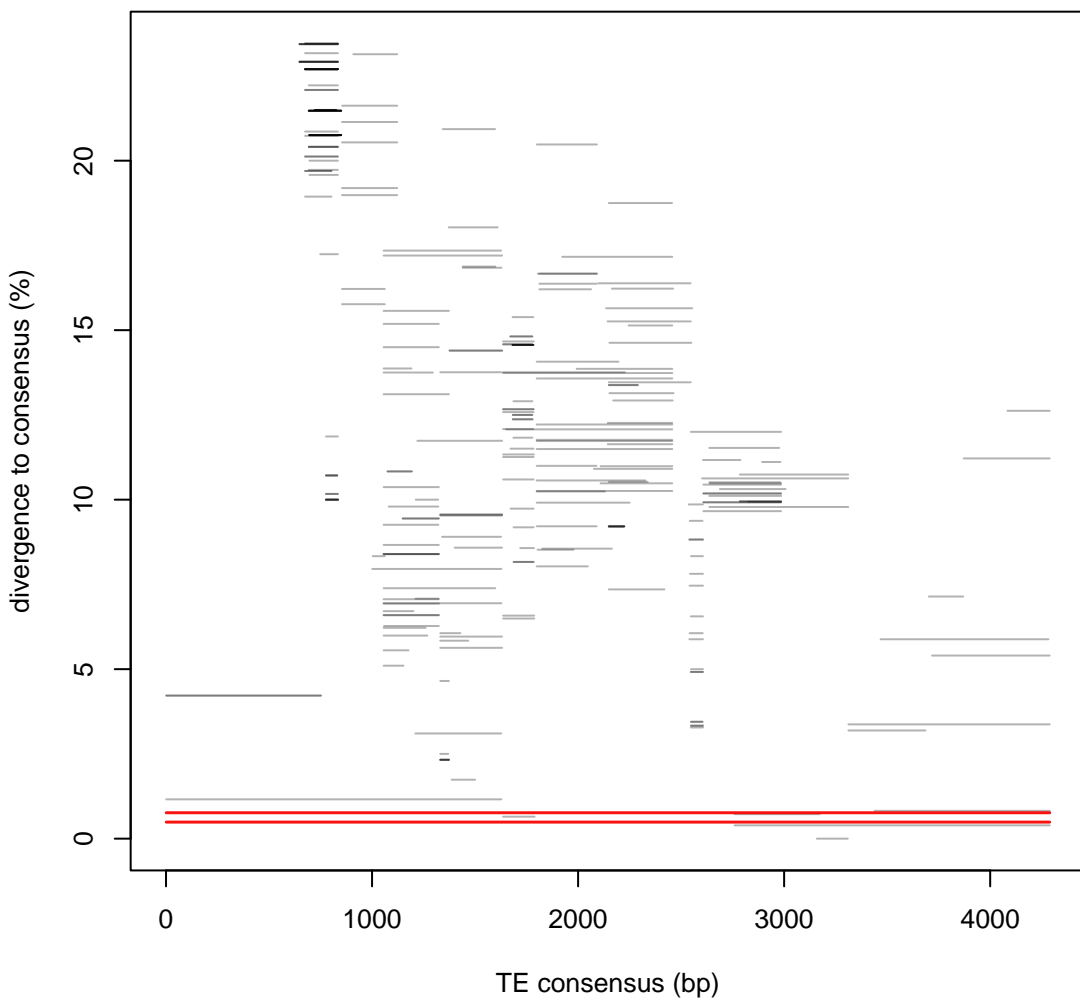
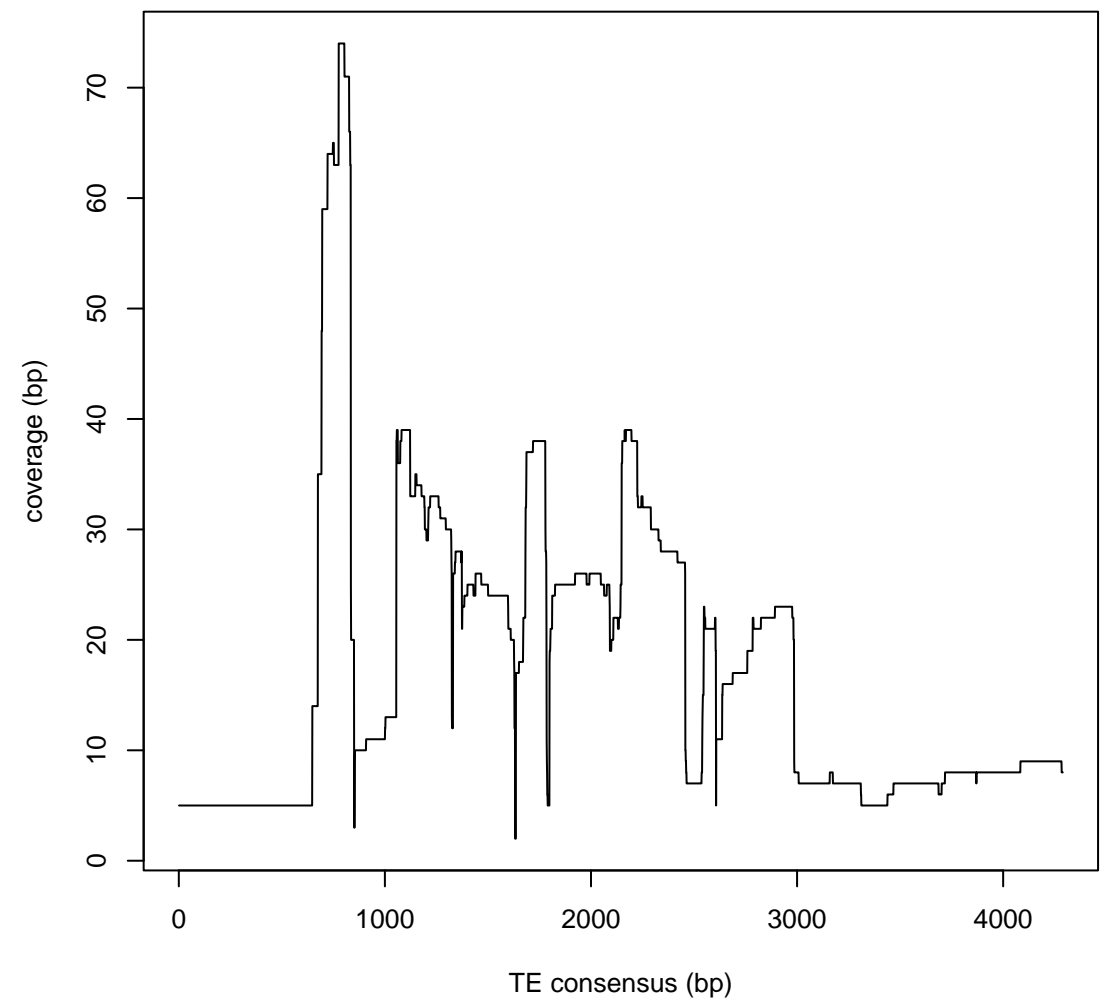


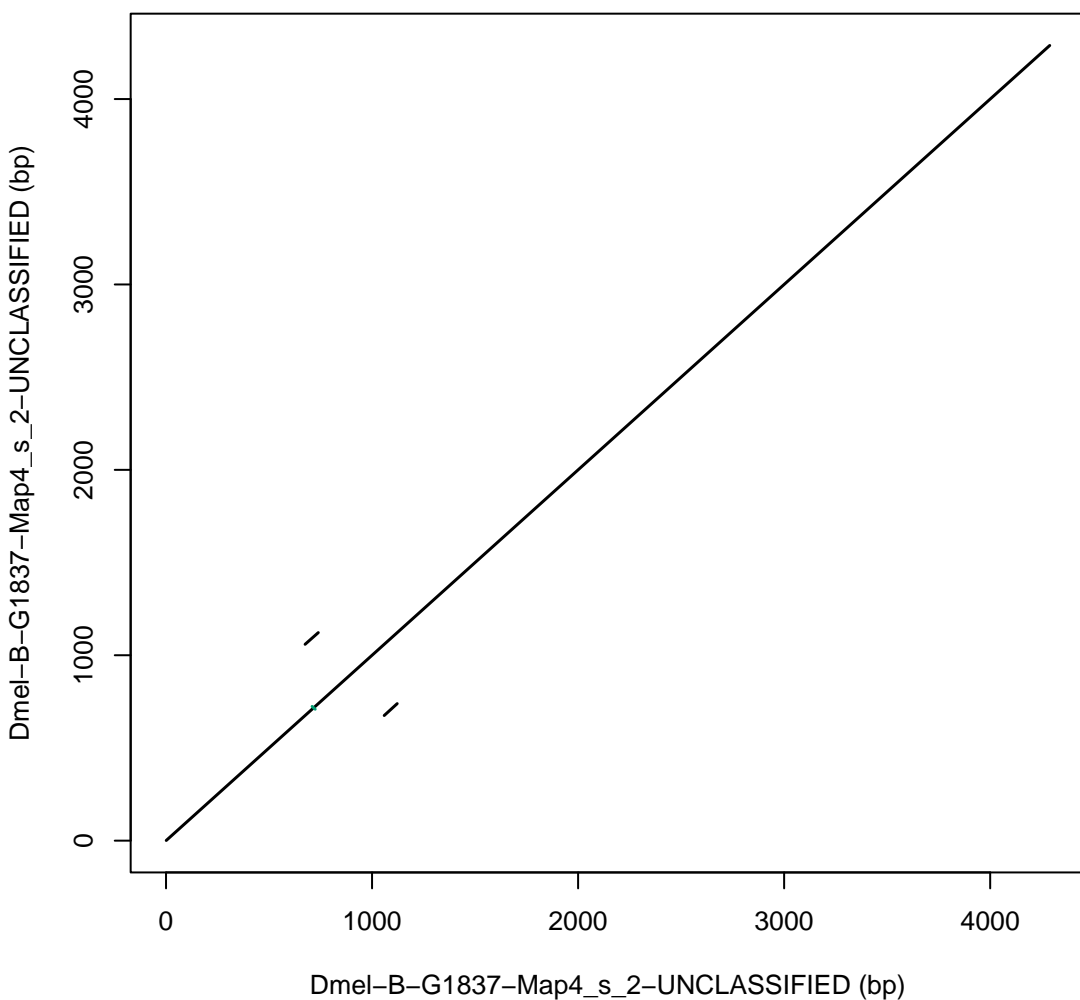
TE: Dmel-B-G1837-Map4_s_2-UNCLASSIFIED
consensus size: 4289bp; fragments: 275; full length: 2 (≥ 3860.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

