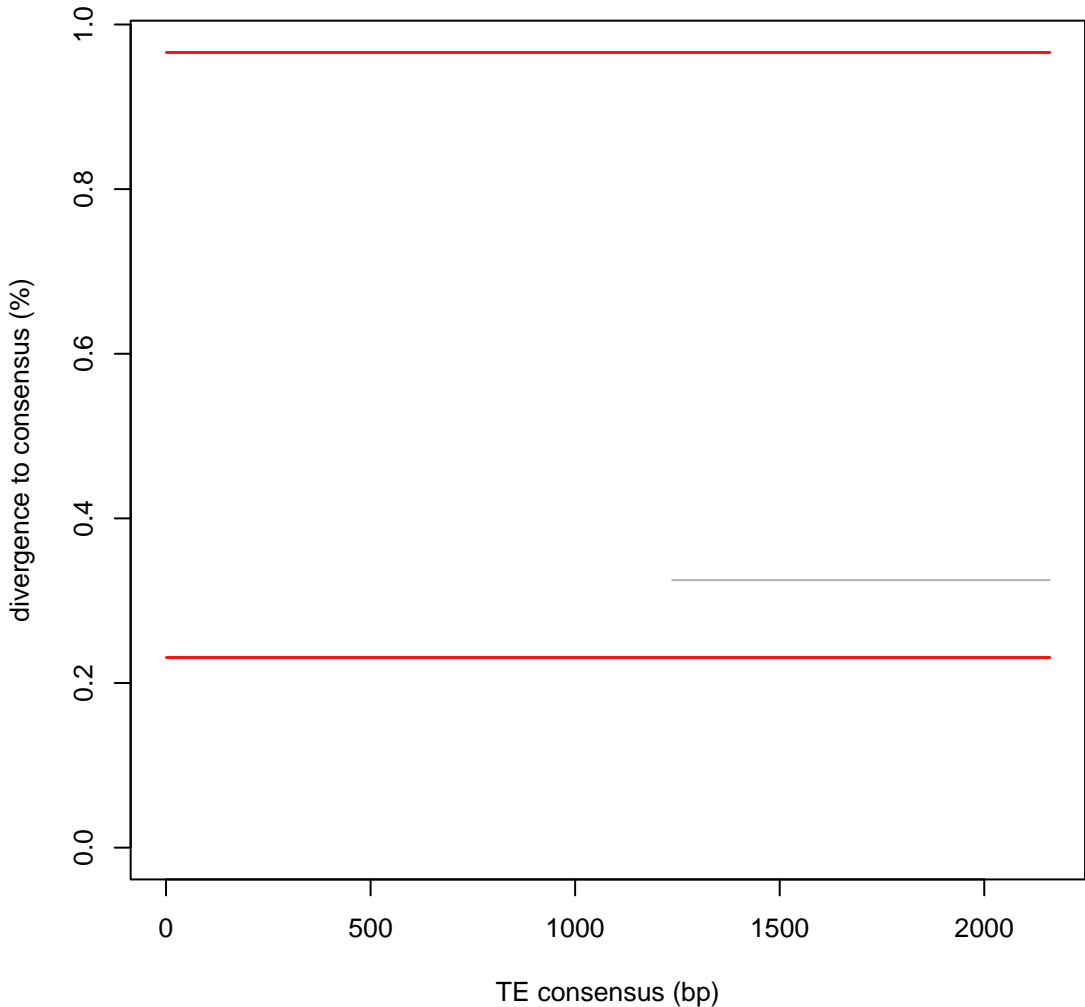
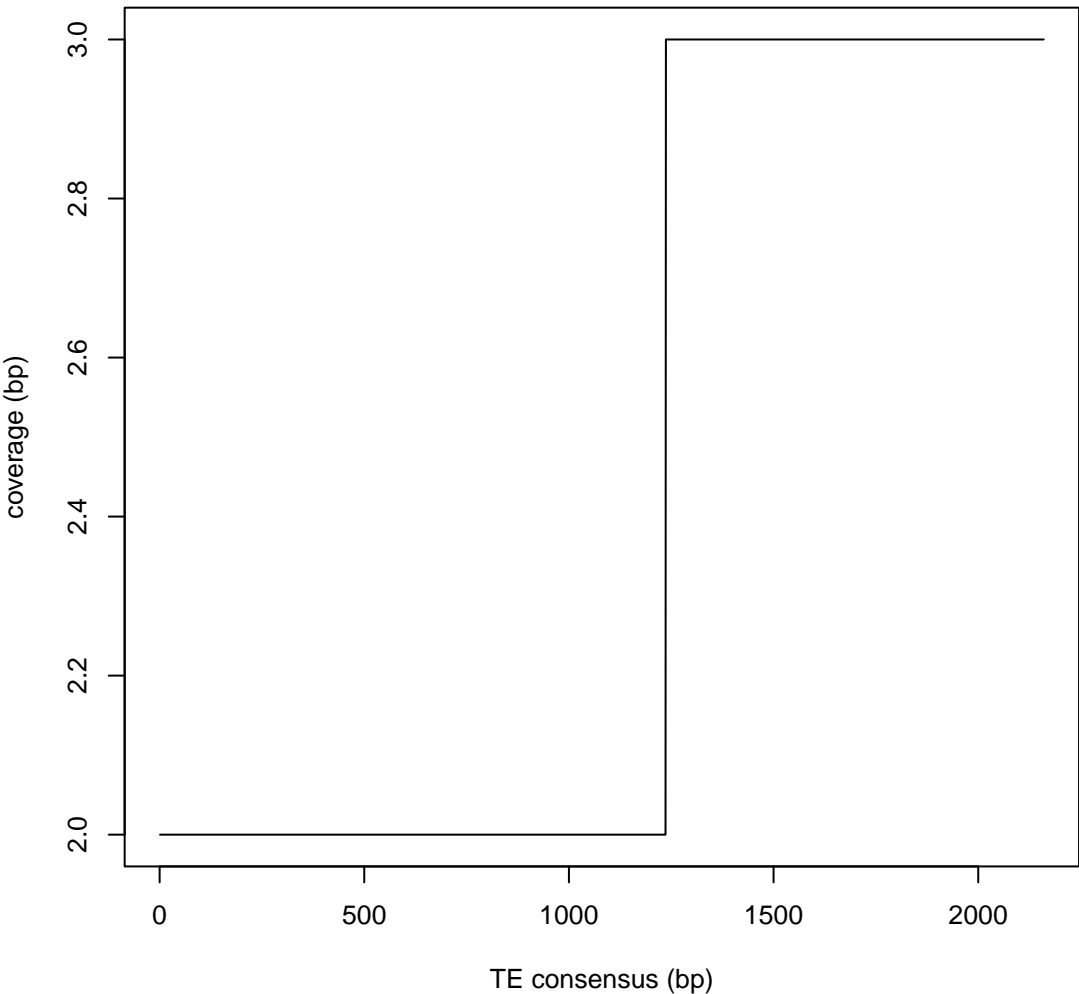


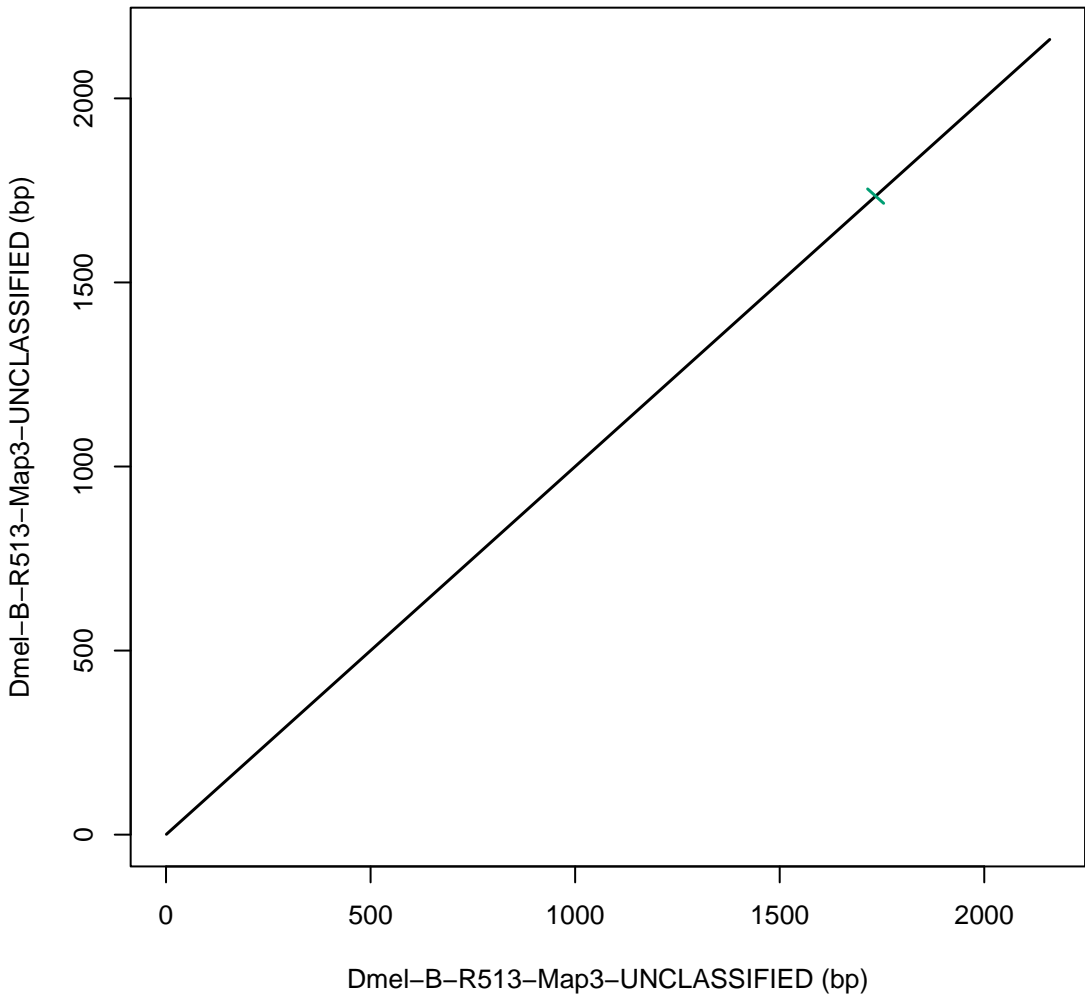
TE: Dmel-B-R513-Map3-UNCLASSIFIED
consensus size: 2160bp; fragments: 3; full length: 2 (>=1944bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

