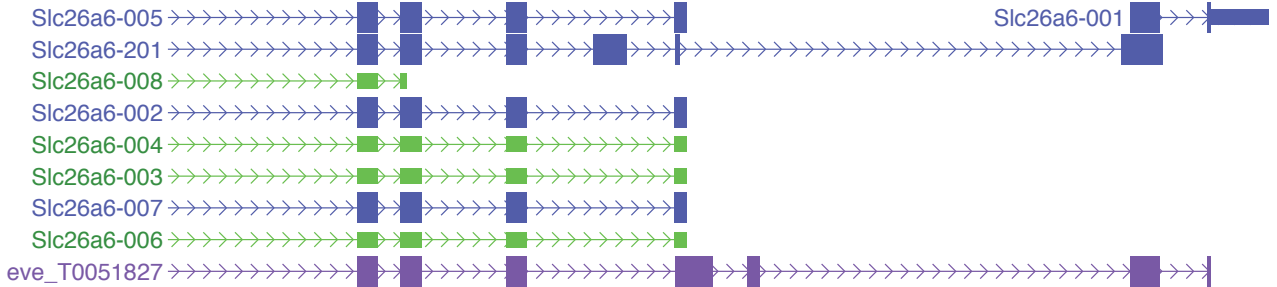
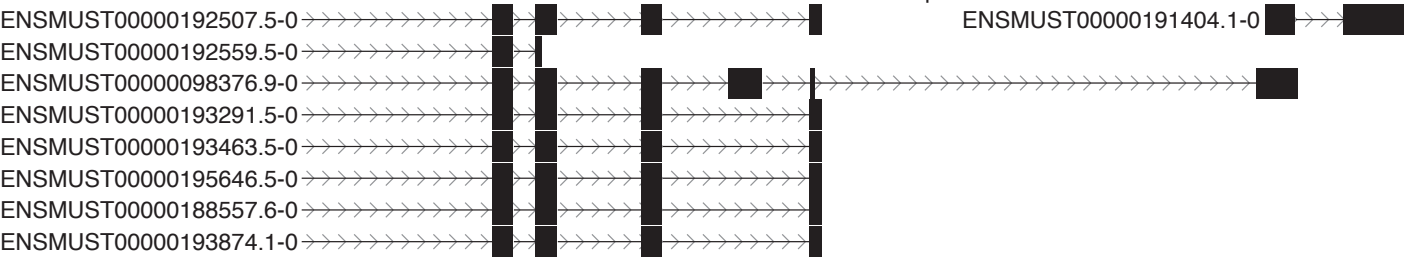


HYBRID_SCAFFOLD_Super-Scaffold_402:1,651,017-1,656,516 (5,500 bp)

CAT Annotation



transMap



GRCm38

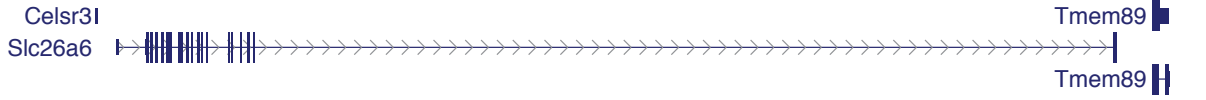


Eve IsoSeq alignments



chr9:108,852,821-108,921,536 (68,716 bp)

Basic Gene Annotation Set from GENCODE Version M14 (Ensembl 89)



Assembly from Fragments

GL456151.2

GL456152.2