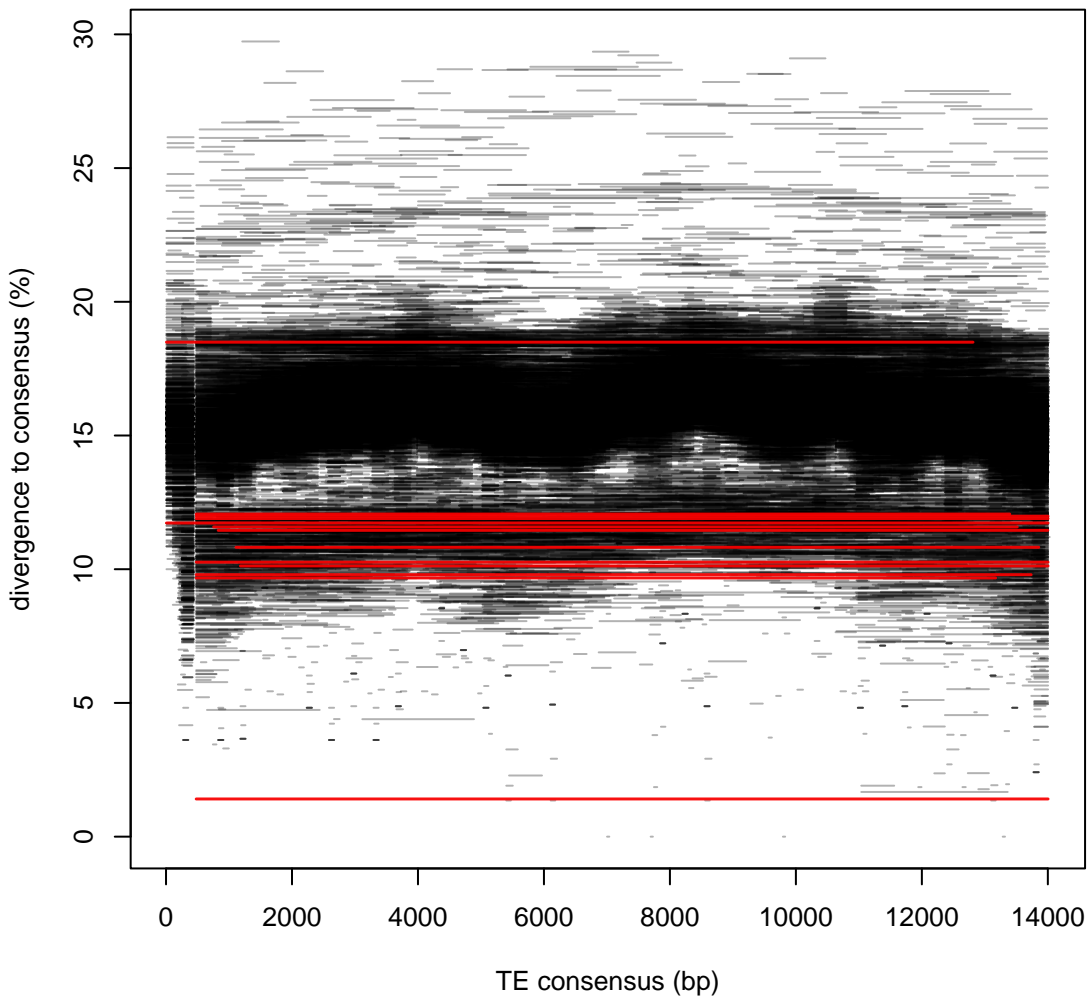
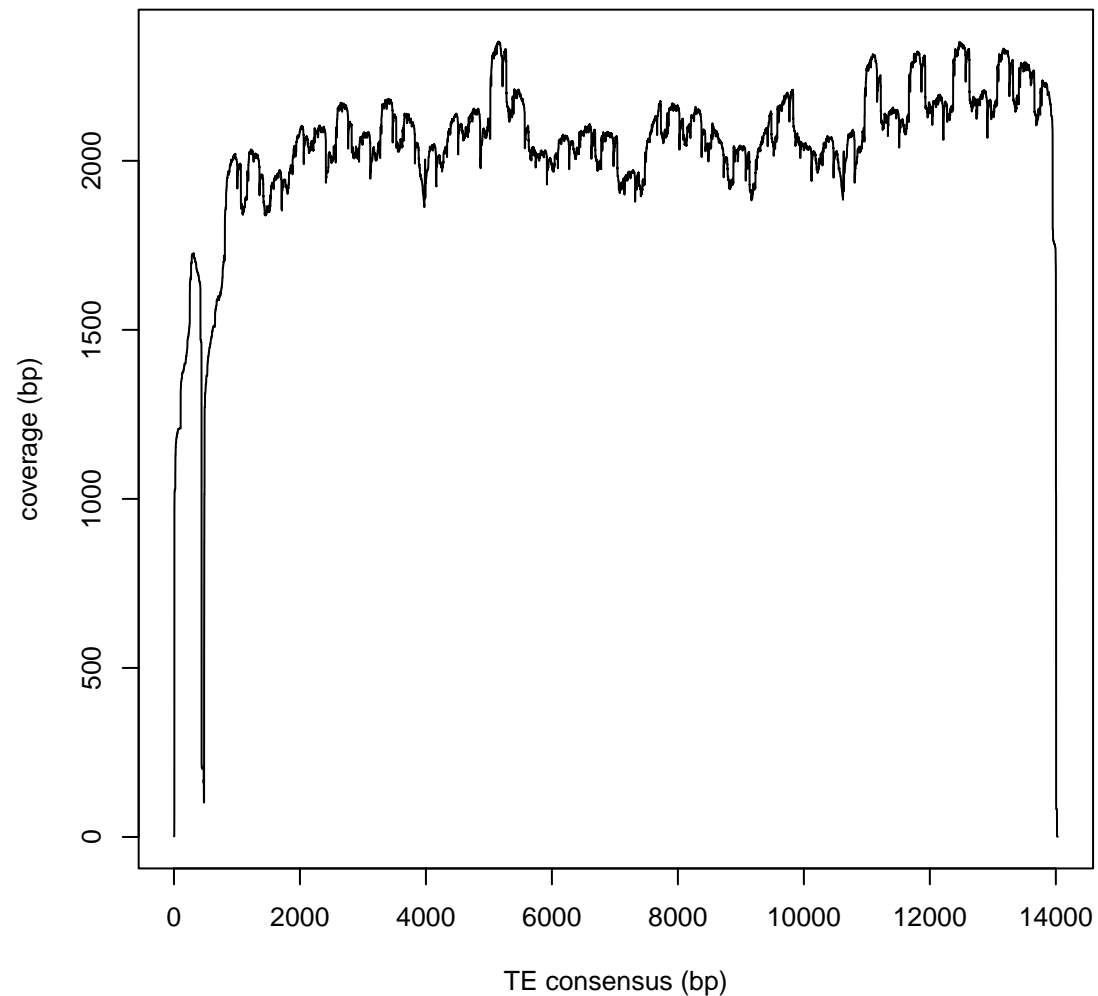


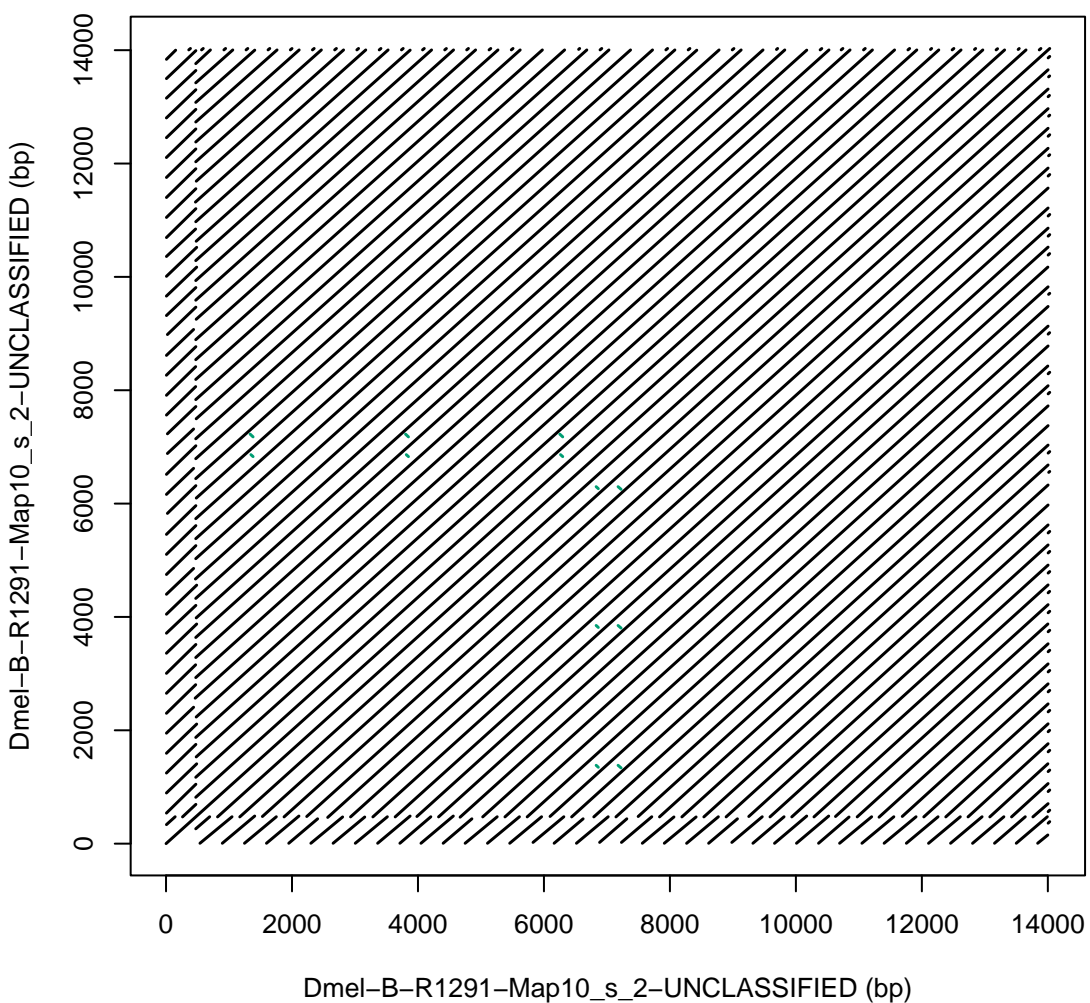
TE: Dmel-B-R1291-Map10_s_2-UNCLASSIFIED
consensus size: 14030bp; fragments: 27440; full length: 13 (>=12627bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

