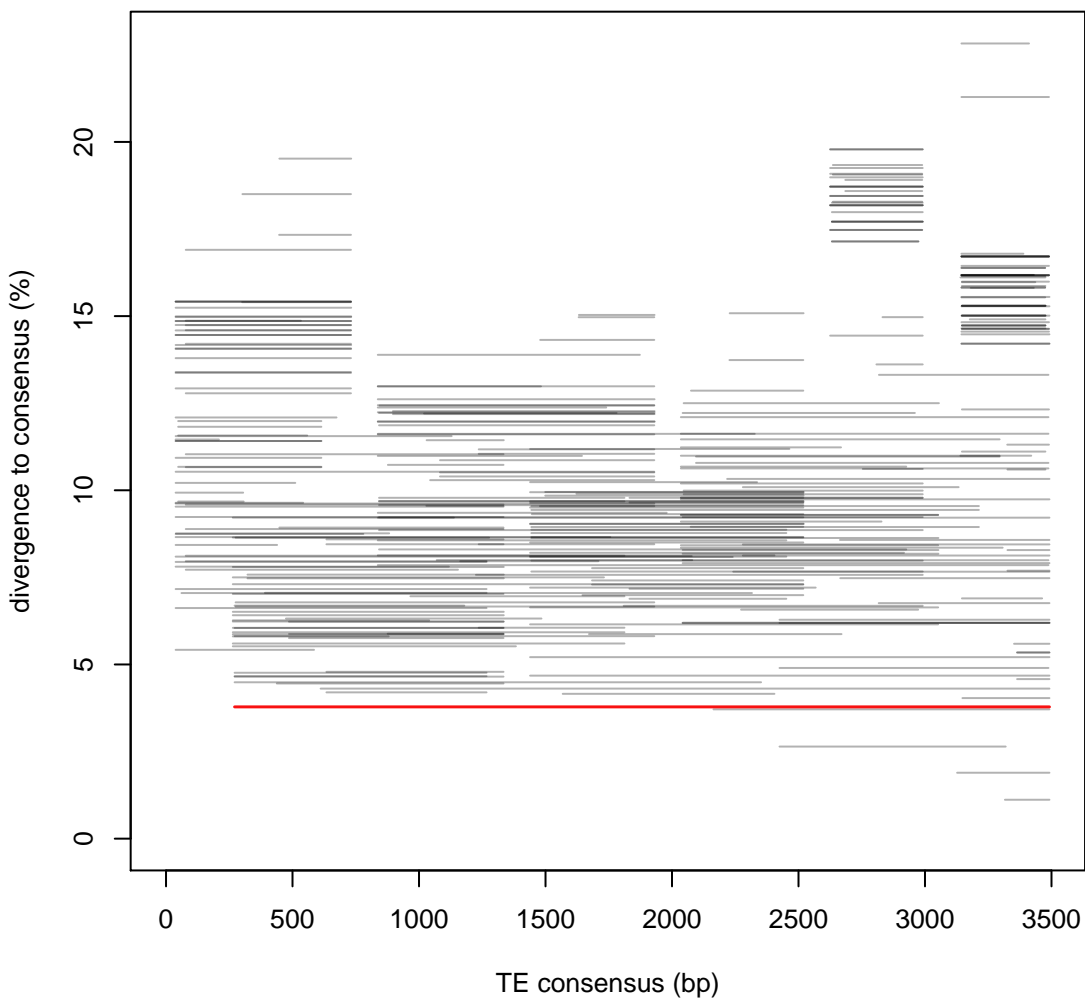
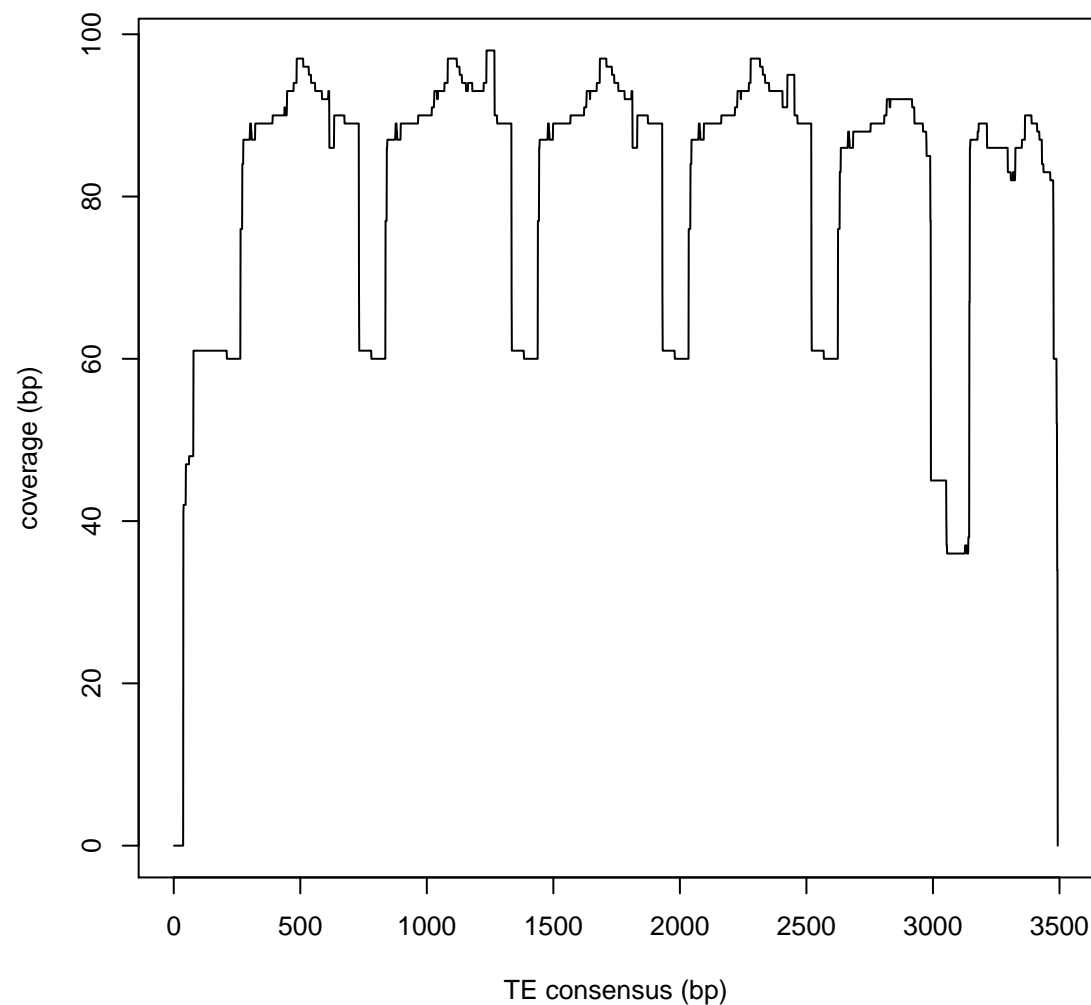


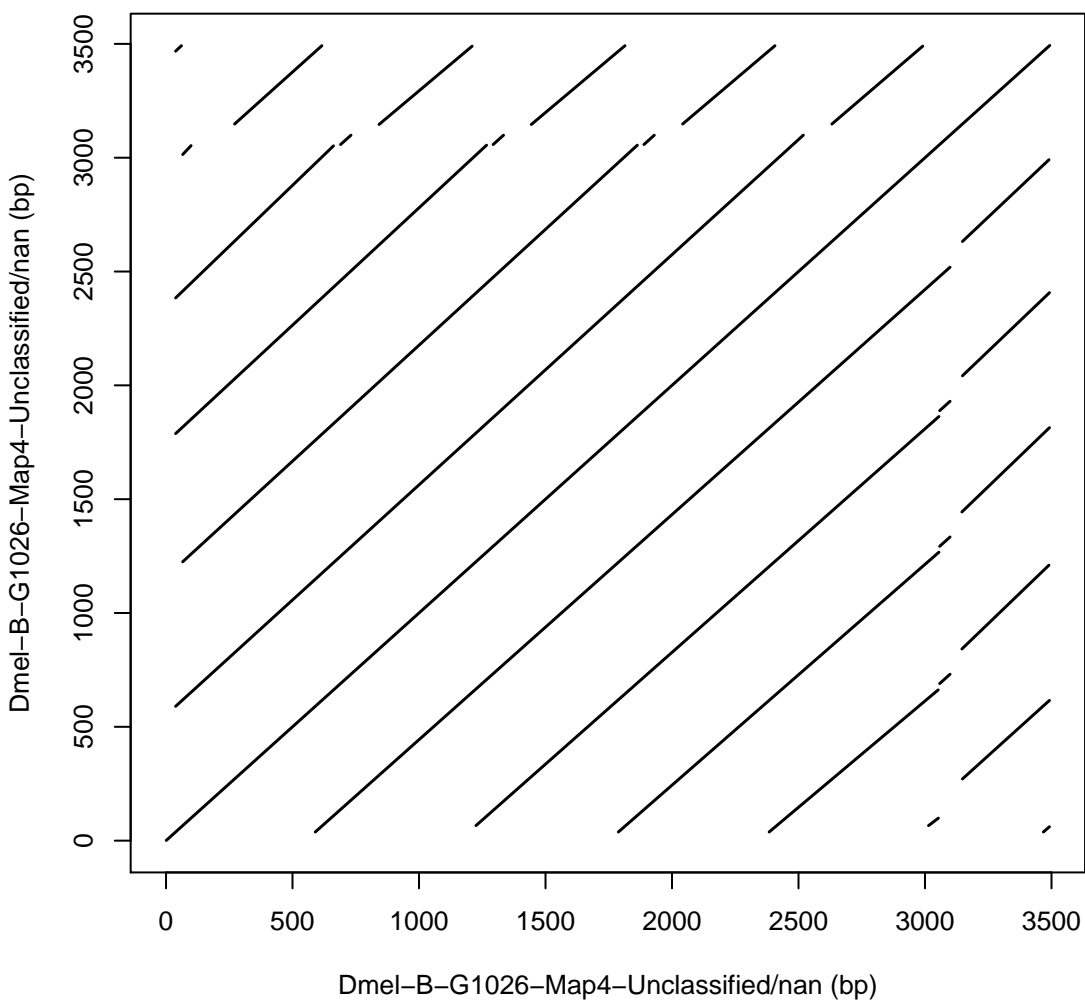
TE: Dmel-B-G1026-Map4-Unclassified/nan
consensus size: 3493bp; fragments: 339; full length: 1 (≥ 3143.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

