

Scale

500 bases

chr5:

176891000

176891500

176892000

--->



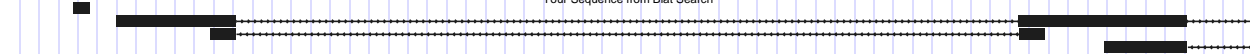
Your Sequence from Blat Search

Reverse_Primer

MG001-38_I07

ALEXA_Seq_Junction

MG001-38_I05



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

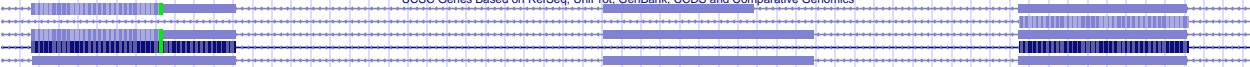
KIAA1931

FLJ10404

KIAA1931

FLJ10404

KIAA1931



Ensembl Gene Predictions

ENST00000398118

ENST00000329540

