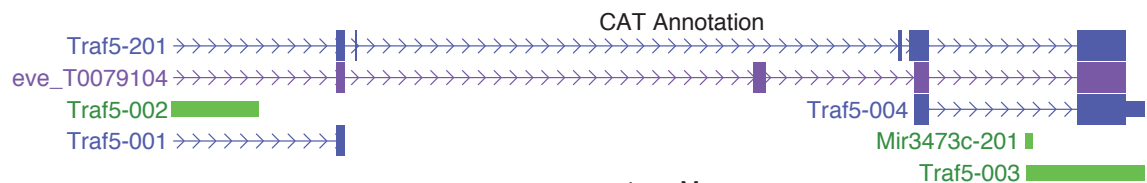


HYBRID_SCAFFOLD_Super-Scaffold_8976:408,462-422,561 (14,100 bp)

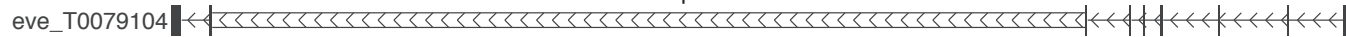


Eve IsoSeq alignments



chr1:191,997,430-192,070,634 (73,205 bp)

Your Sequence from Blat Search



Basic Gene Annotation Set from GENCODE Version M16 (Ensembl 91)



Assembly from Fragments

