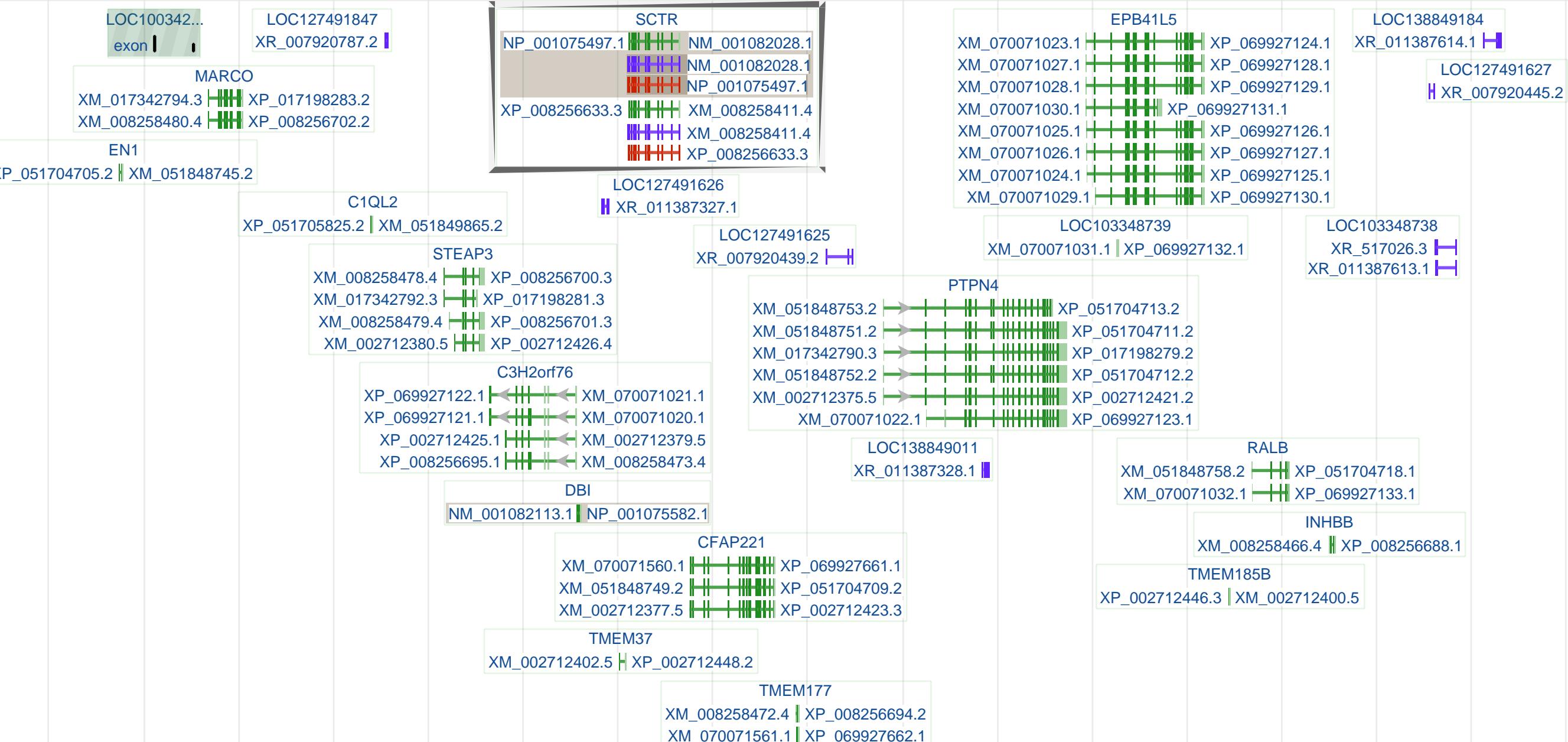


## NC\_091434.1:116347594..118328360 Oryctolagus cuniculus chromosome 3, mOryCun1.1, whole genome shotgun sequence

116,400 K 116,500 K 116,600 K 116,700 K 116,800 K 116,900 K 117 M 117,100 K 117,200 K 117,300 K 117,400 K 117,500 K 117,600 K 117,700 K 117,800 K 117,900 K 118 M 118,100 K 118,200 K

Genes, NCBI RefSeq Annotation GCF\_964237555.1-RS\_2024\_11



RNA-seq exon coverage, aggregate (filtered), NCBI RefSeq Annotation GCF\_964237555.1-RS\_2024\_11 - log 2 scaled



RNA-seq intron-spanning reads, aggregate (filtered), NCBI RefSeq Annotation GCF\_964237555.1-RS\_2024\_11 - log 2 scaled



RNA-seq intron features, aggregate (filtered), NCBI RefSeq Annotation GCF\_964237555.1-RS\_2024\_11

