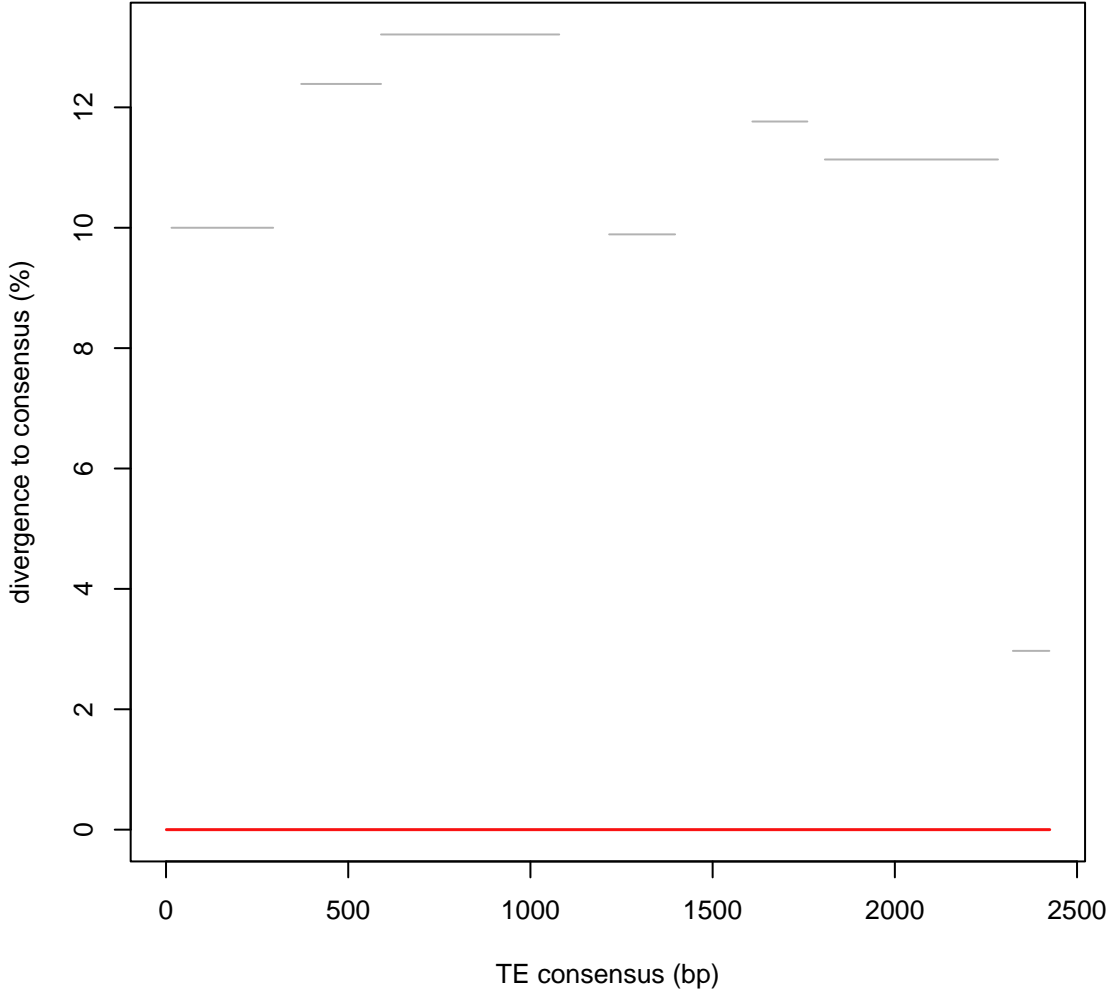
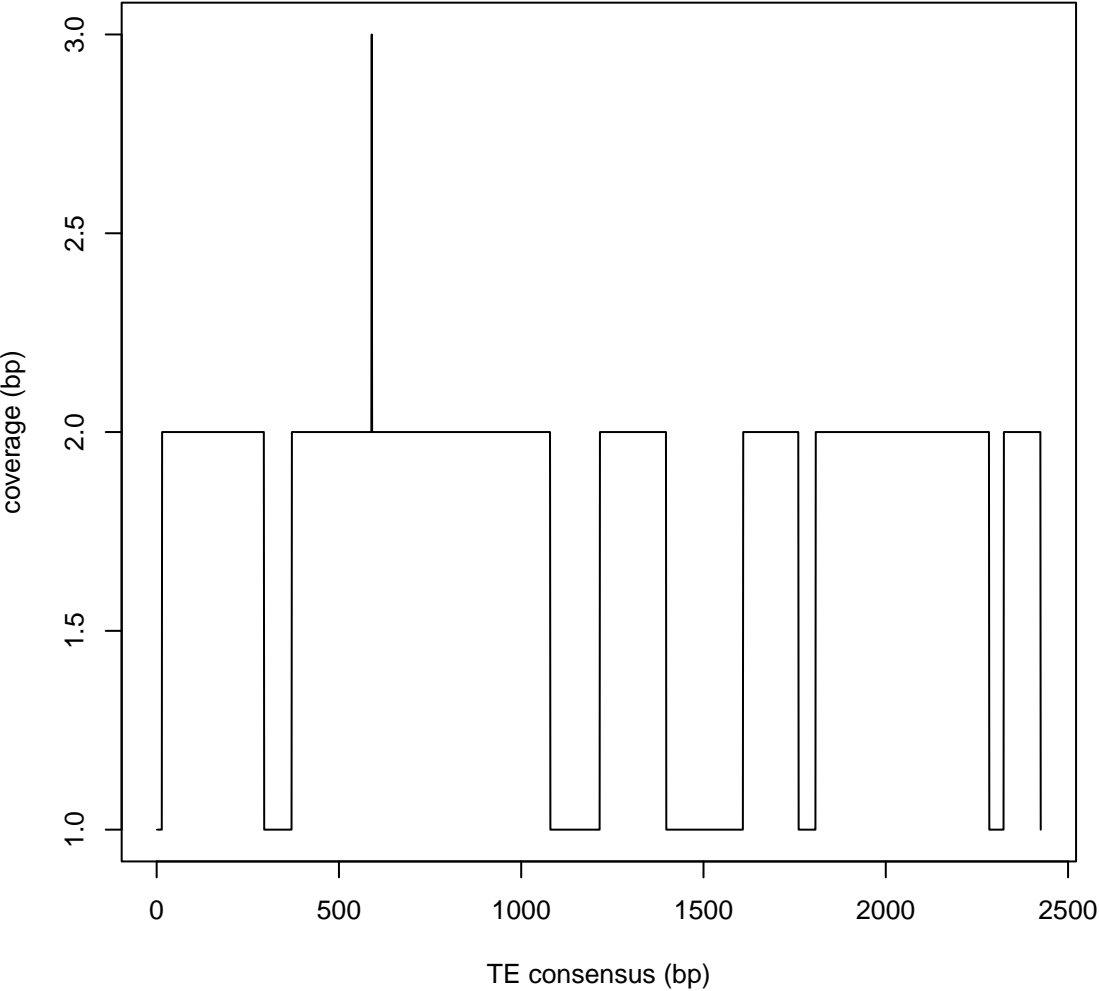


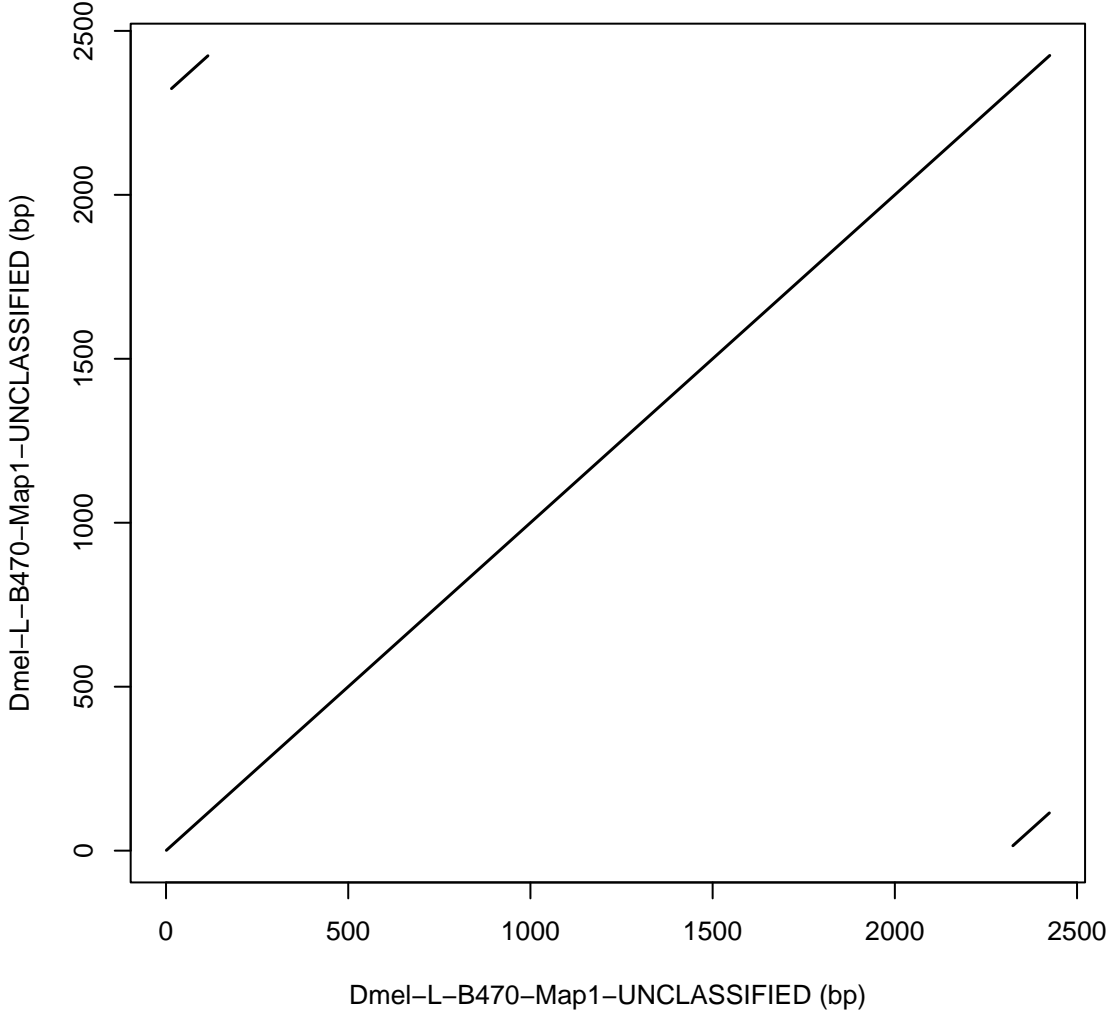
TE: Dmel-L-B470-Map1-UNCLASSIFIED
consensus size: 2425bp; fragments: 8; full length: 1 (>=2182.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

