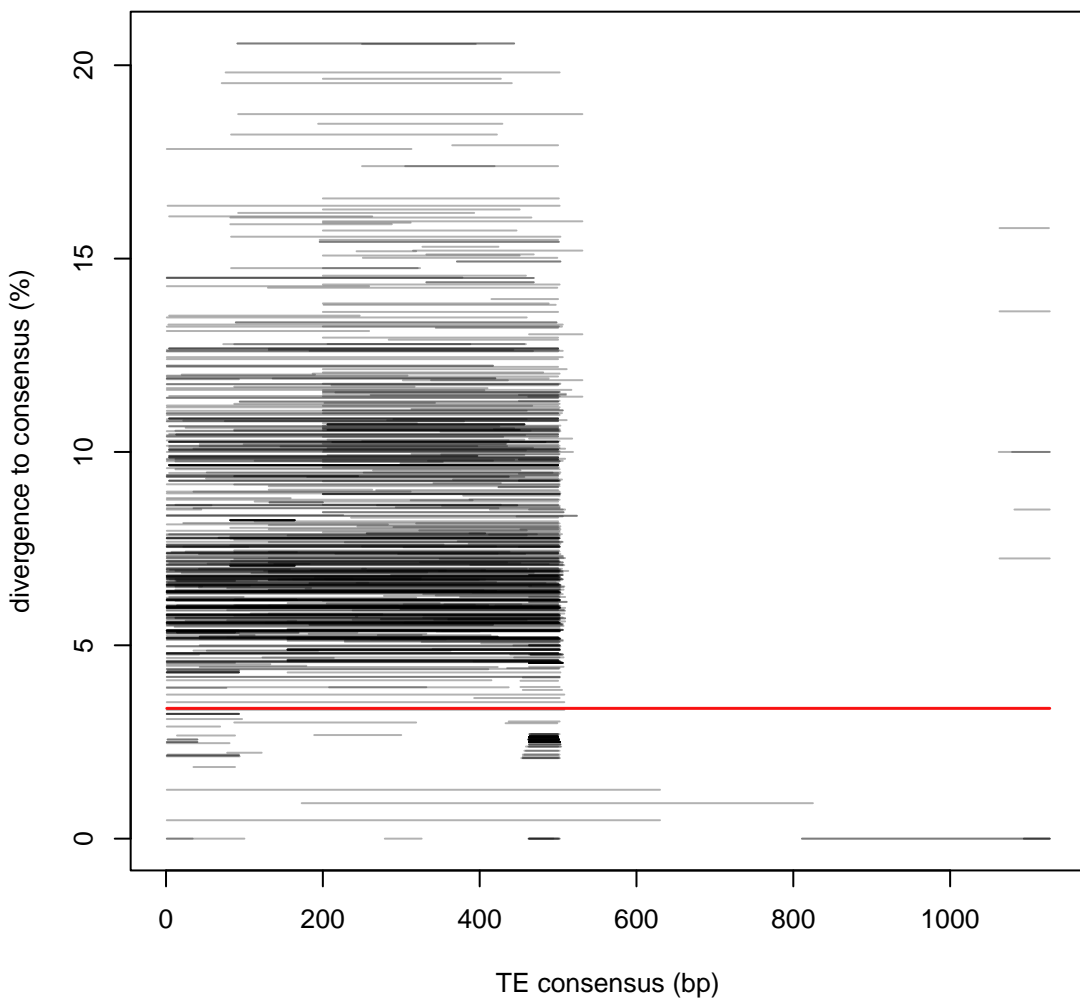
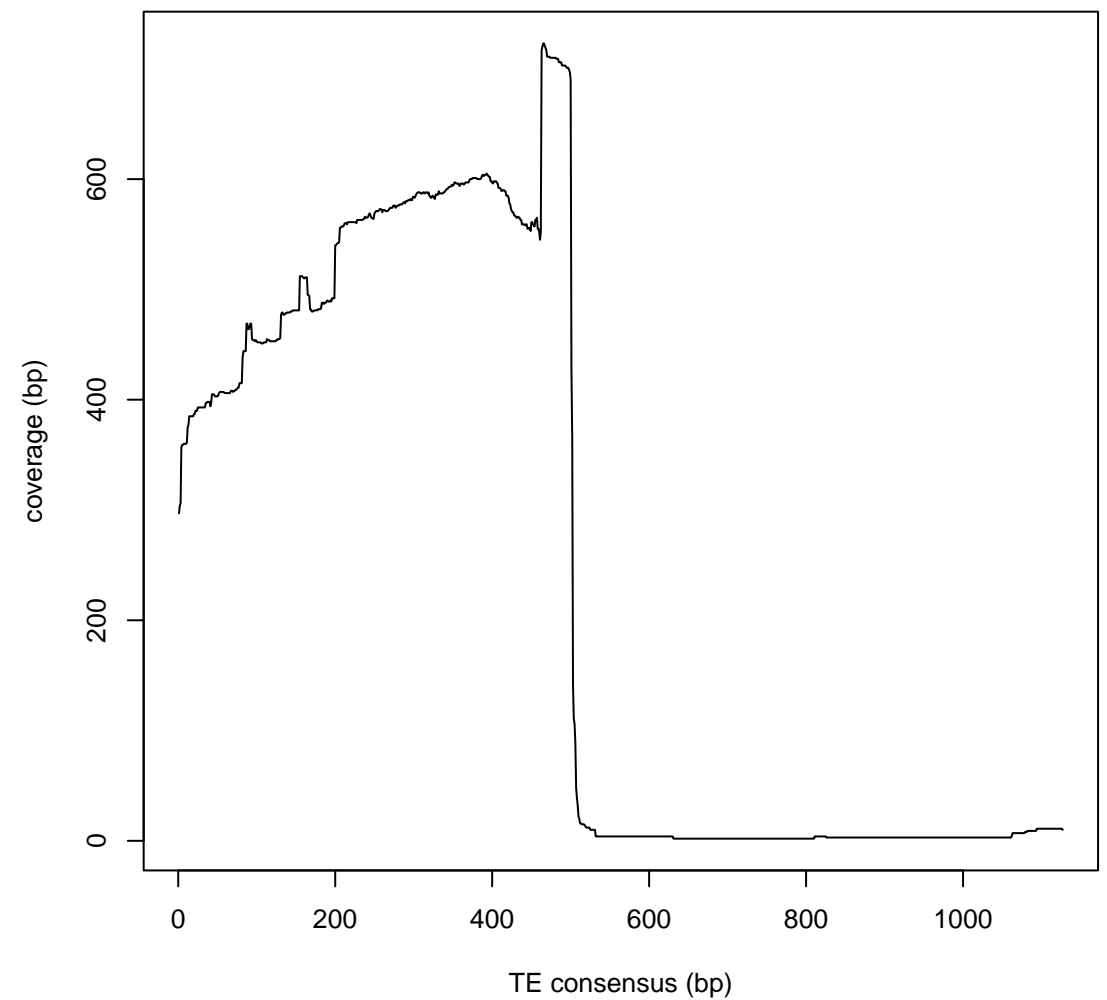


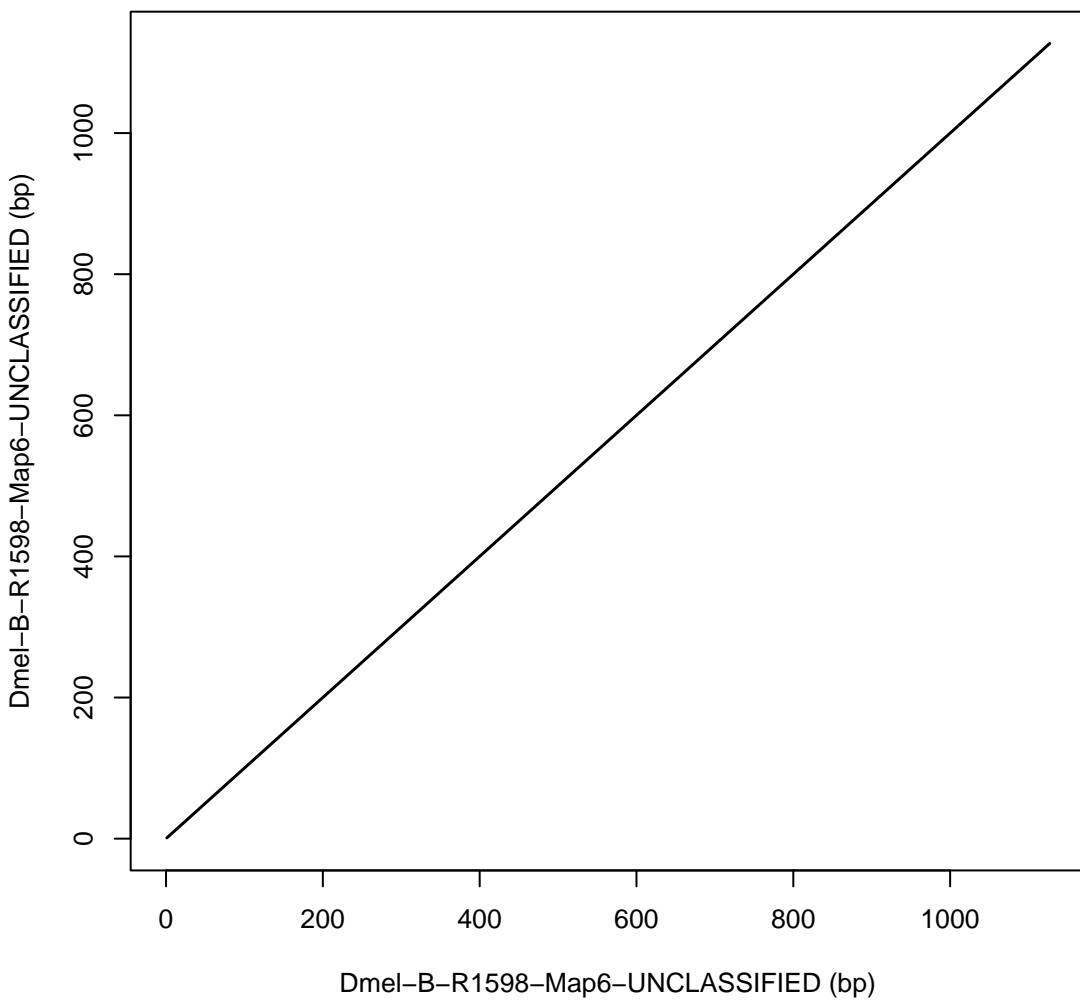
TE: Dmel-B-R1598-Map6-UNCLASSIFIED
consensus size: 1127bp; fragments: 972; full length: 1 (≥ 1014.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

