

NW_027185390.1:2380131..2380193 Embiotoca jacksoni isolate LCO1_2020_lvr unplaced genomic scaffold, fEmbJac1.0.p SCAF_38, whole genome shotgun sequence

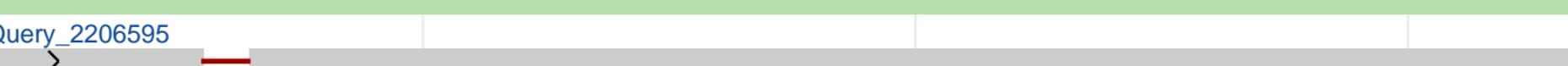
2,380,140 2,380,150 2,380,160 2,380,170 2,380,180 2,380,190

A T G C G C C T T G G G G C T T G G A C G T A C G T A A C A C A T C C T G A T G G G G A A A A G G A A C C T G T T G A C A A C T G A C T G G T A C C C C T T T C C T T G G A C A A A C T G T

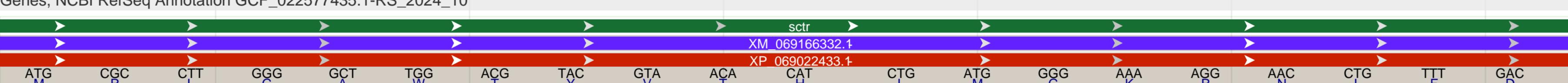
(U) Alignment for group templates, BLAST Results for: XM_069166332.1:40-1458 PREDICTED: Embiotoca...



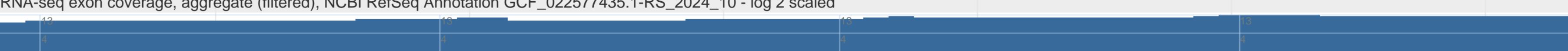
(U) BLAST Results for: XM_069166332.1:40-1458 PREDICTED: Embiotoca...



Genes, NCBI RefSeq Annotation GCF_022577435.1-RS_2024_10



RNA-seq exon coverage, aggregate (filtered), NCBI RefSeq Annotation GCF_022577435.1-RS_2024_10 - log 2 scaled



RNA-seq intron-spanning reads, aggregate (filtered), NCBI RefSeq Annotation GCF_022577435.1-RS_2024_10 - log 2 scaled



RNA-seq intron features, aggregate (filtered), NCBI RefSeq Annotation GCF_022577435.1-RS_2024_10