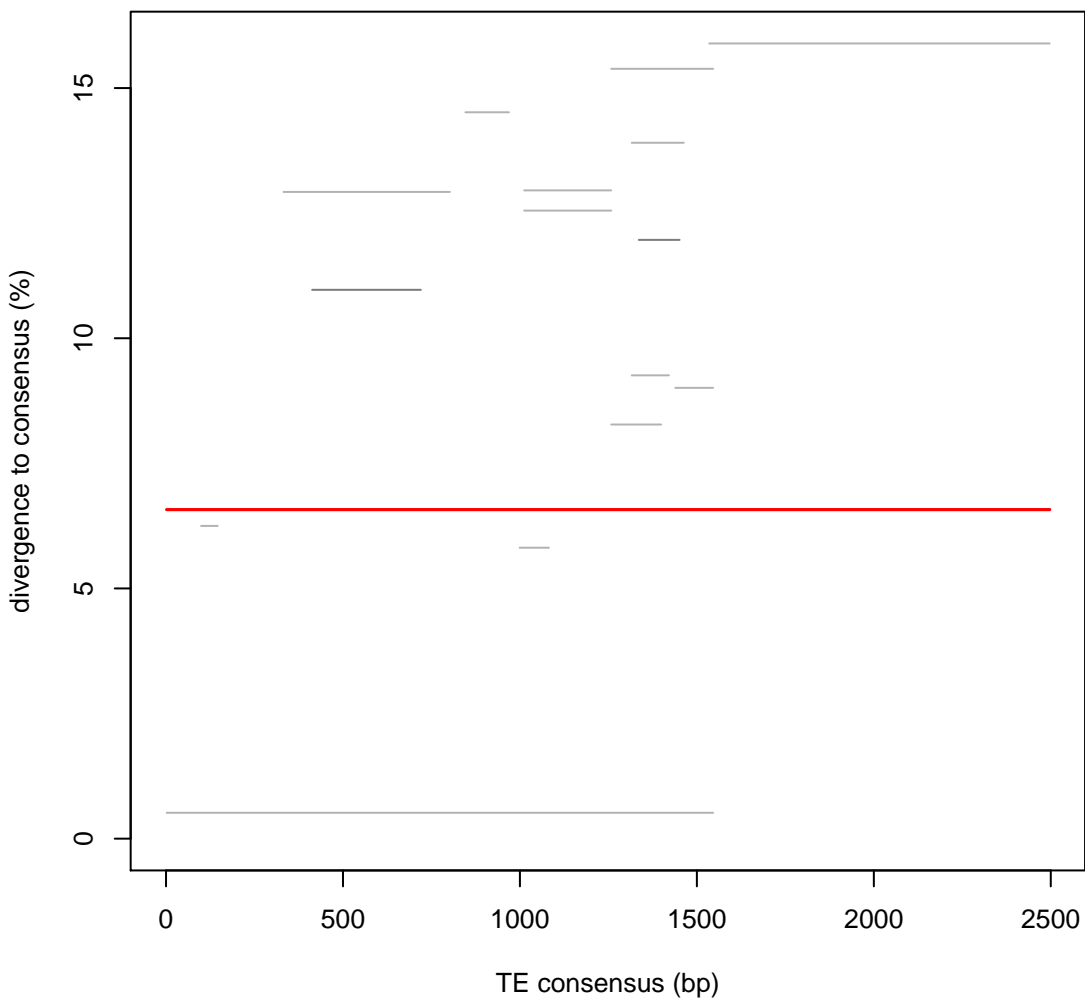
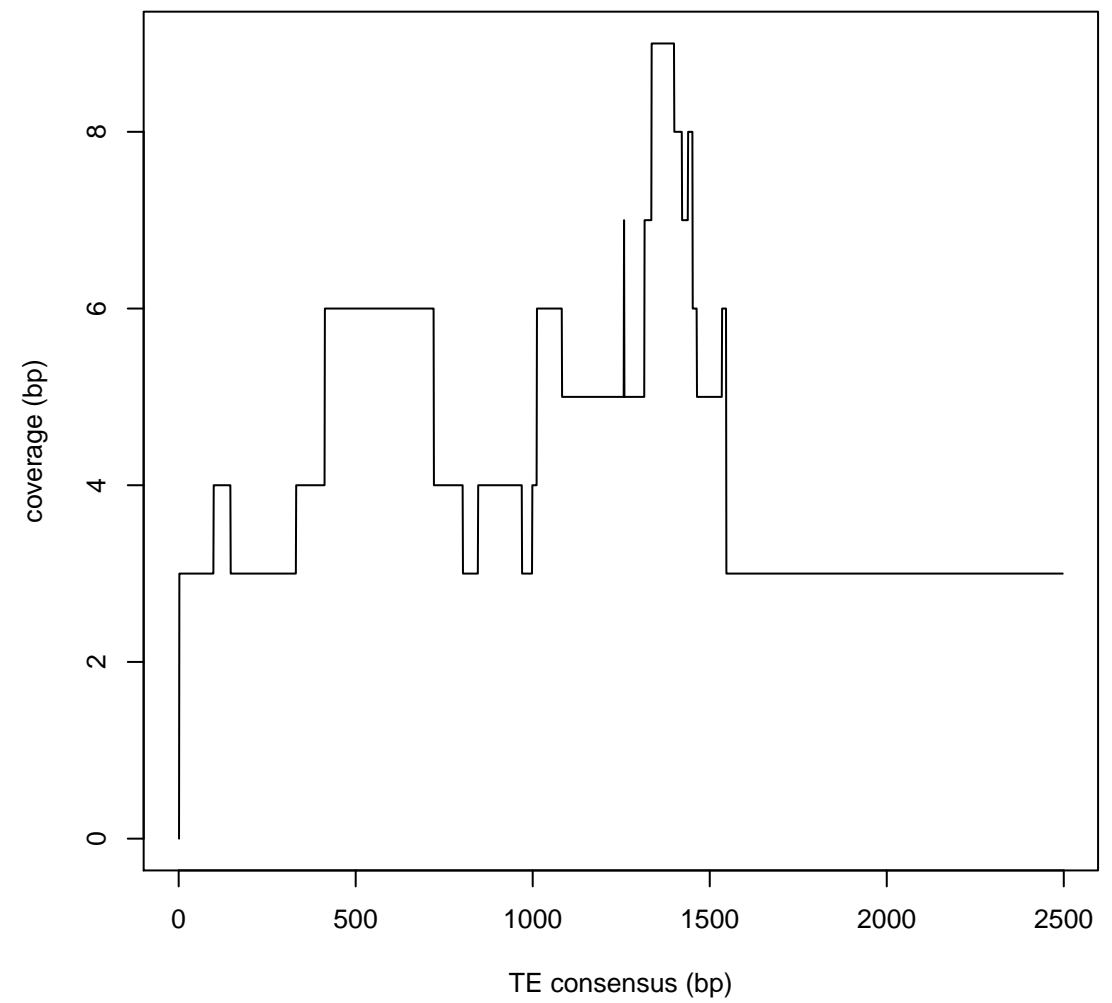


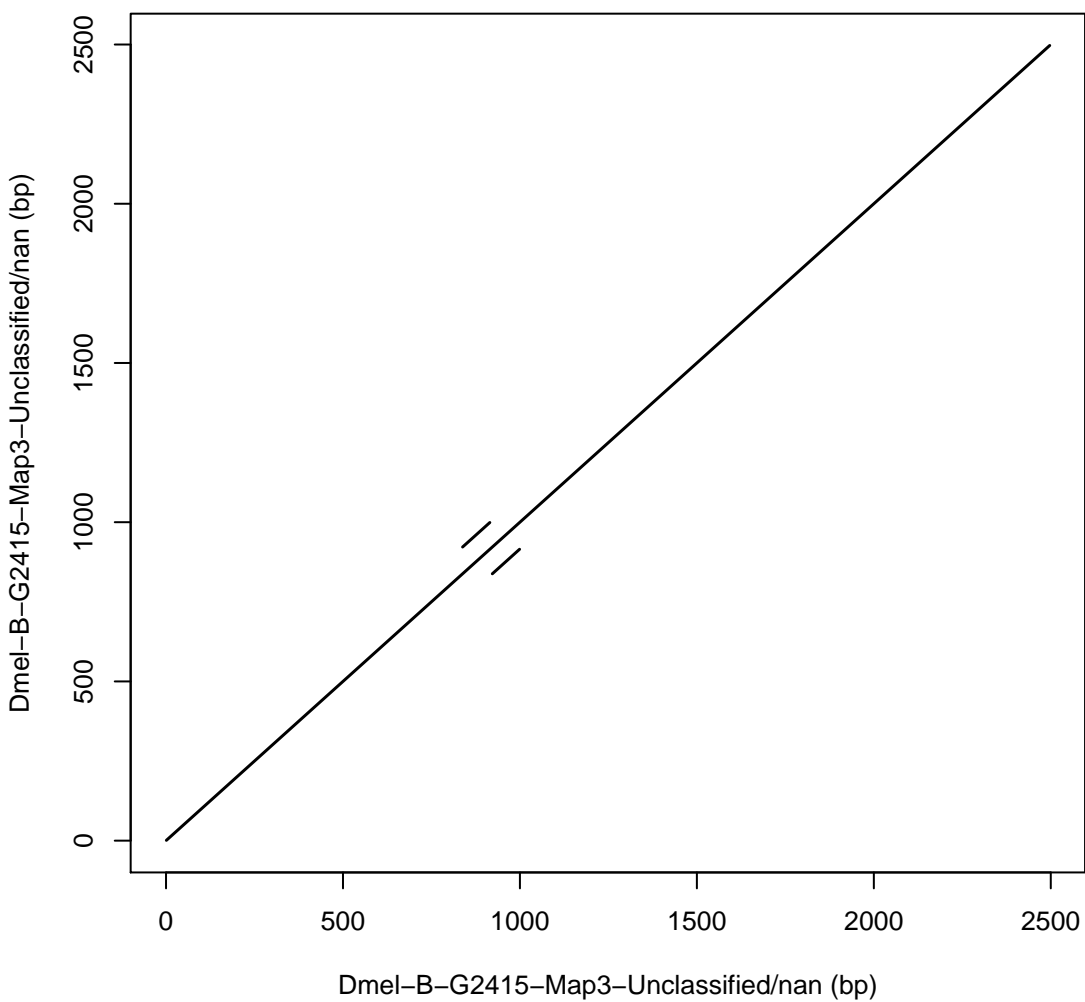
TE: Dmel-B-G2415-Map3-Unclassified/nan
consensus size: 2497bp; fragments: 19; full length: 2 (≥ 2247.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

