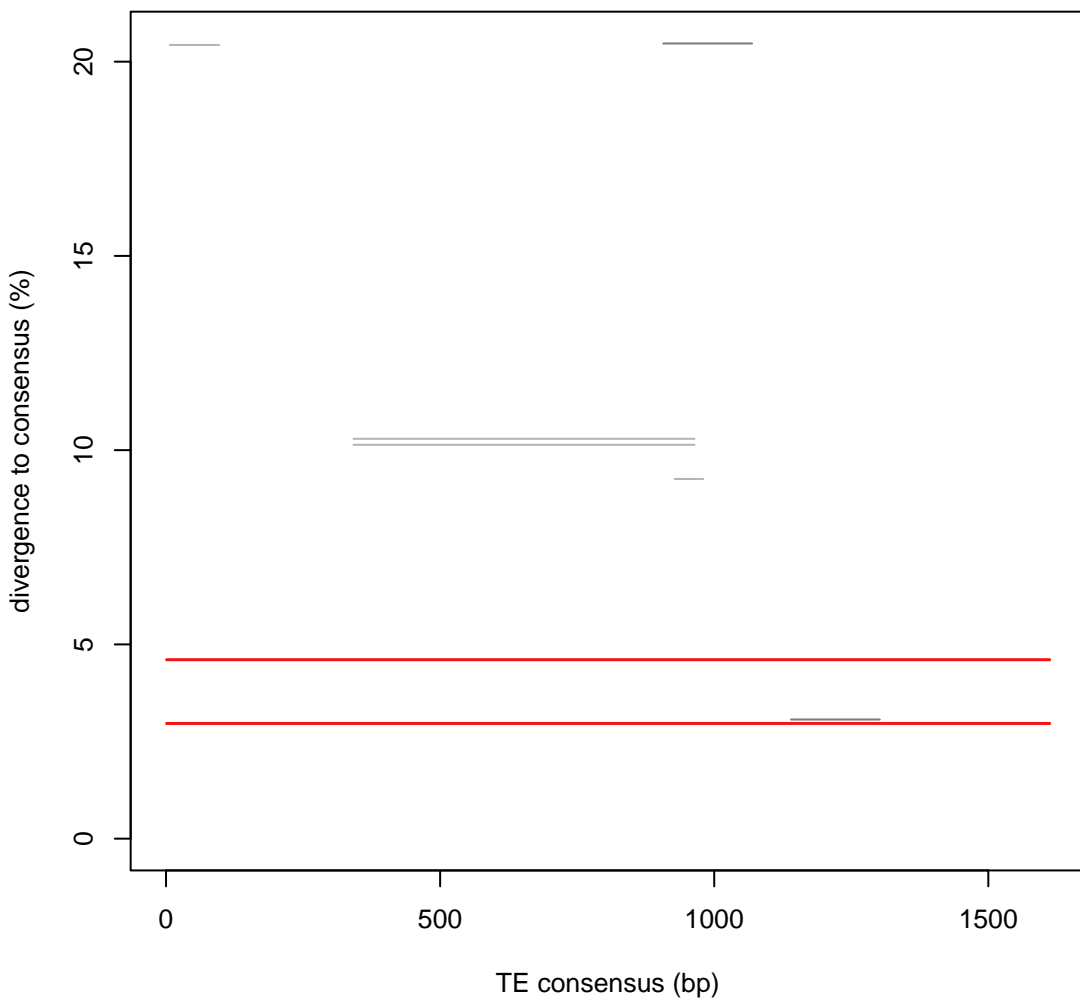
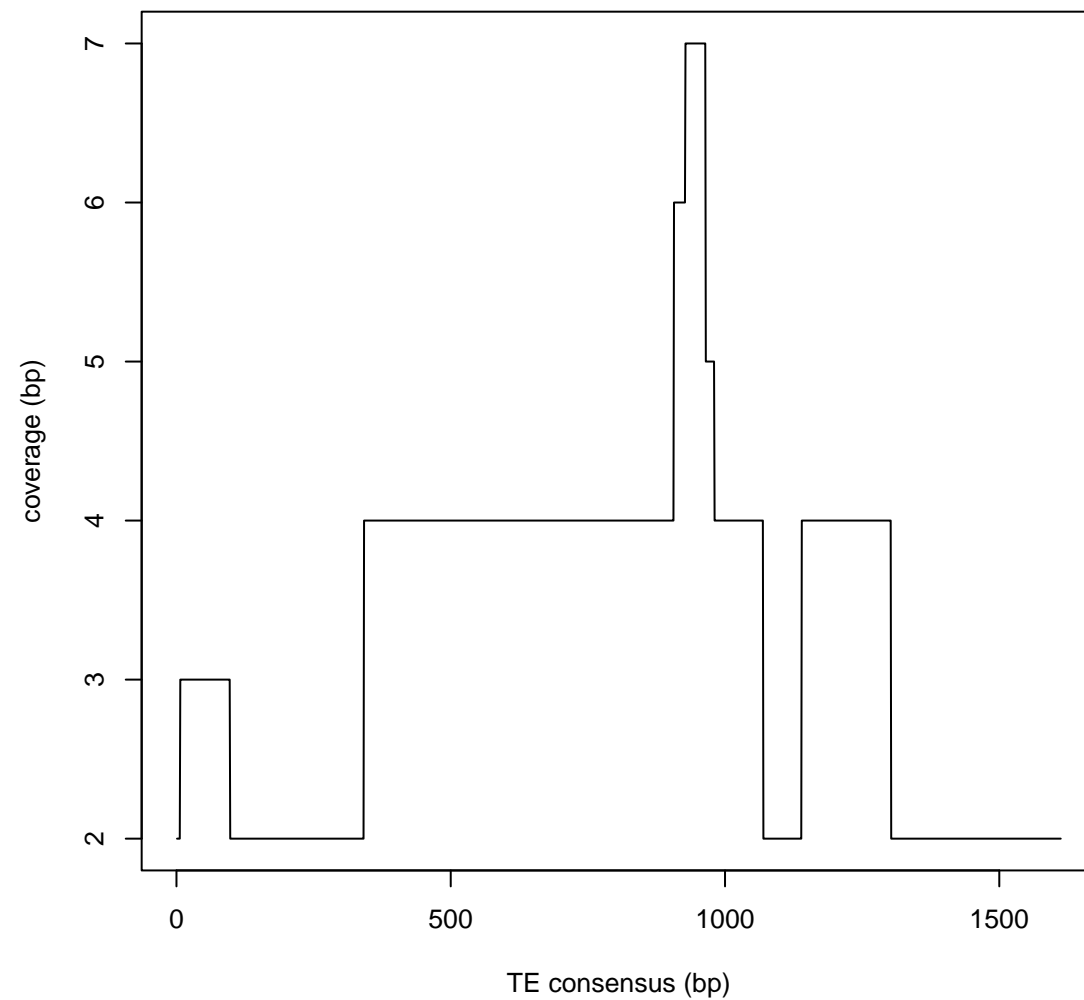


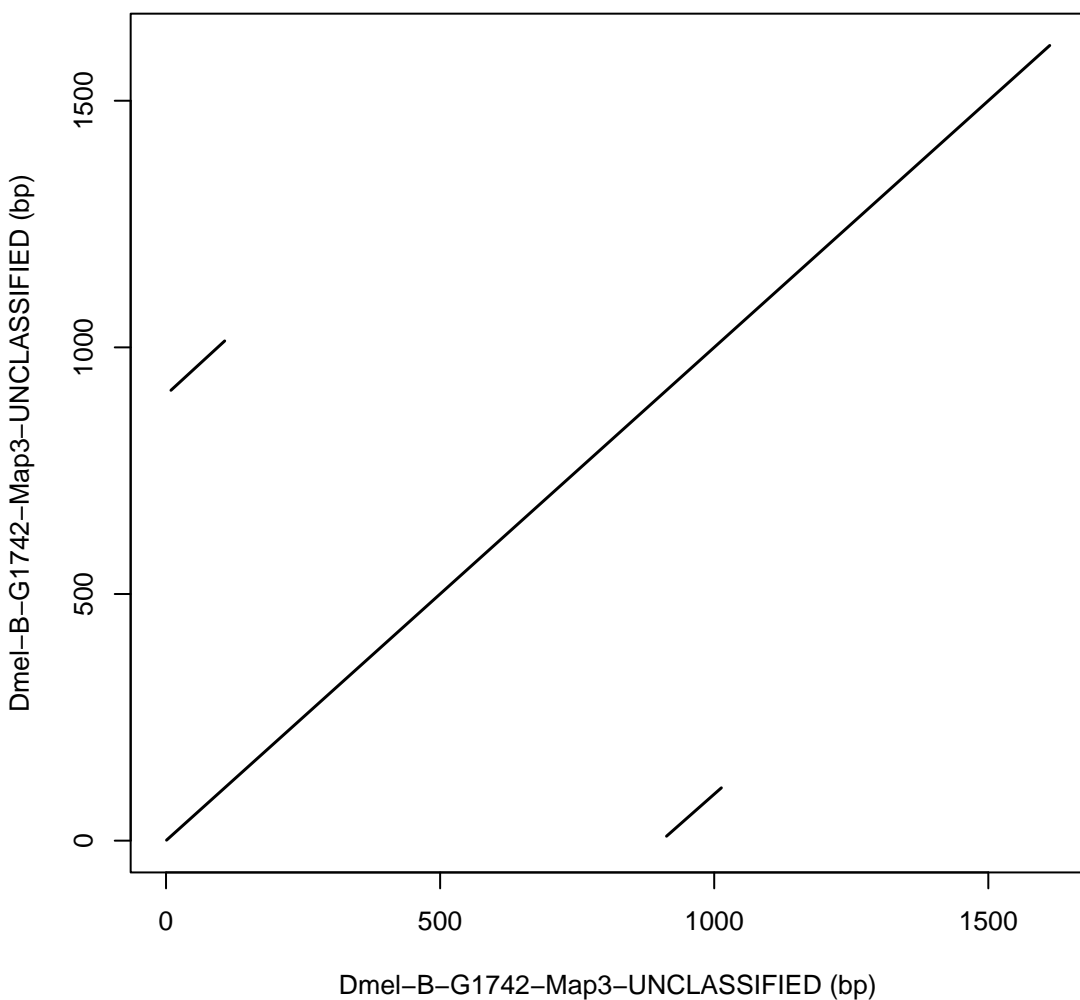
TE: Dmel-B-G1742-Map3-UNCLASSIFIED
consensus size: 1612bp; fragments: 10; full length: 2 (≥ 1450.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

