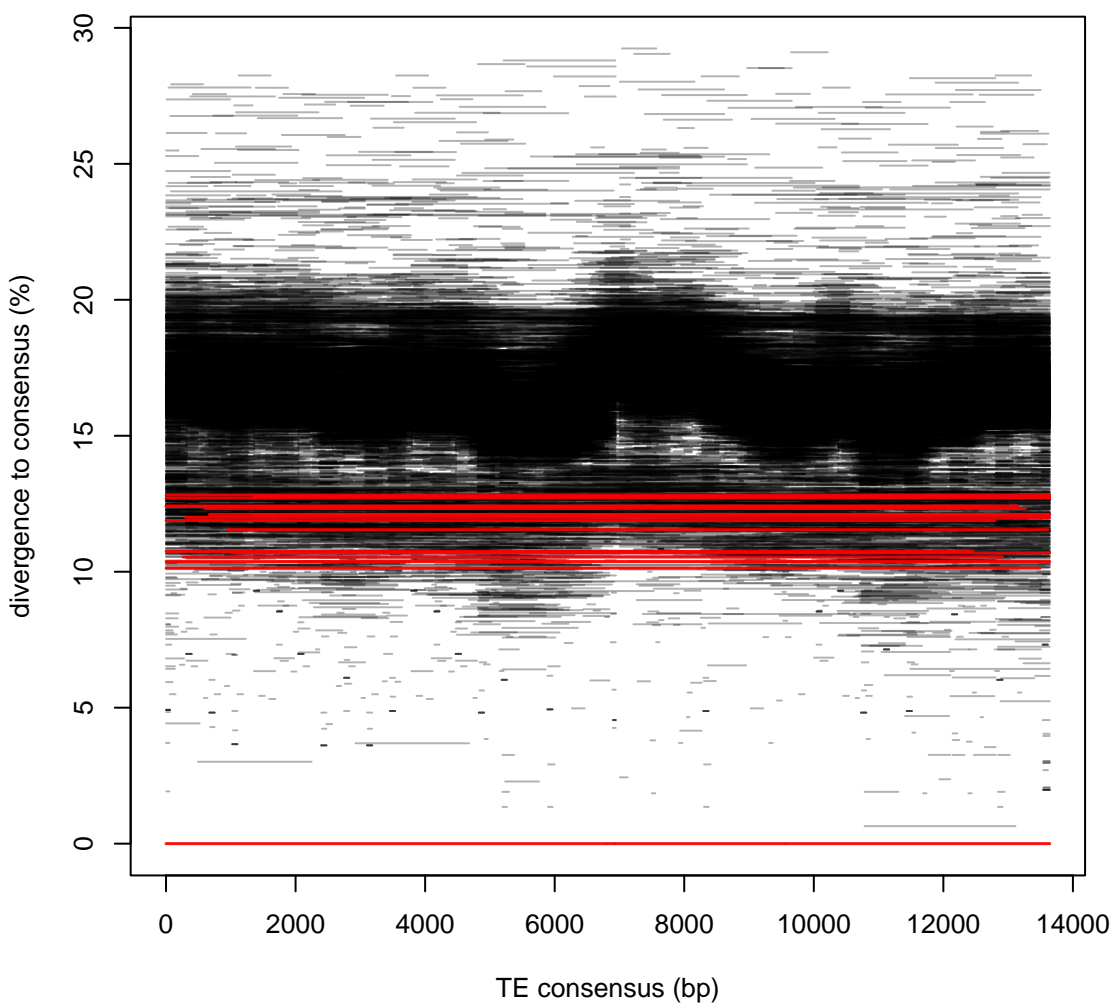
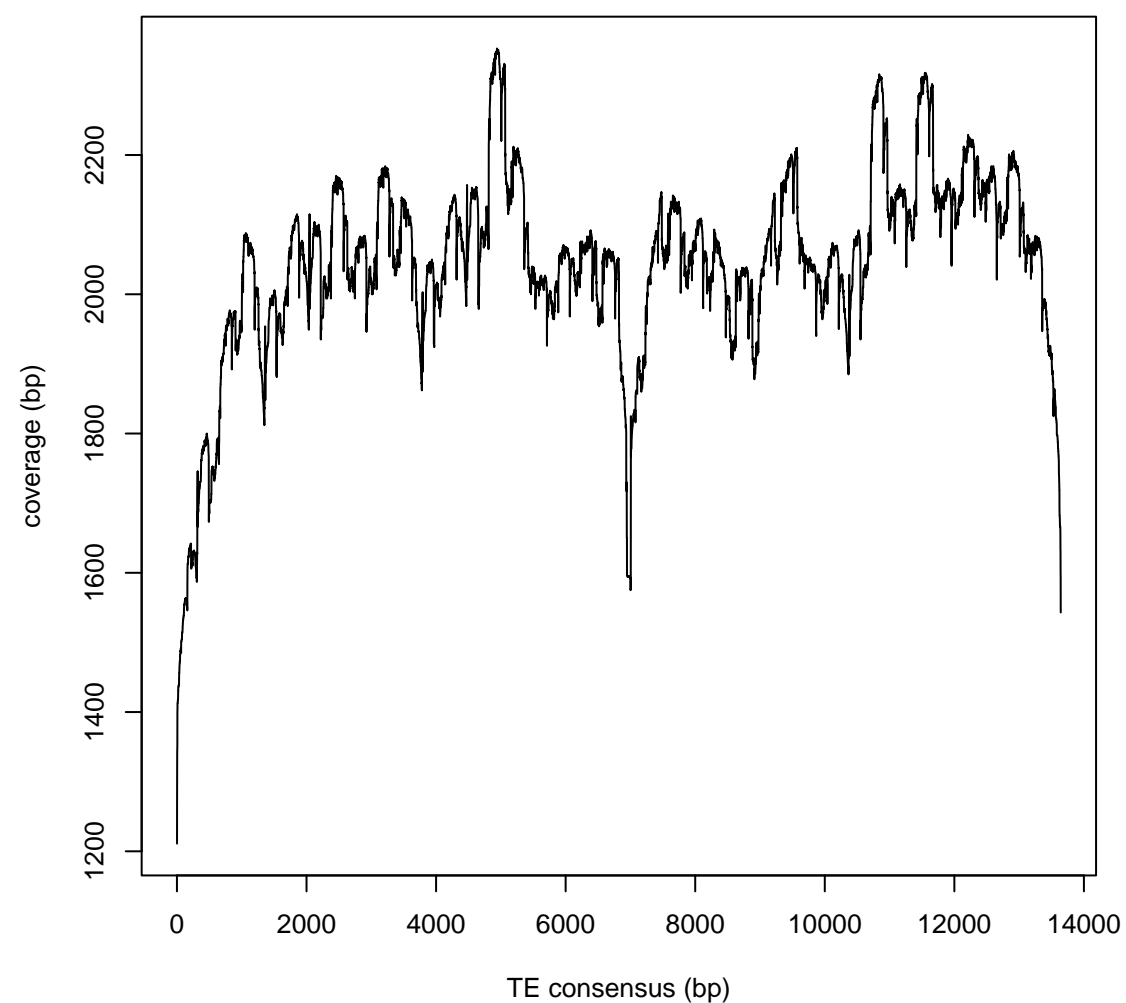


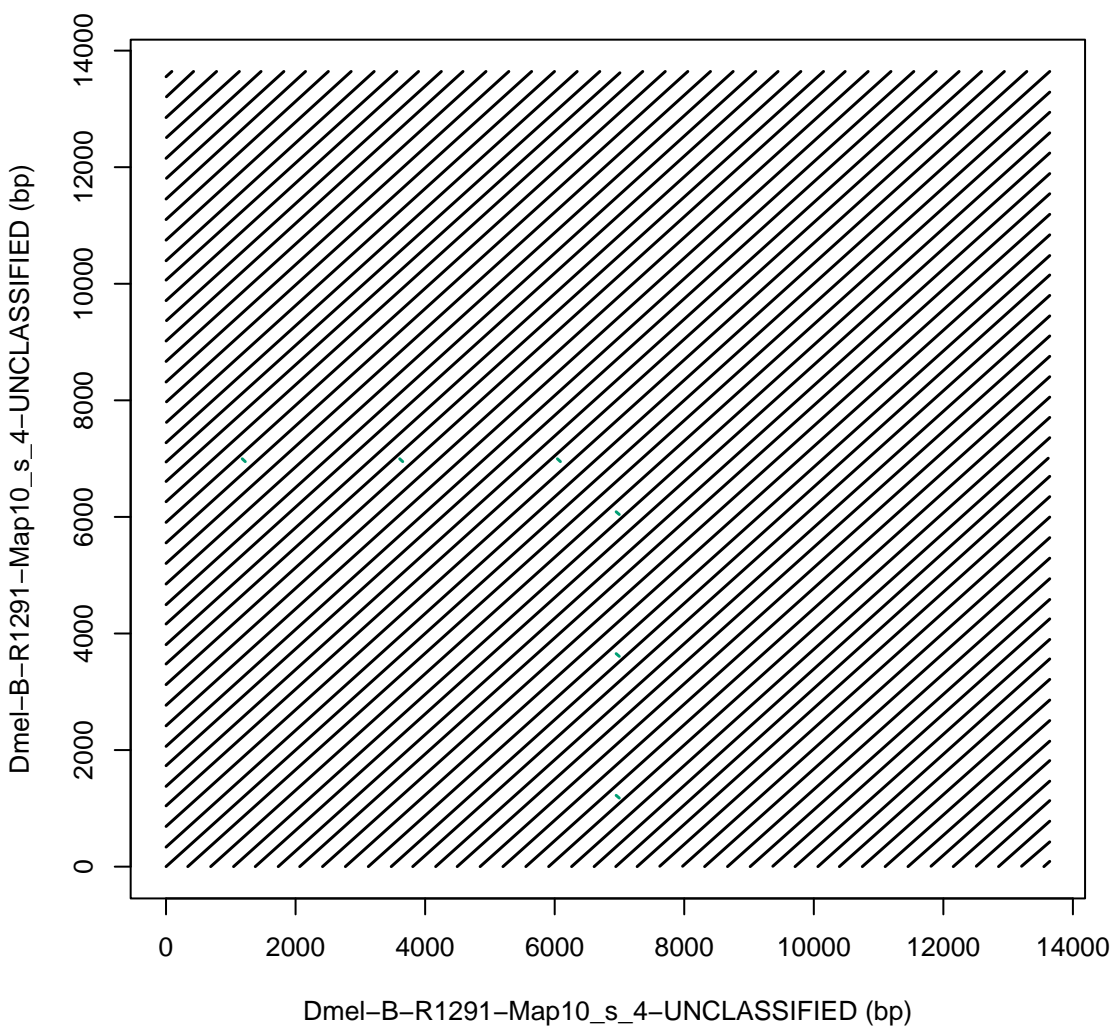
TE: Dmel-B-R1291-Map10_s_4-UNCLASSIFIED
consensus size: 13642bp; fragments: 25214; full length: 17 (≥ 12277.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

