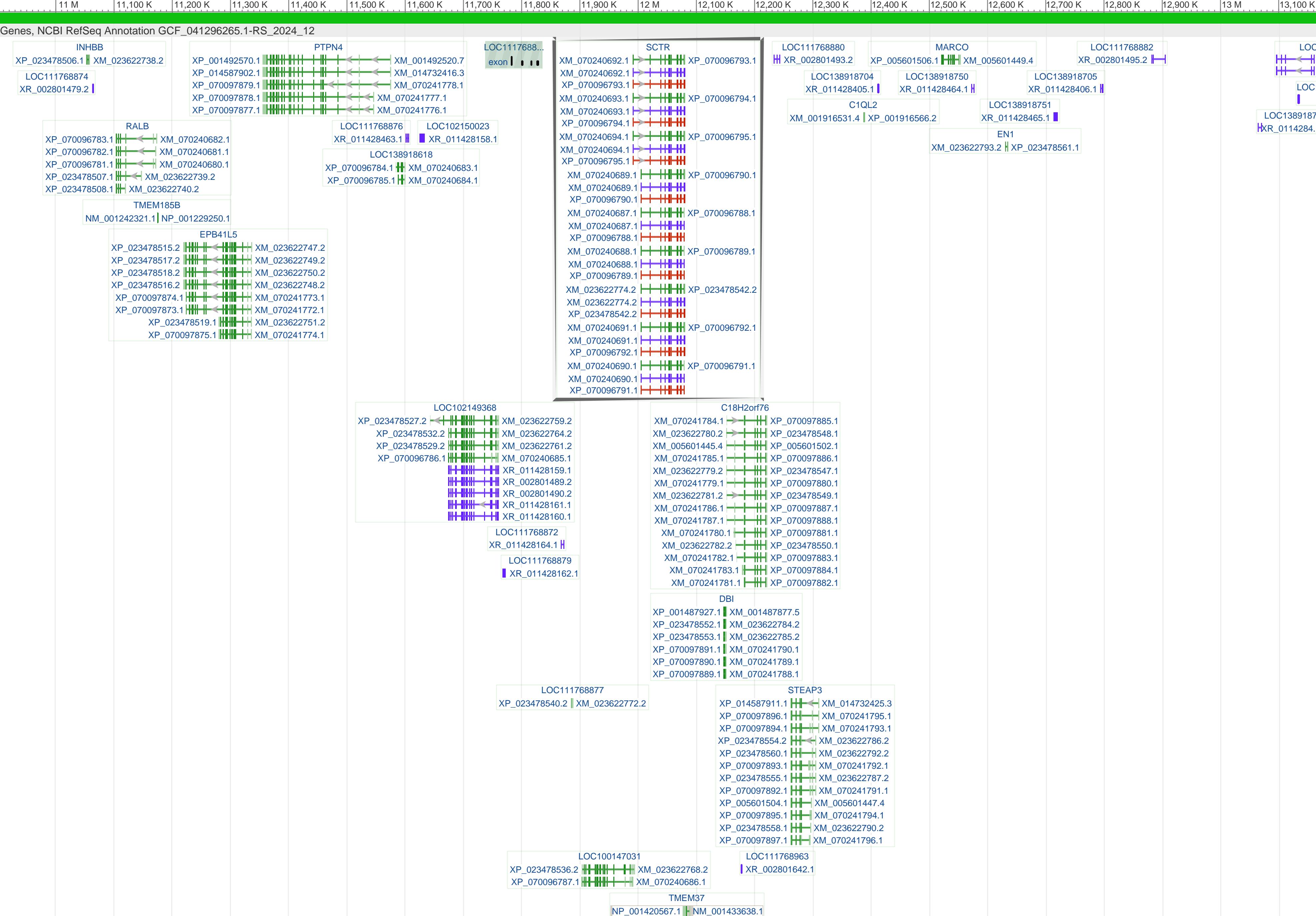


NC\_091701.1:10904146..13165606 Equus caballus isolate H\_3958 breed thoroughbred chromosome 18, TB-T2T, whole genome shotgun sequence



RNA-seq exon coverage, aggregate (filtered), NCBI RefSeq Annotation GCF\_041296265.1-RS\_2024\_12 - log 2 scaled



RNA-seq intron-spanning reads, aggregate (filtered), NCBI RefSeq Annotation GCF\_041296265.1-RS\_2024\_12 - log 2 scale

