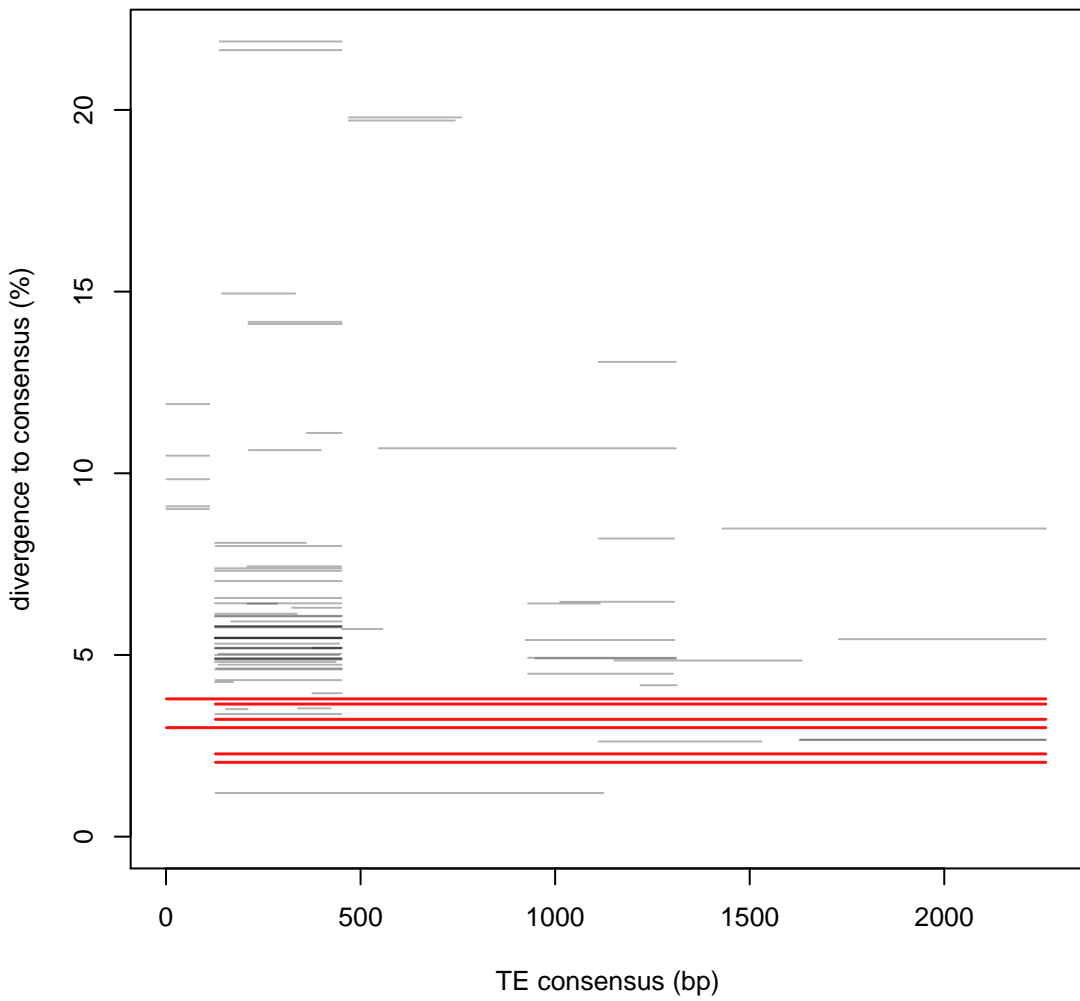
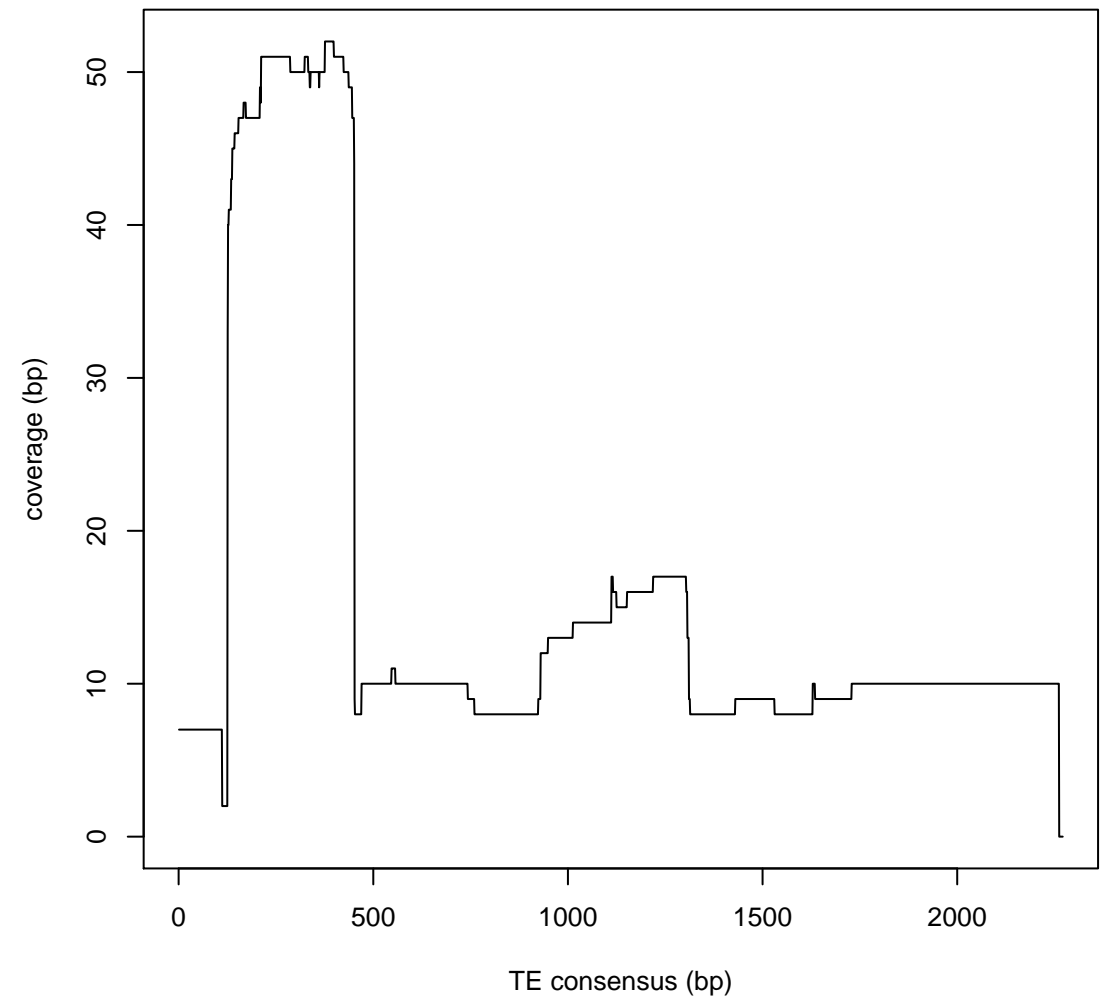


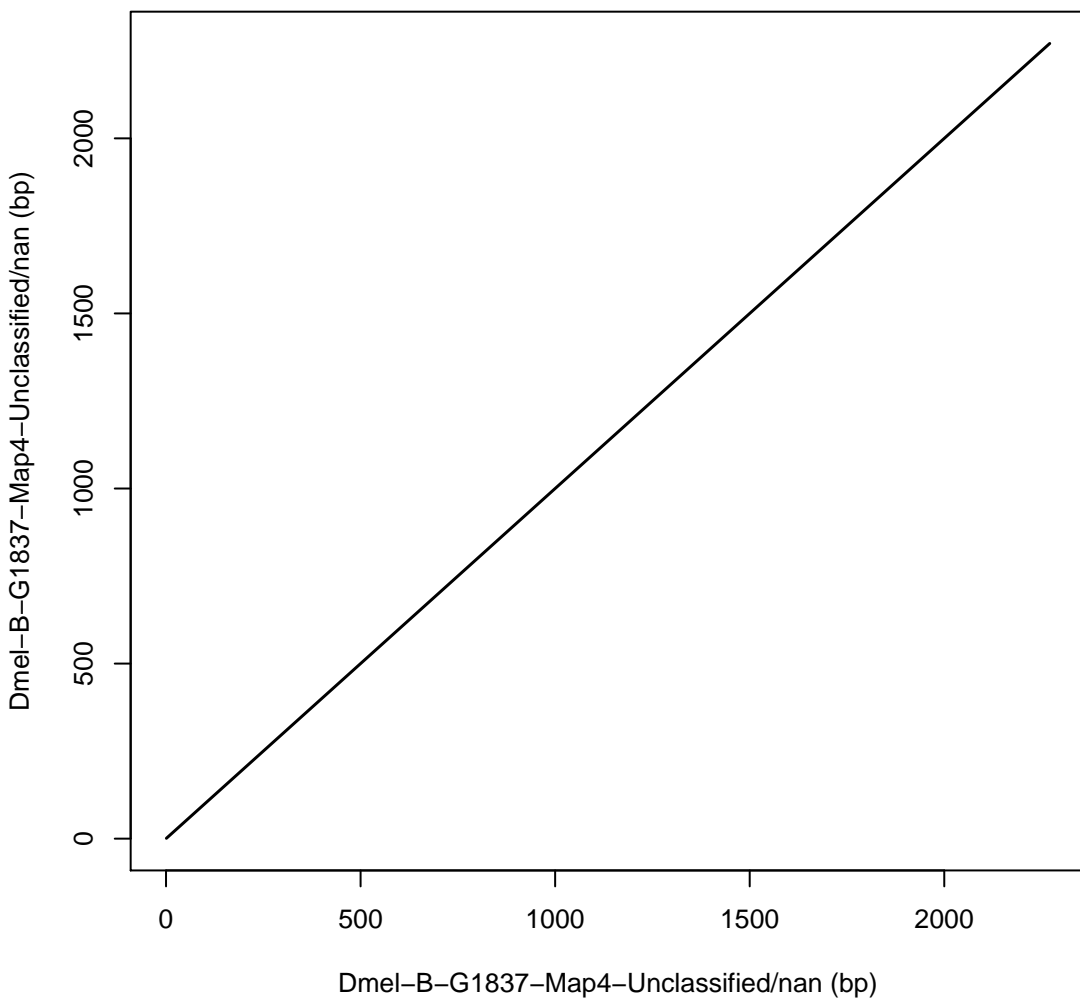
TE: Dmel-B-G1837-Map4-Unclassified/nan
consensus size: 2271bp; fragments: 82; full length: 6 (≥ 2043.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

