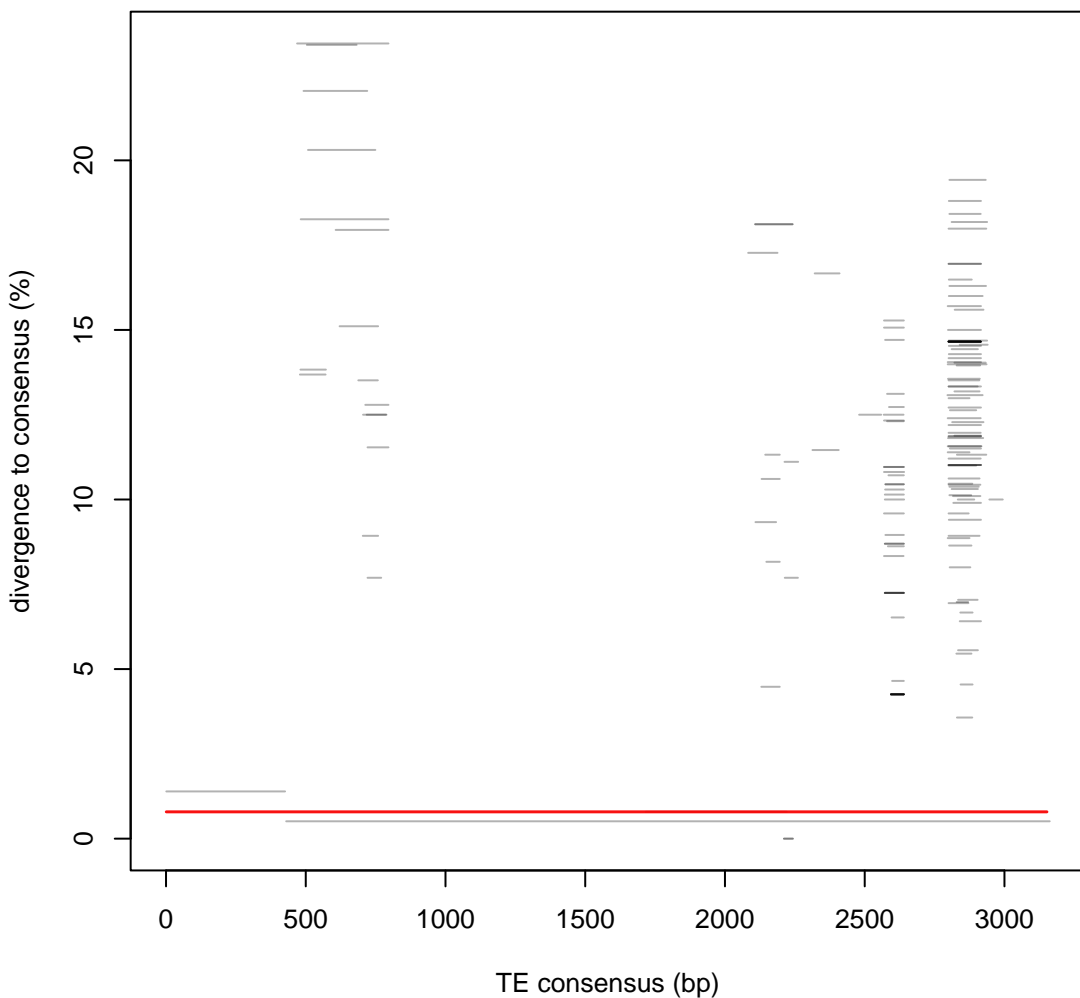
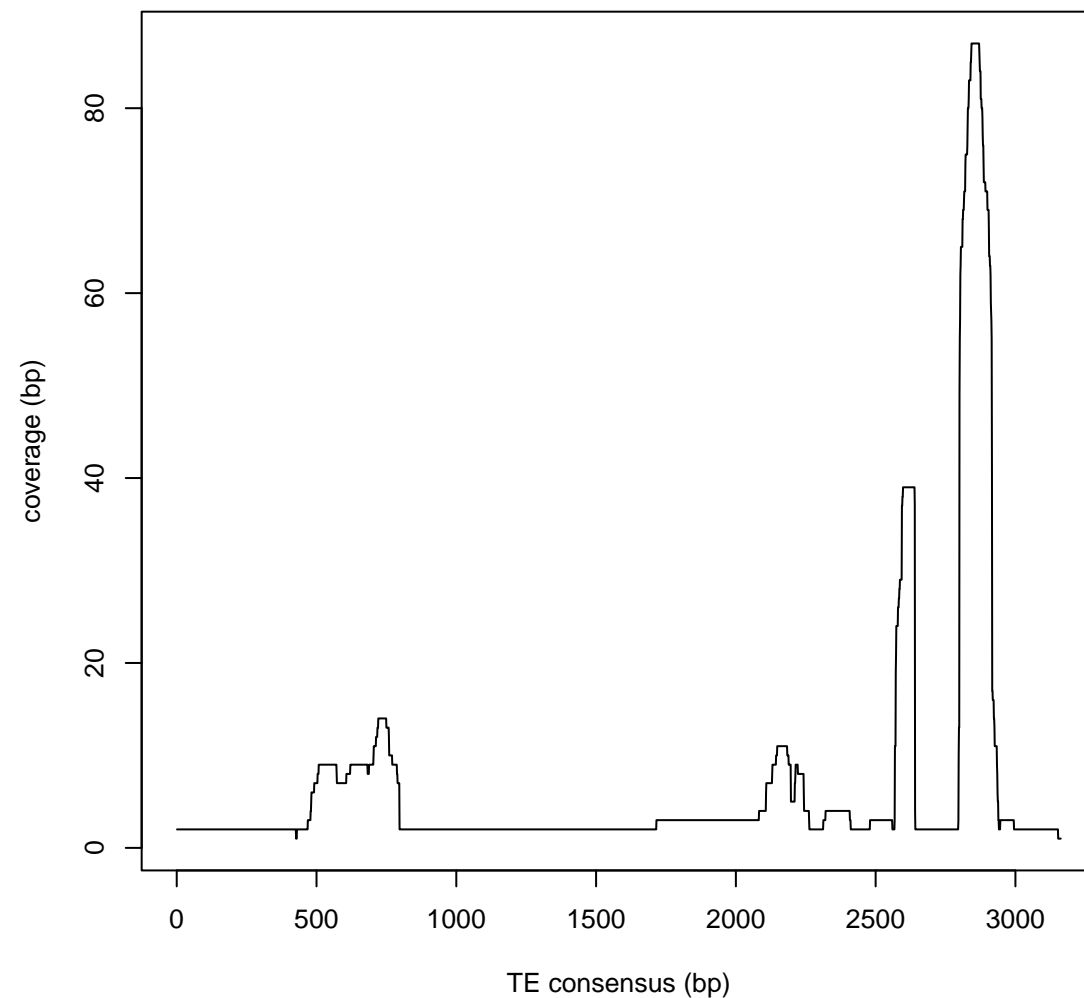


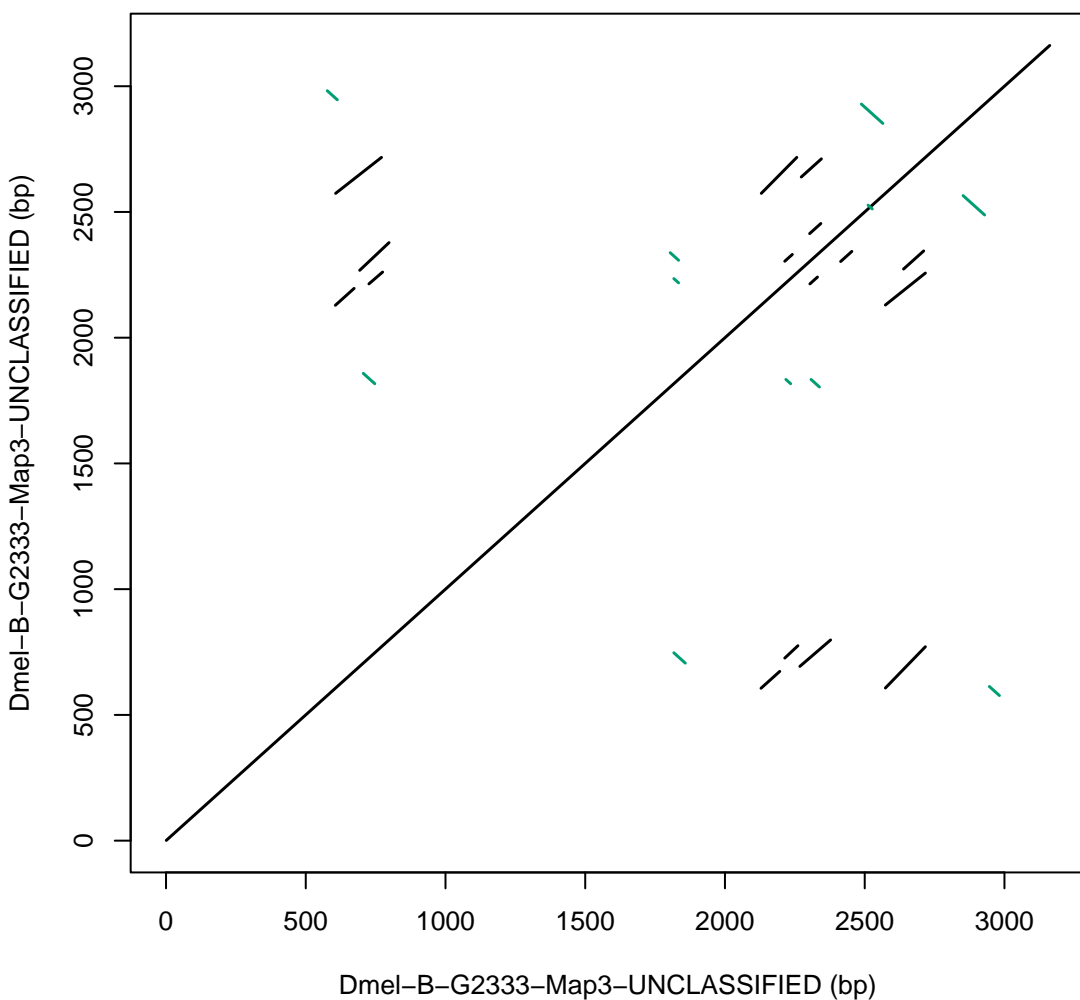
TE: Dmel-B-G2333-Map3-UNCLASSIFIED
consensus size: 3162bp; fragments: 158; full length: 1 (≥ 2845.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

