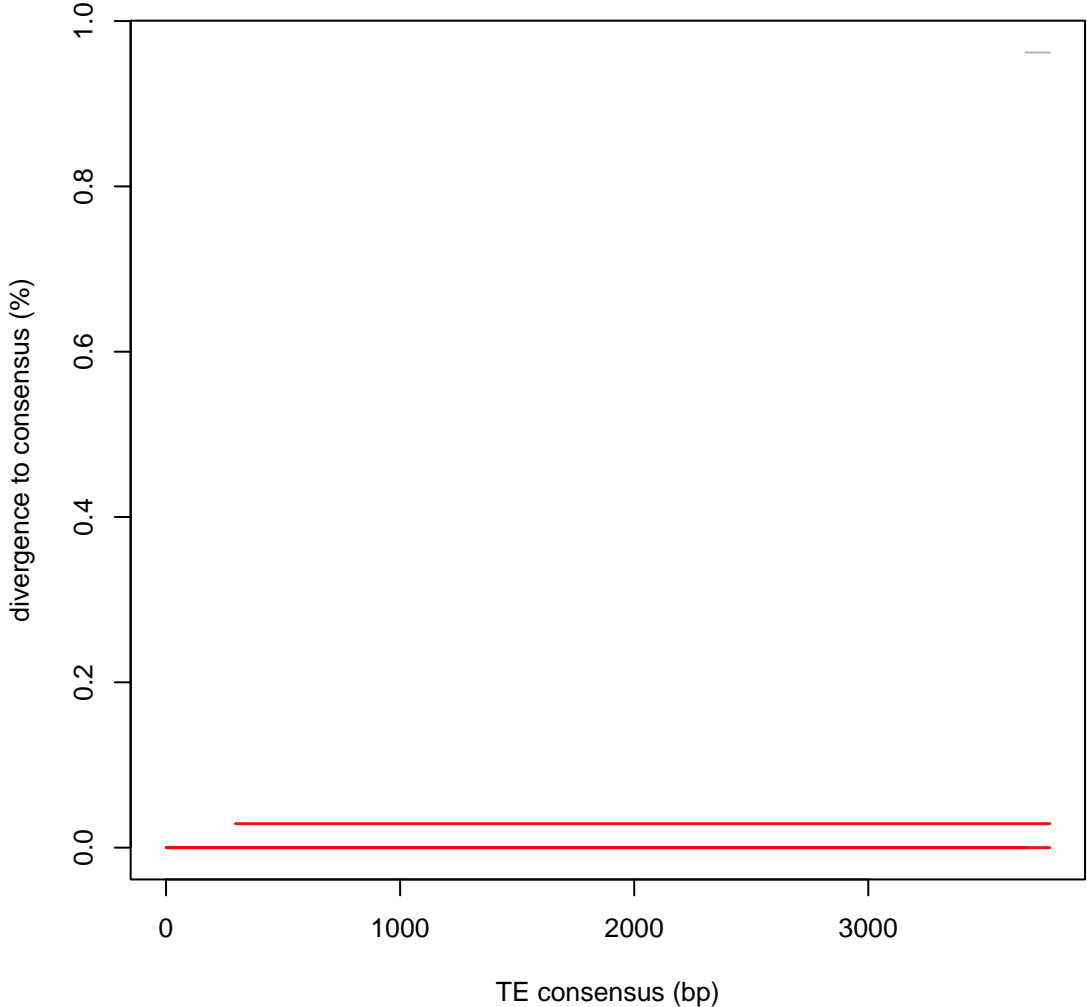
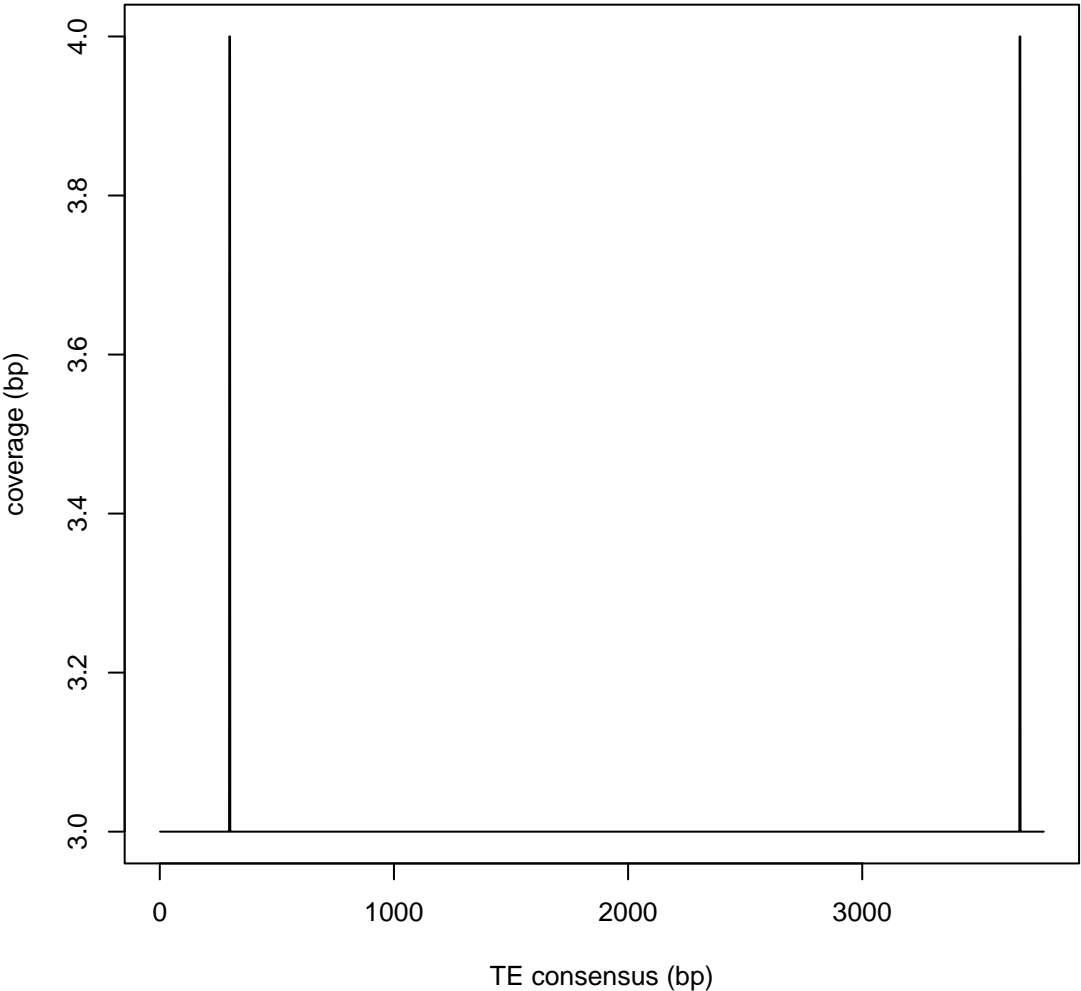


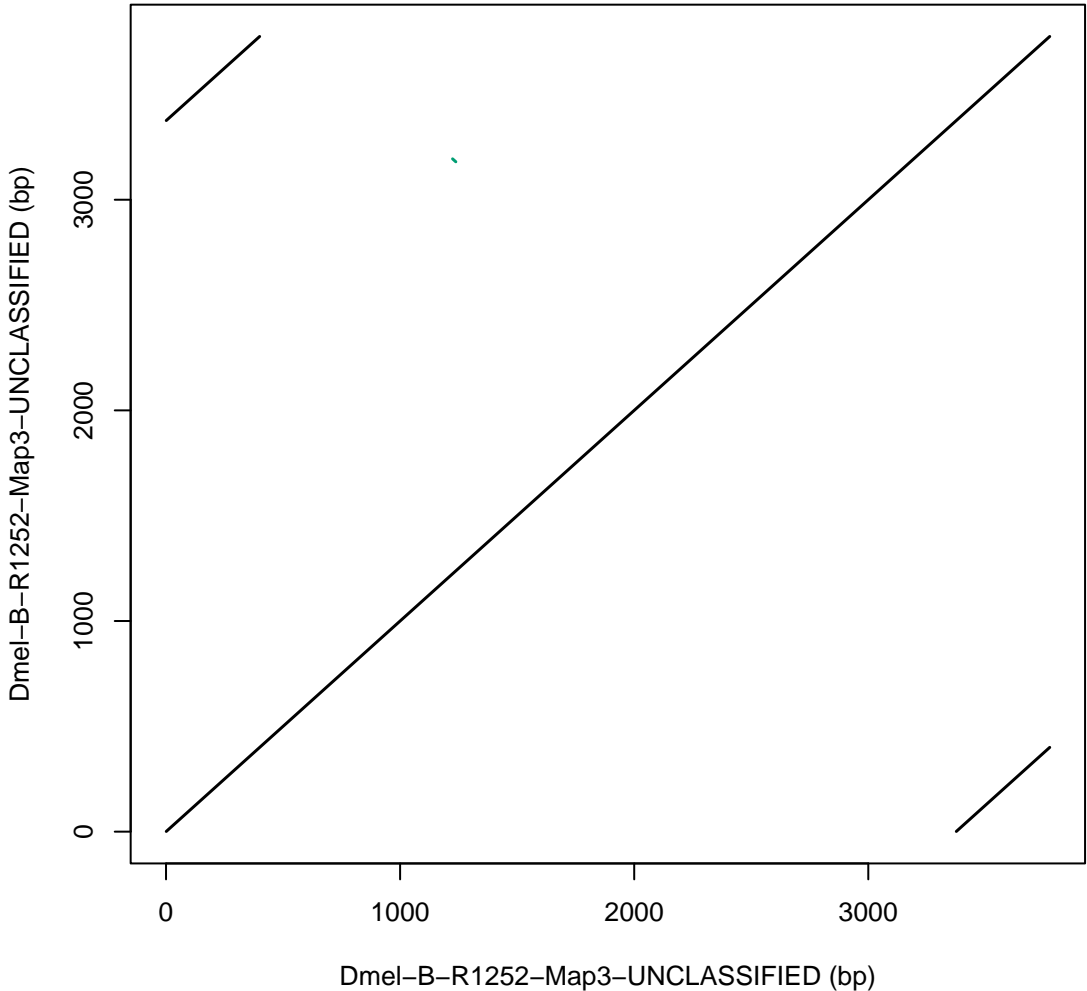
TE: Dmel-B-R1252-Map3-UNCLASSIFIED
consensus size: 3775bp; fragments: 5; full length: 3 (>=3397.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

