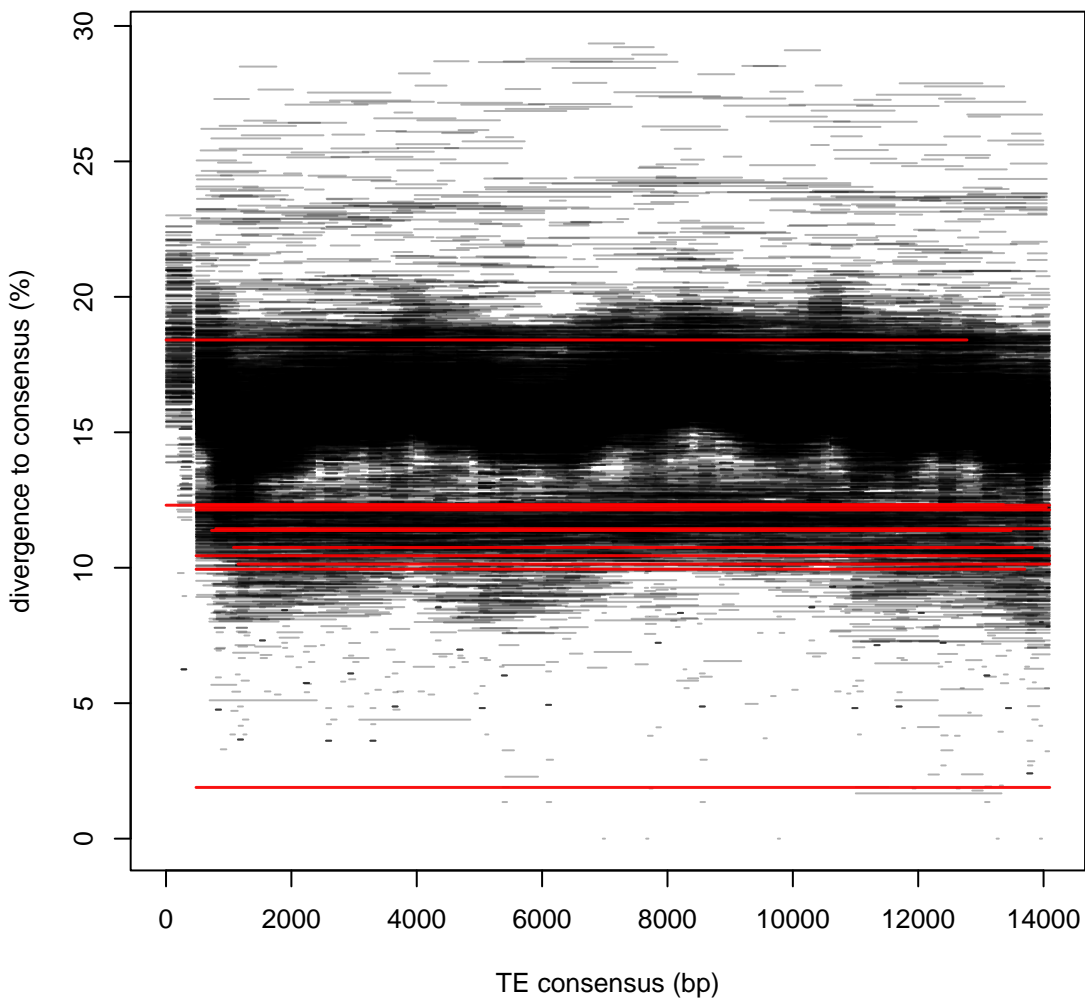
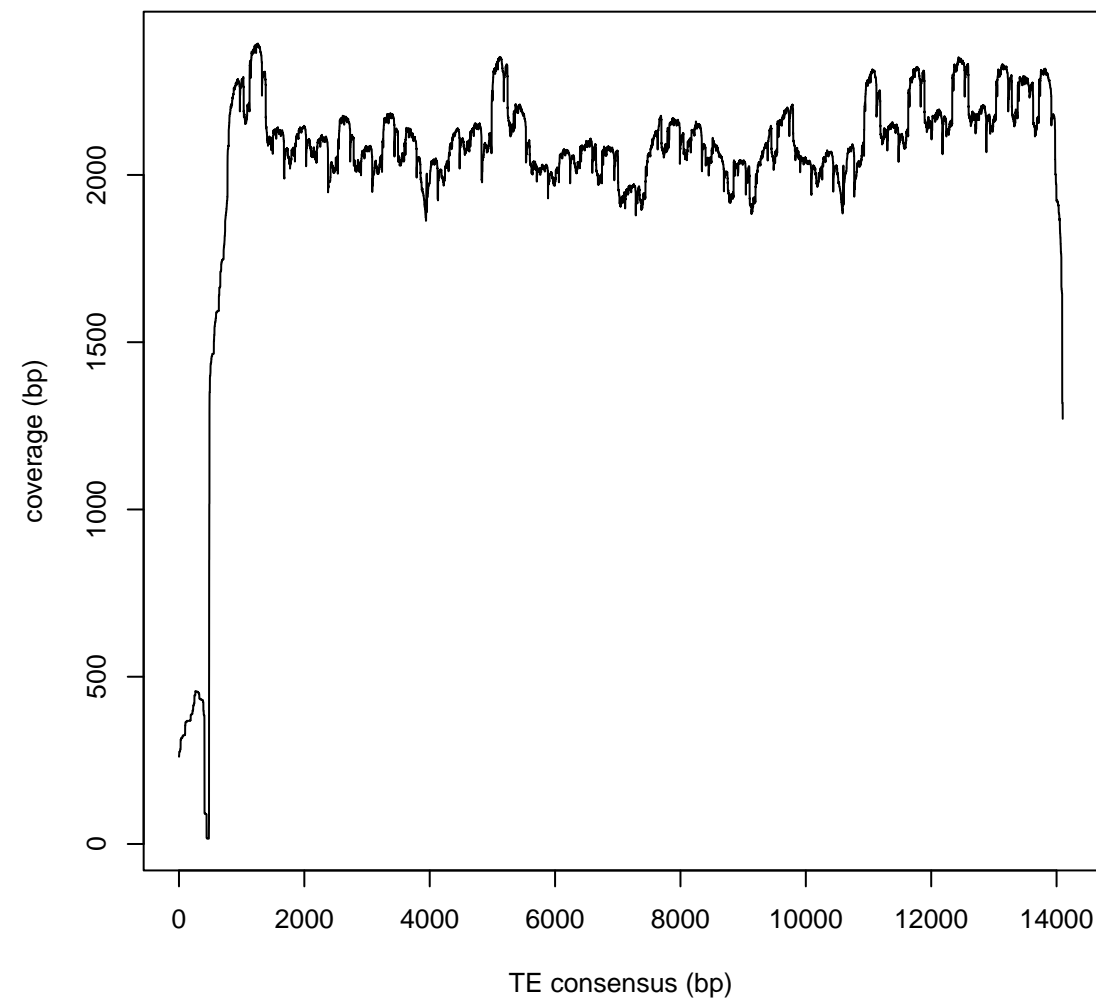


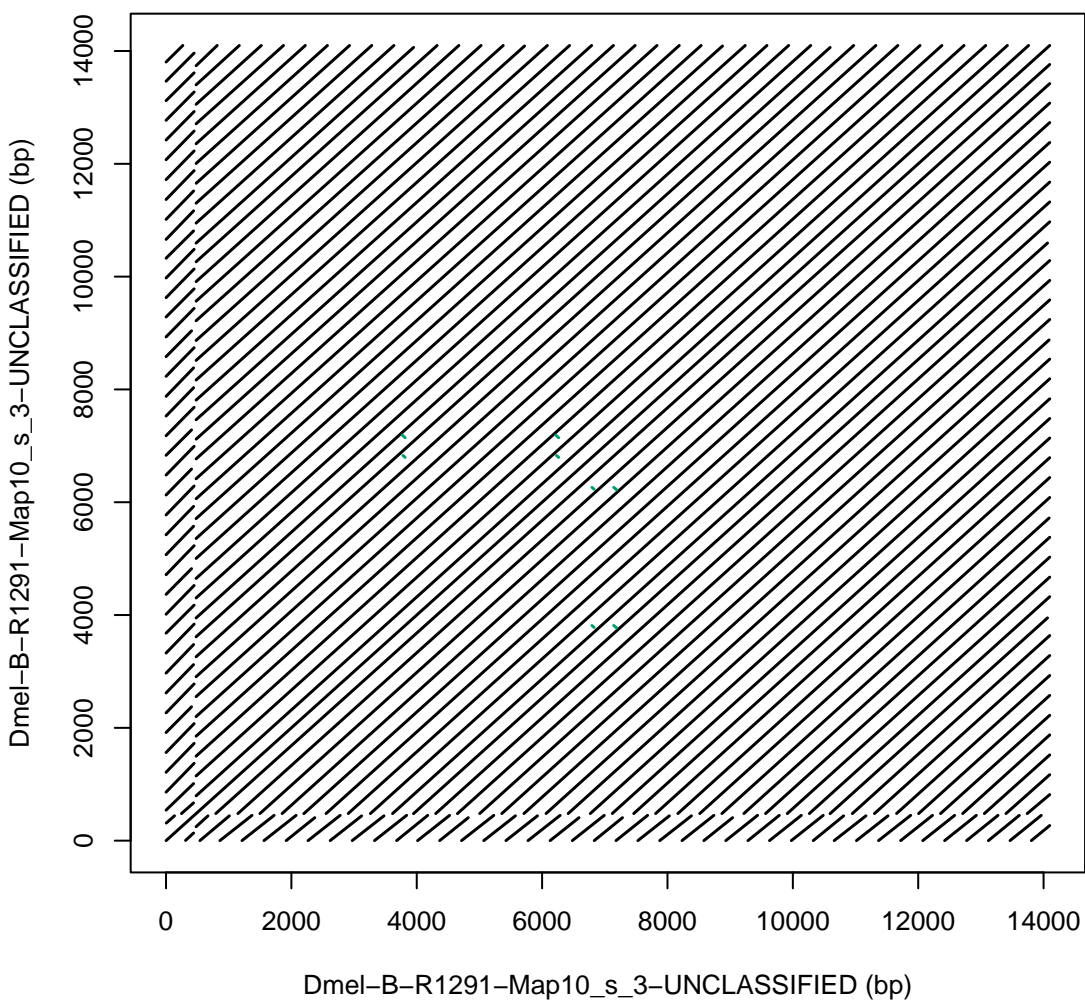
TE: Dmel-B-R1291-Map10_s_3-UNCLASSIFIED
consensus size: 14098bp; fragments: 26784; full length: 12 (≥ 12688.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

