

Scale

chr10:
-->

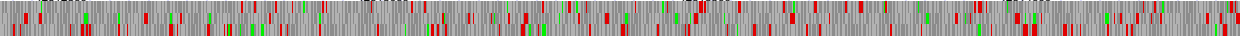
500 bases

42612500

42613000

42613500

42614000

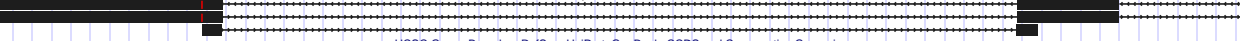


Your Sequence from Blat Search

MG001-38_M03

MG001-38_M01

ALEXA_Seq_Junction



UCSC Genes Based on RefSeq, UniProt, GenBank, CCDS and Comparative Genomics

BMS1



Ensembl Gene Predictions

ENST00000374518

