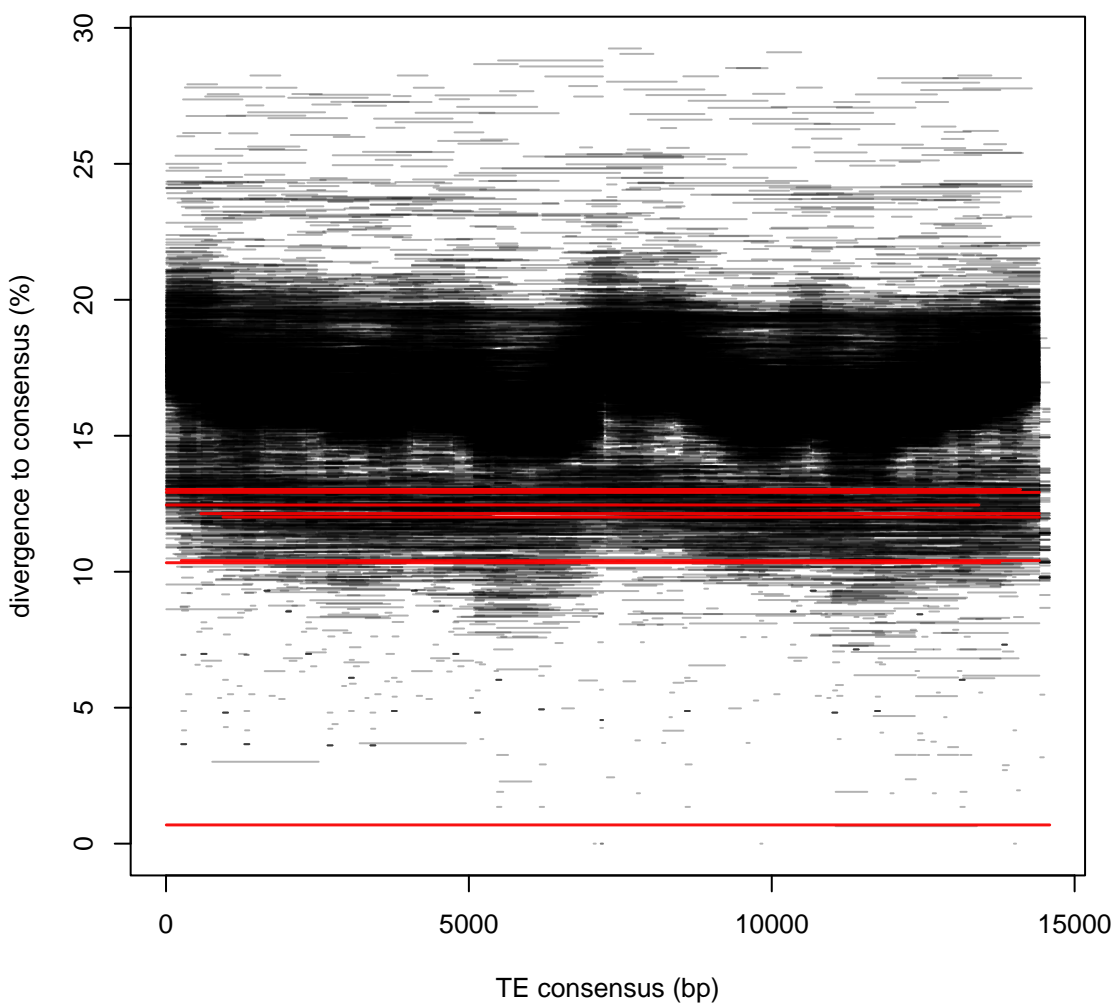
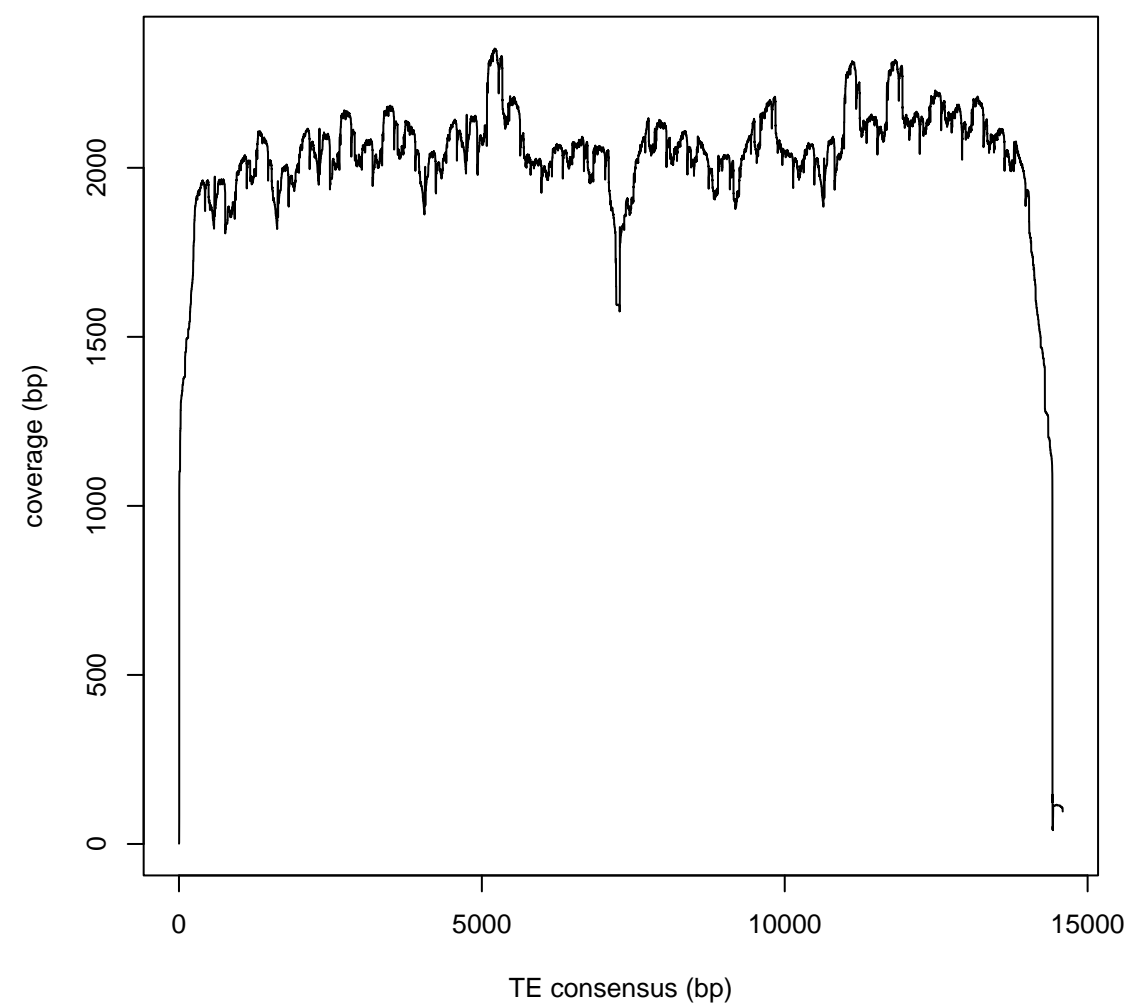


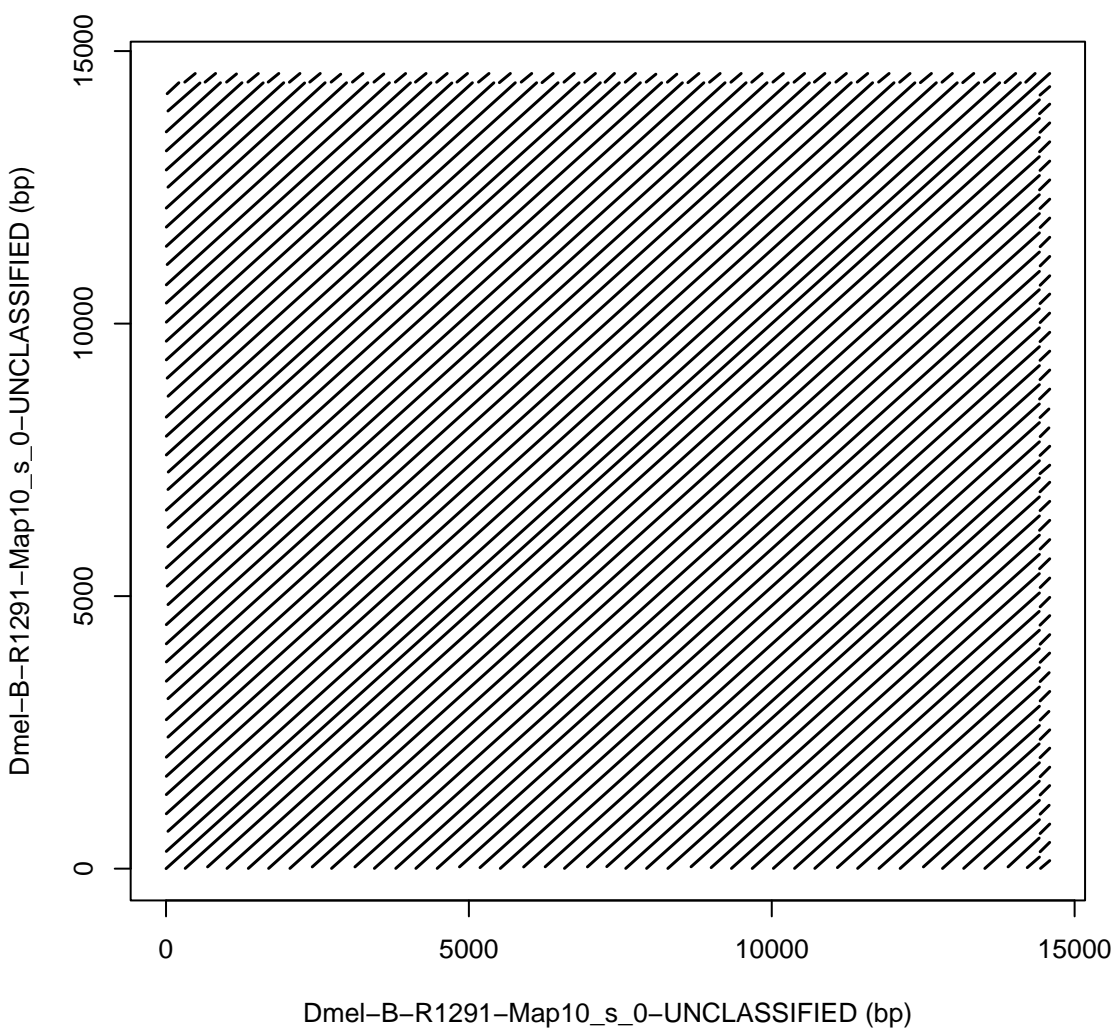
TE: Dmel-B-R1291-Map10_s_0-UNCLASSIFIED
consensus size: 14589bp; fragments: 26213; full length: 8 (≥ 13130.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

