted genes in sequence fosmid_2602K20

0.0 2.0 4.0 6.0 8.0 10.0 12.0 14.0 16.0 18.0 20.0 kb



20.0 22.0 24.0 26.0 28.0 30.0 32.0 34.0 36.0 38.0 40.0 kb



40.0 kb

Key:



Initial
exon



Internal
exon



Terminal
exon



Single-exon
gene



Optimal exon



Suboptimal exon