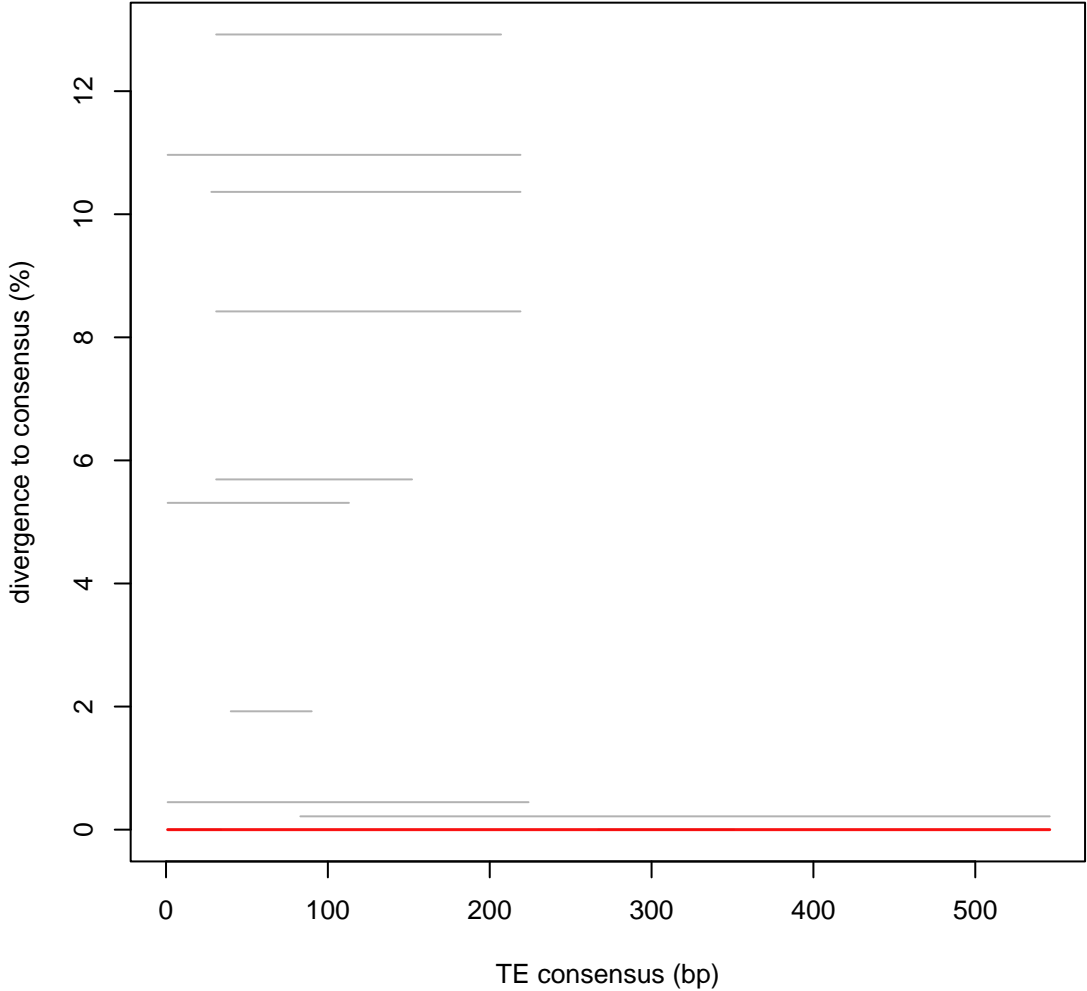
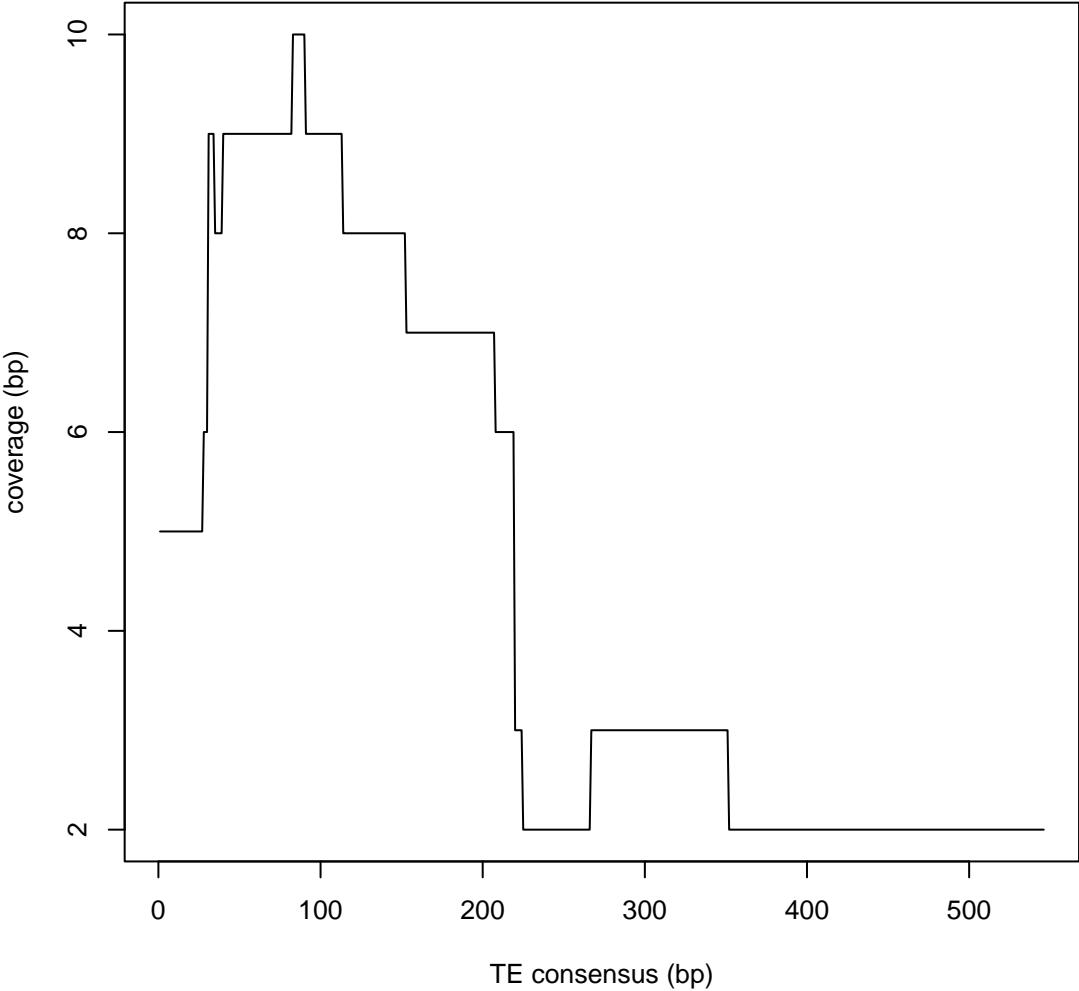


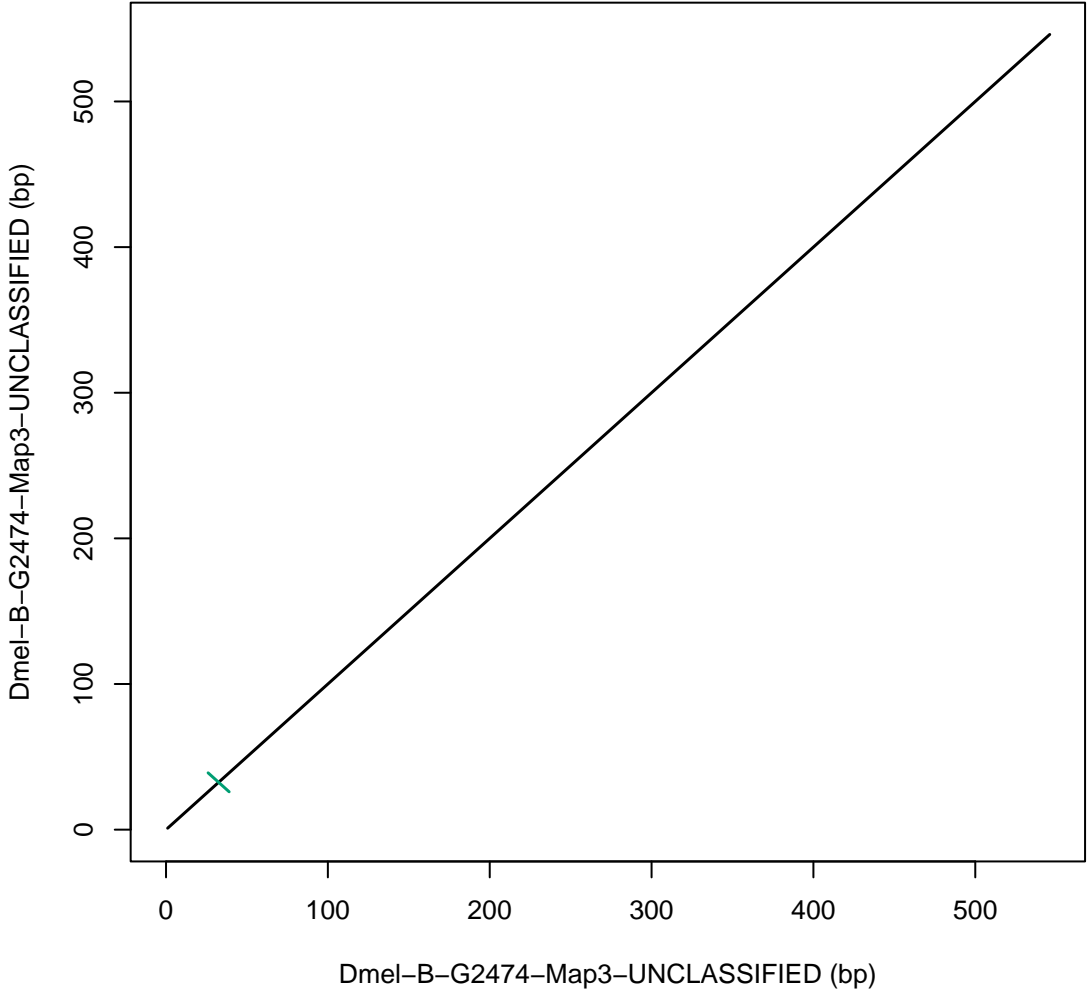
TE: Dmel-B-G2474-Map3-UNCLASSIFIED
consensus size: 546bp; fragments: 12; full length: 1 (>=491.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

