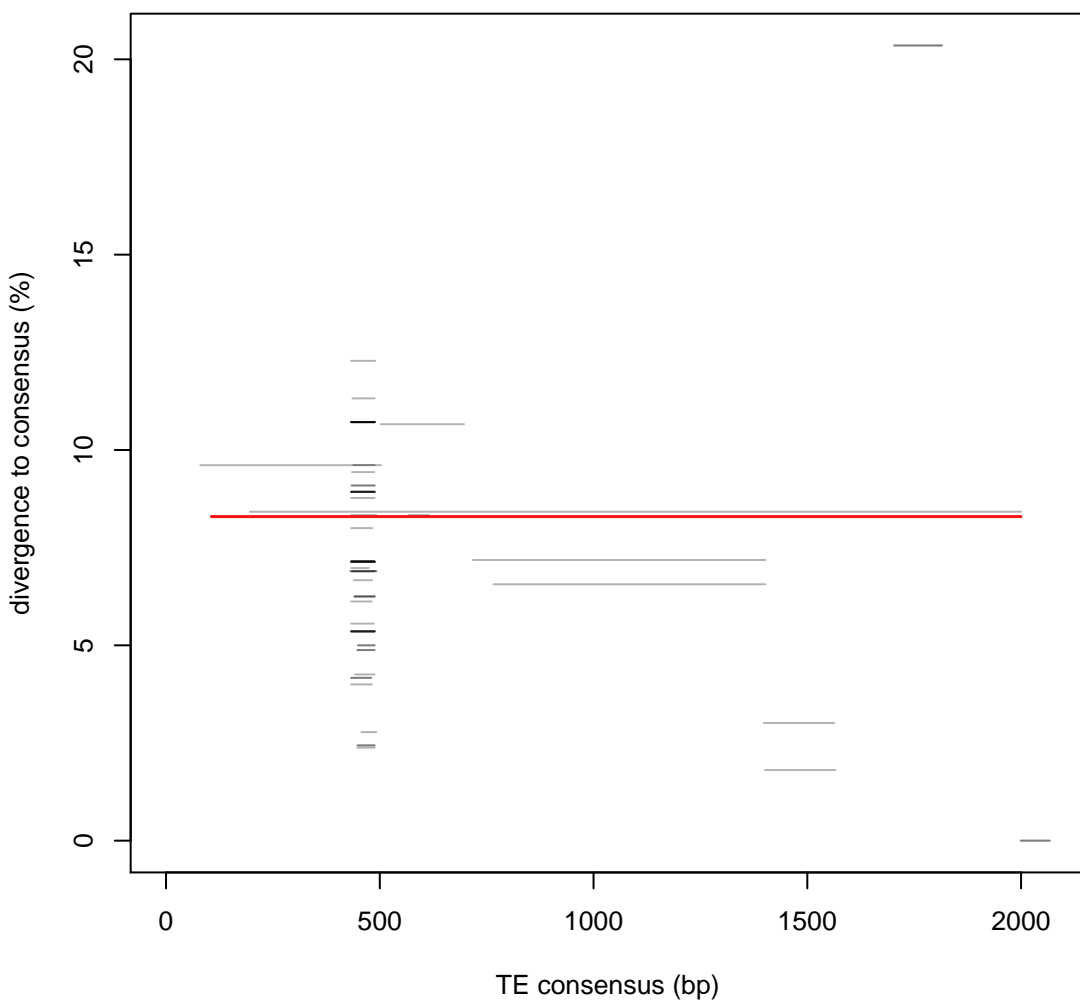
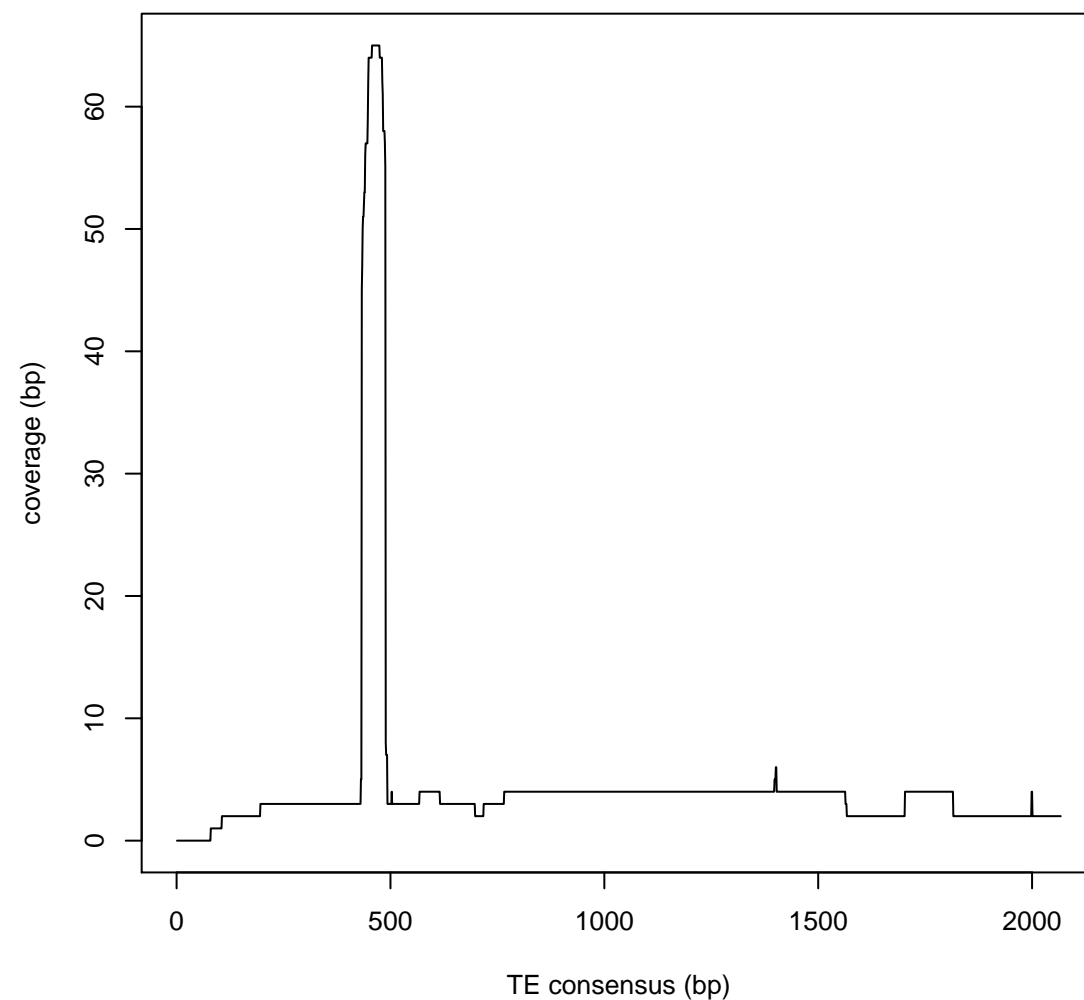


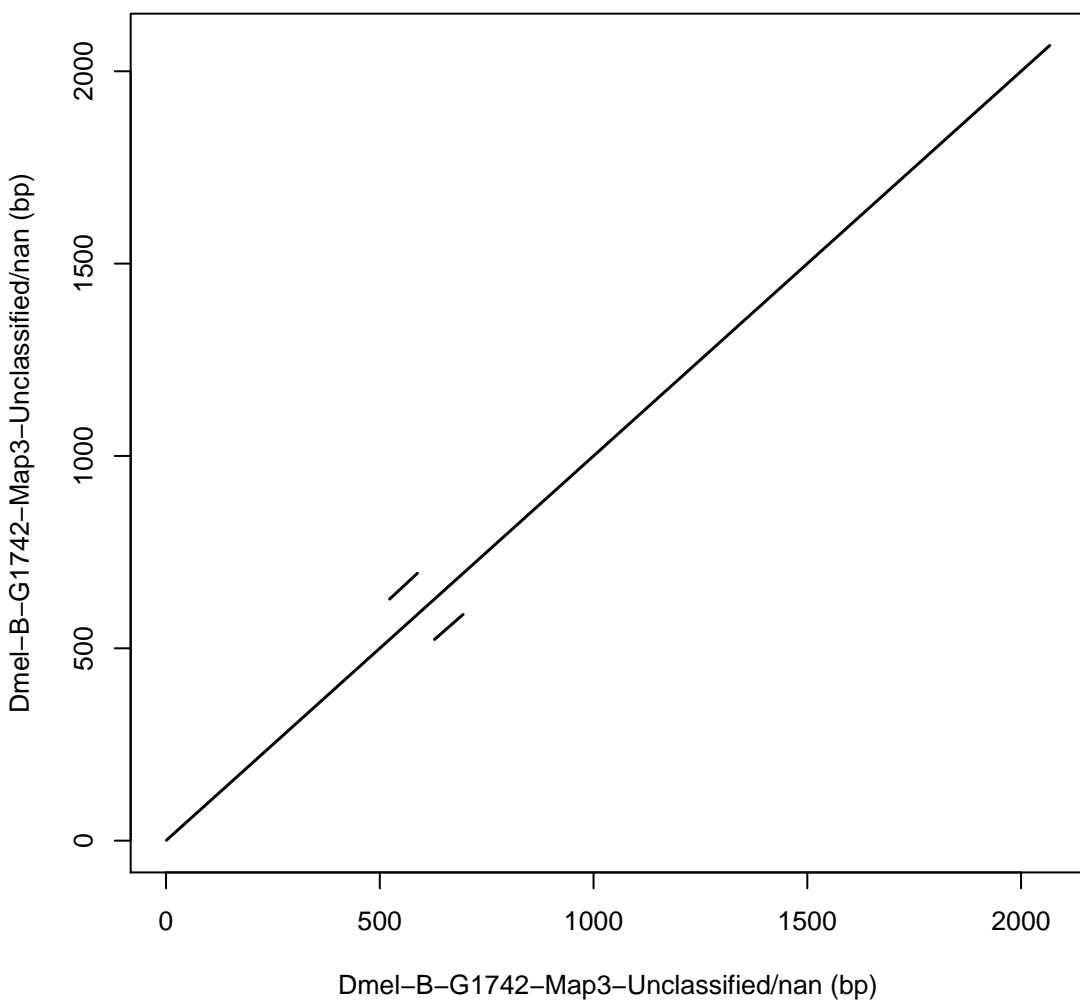
TE: Dmel-B-G1742-Map3-Unclassified/nan
consensus size: 2067bp; fragments: 75; full length: 1 (≥ 1860.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

