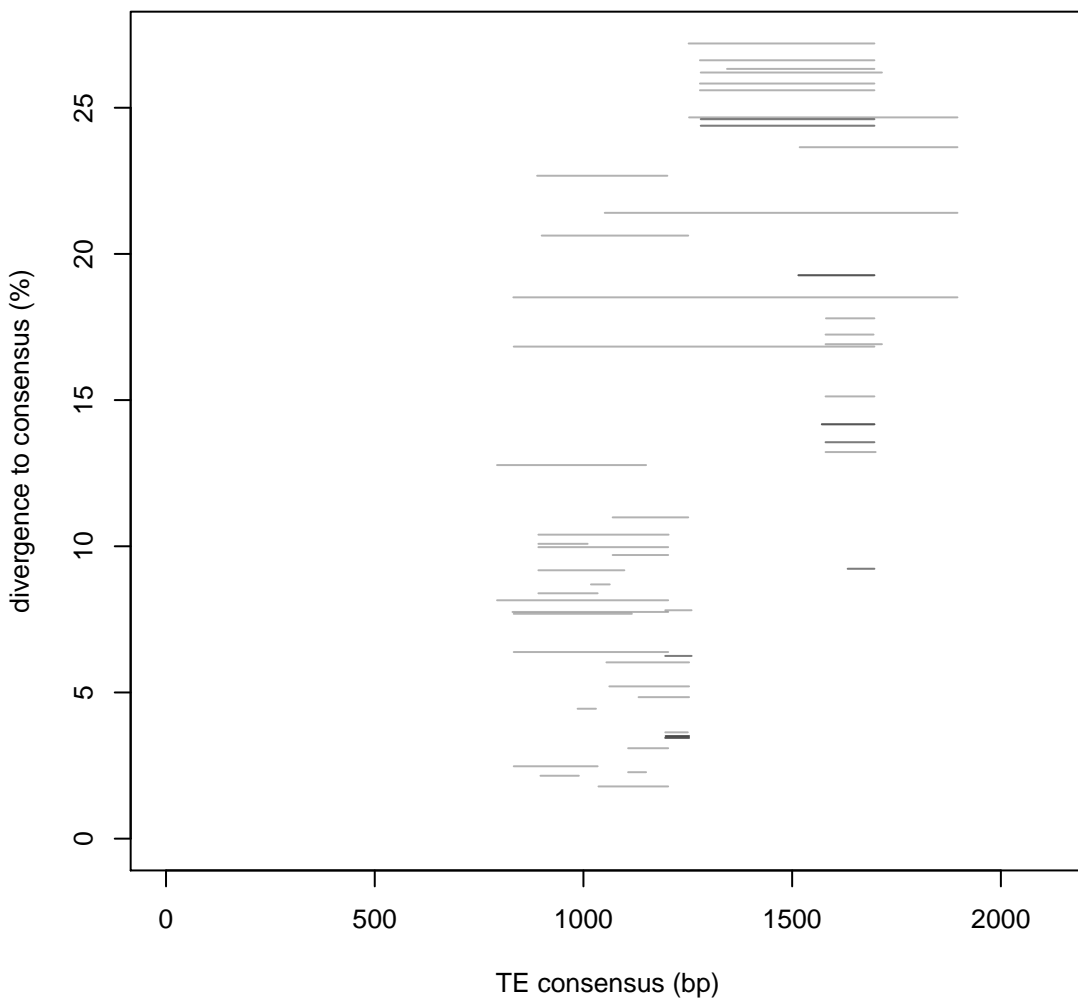
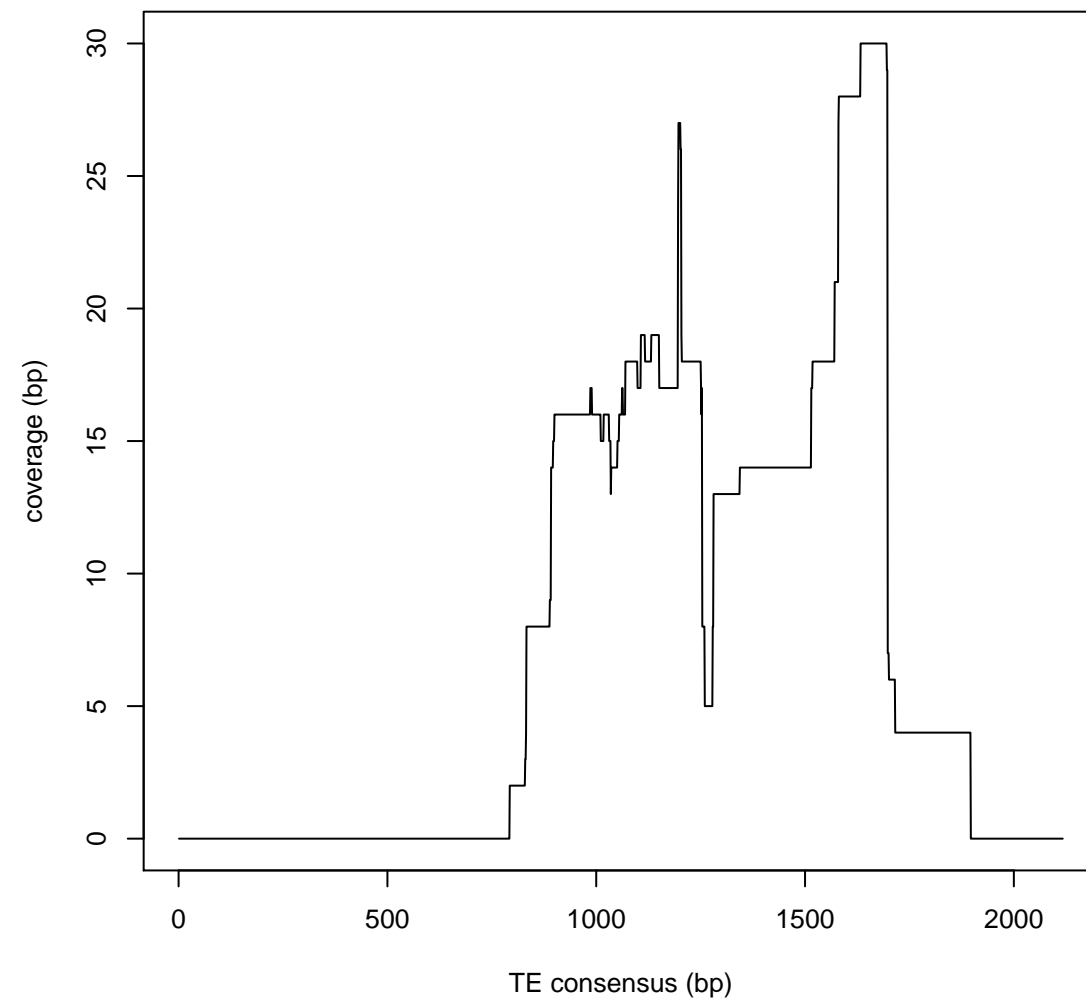


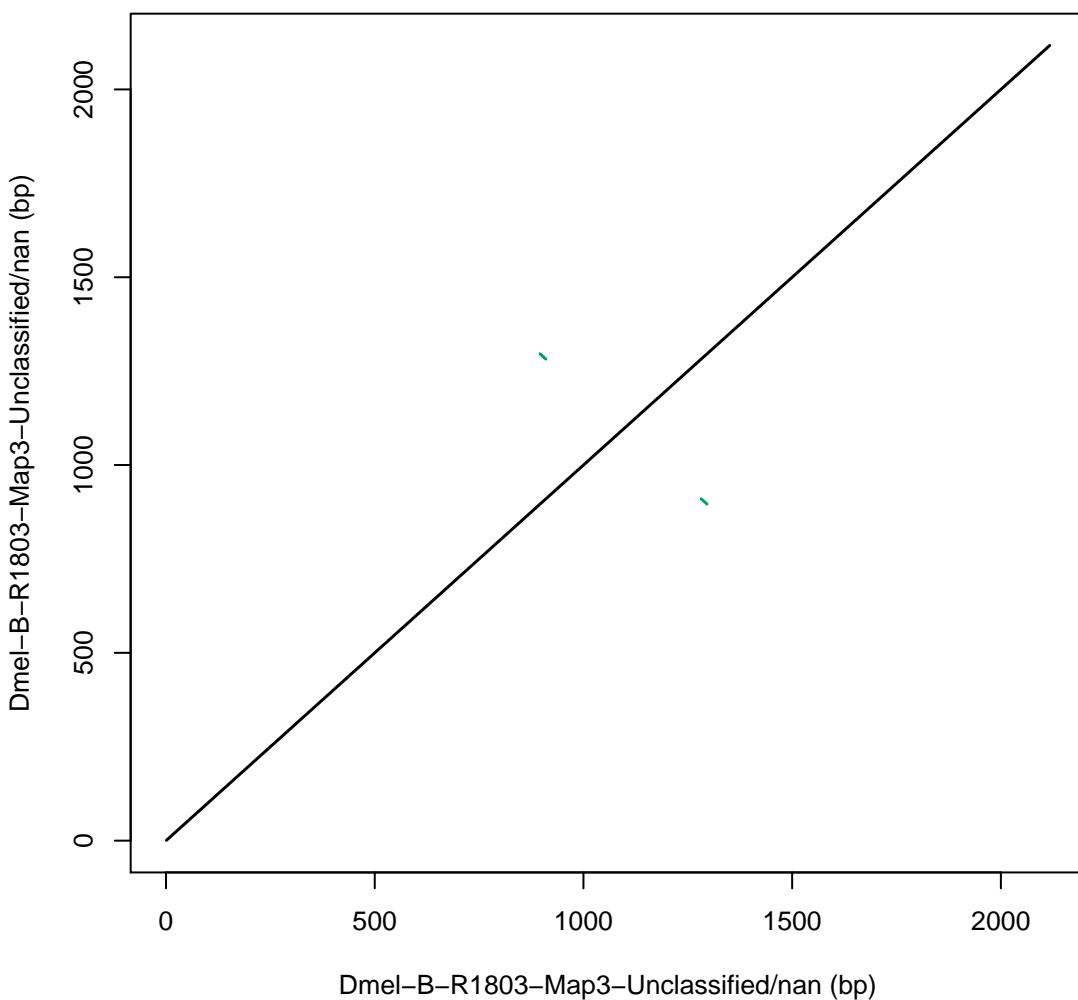
TE: Dmel-B-R1803-Map3-Unclassified/nan
consensus size: 2117bp; fragments: 64; full length: 0 (≥ 1905.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

