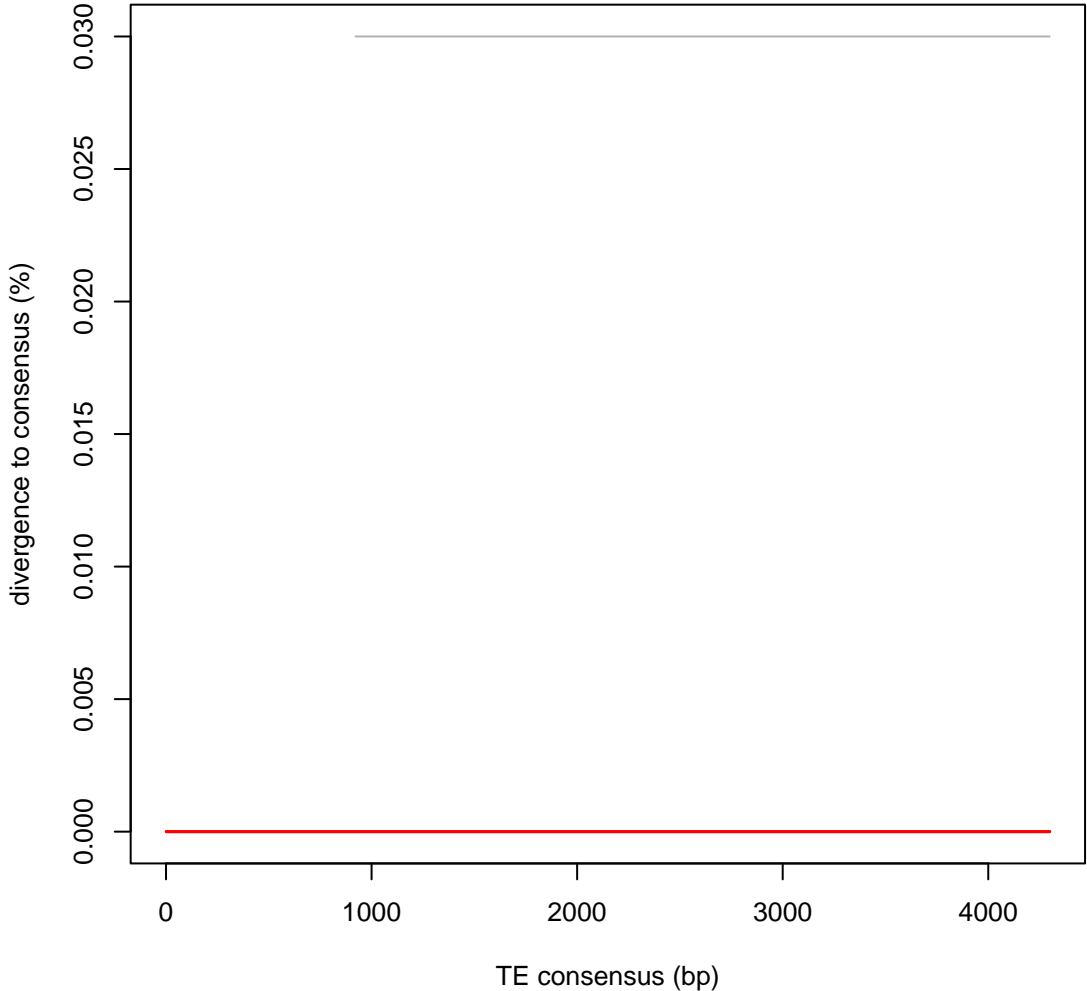
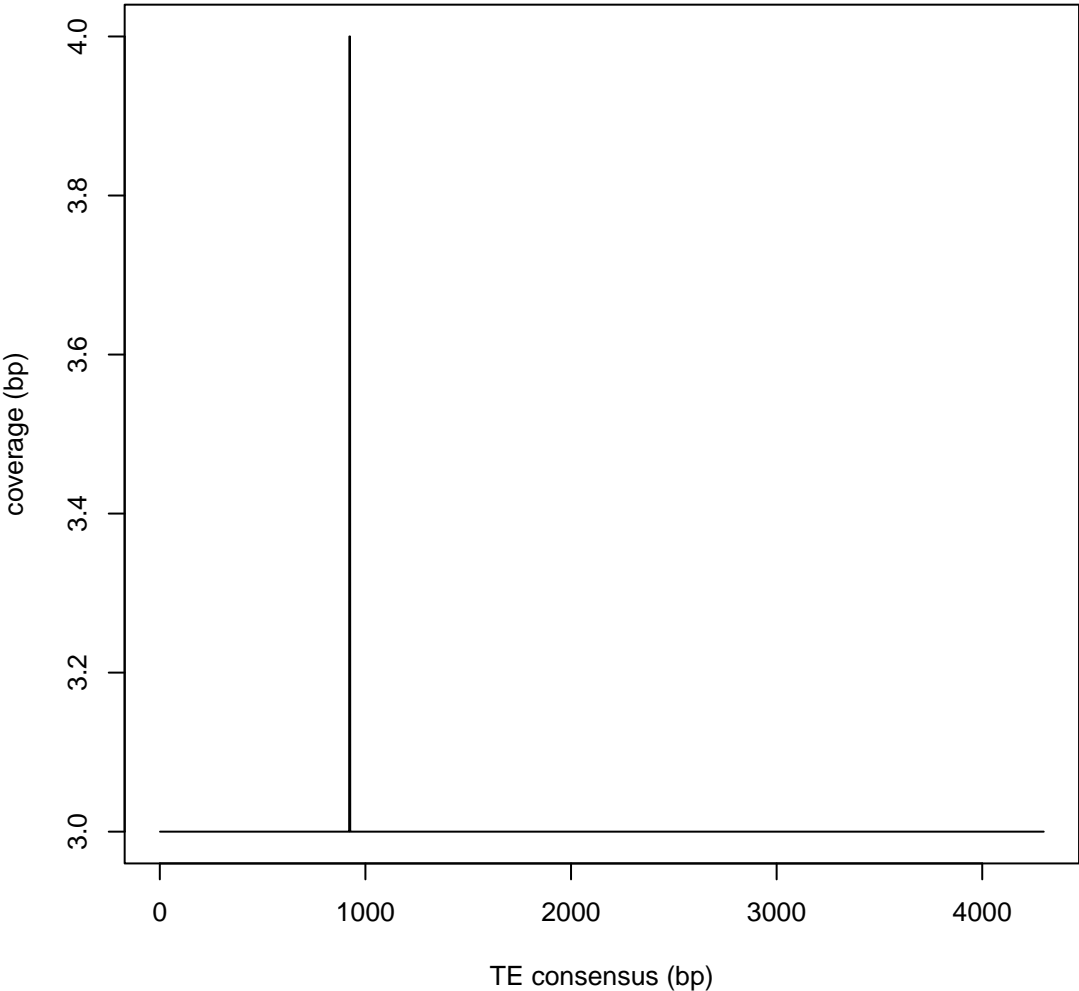


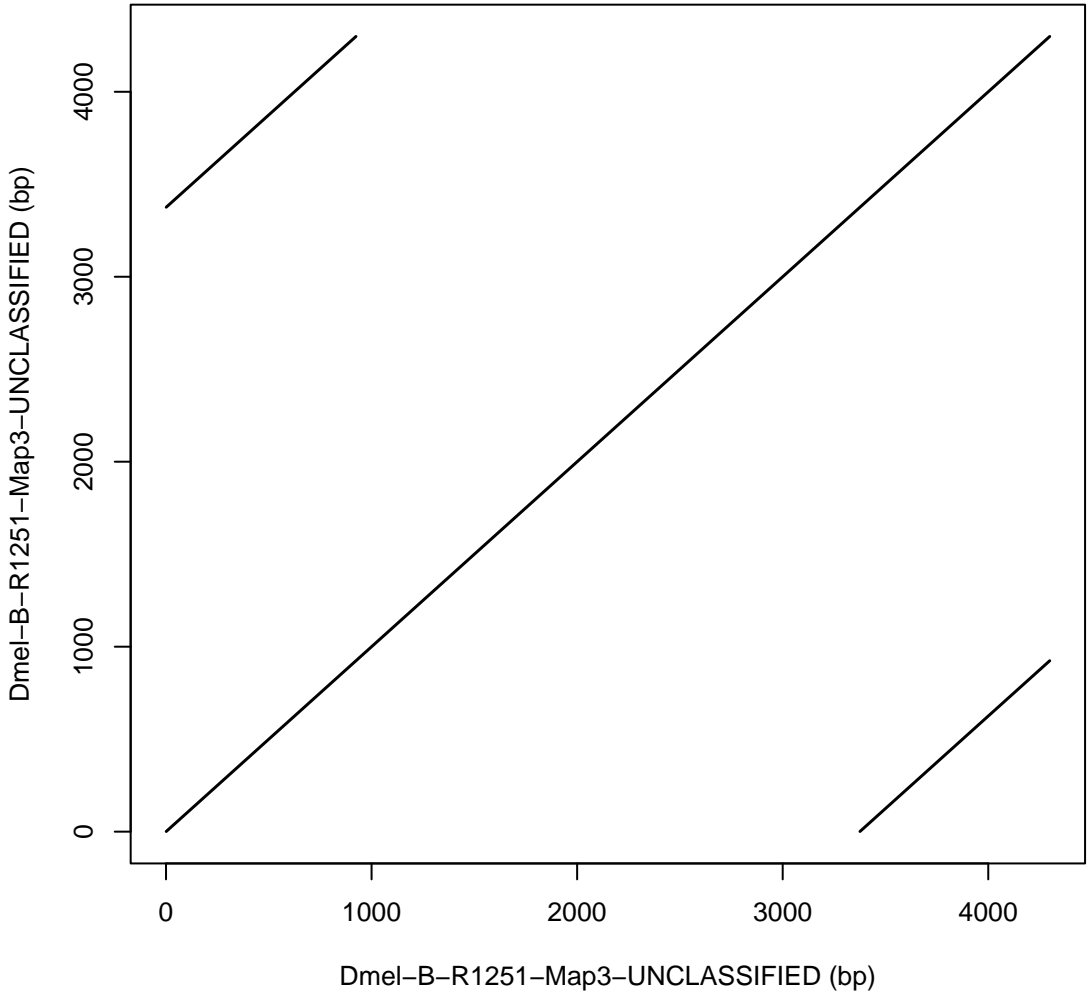
TE: Dmel-B-R1251-Map3-UNCLASSIFIED
consensus size: 4299bp; fragments: 4; full length: 2 (>=3869.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

