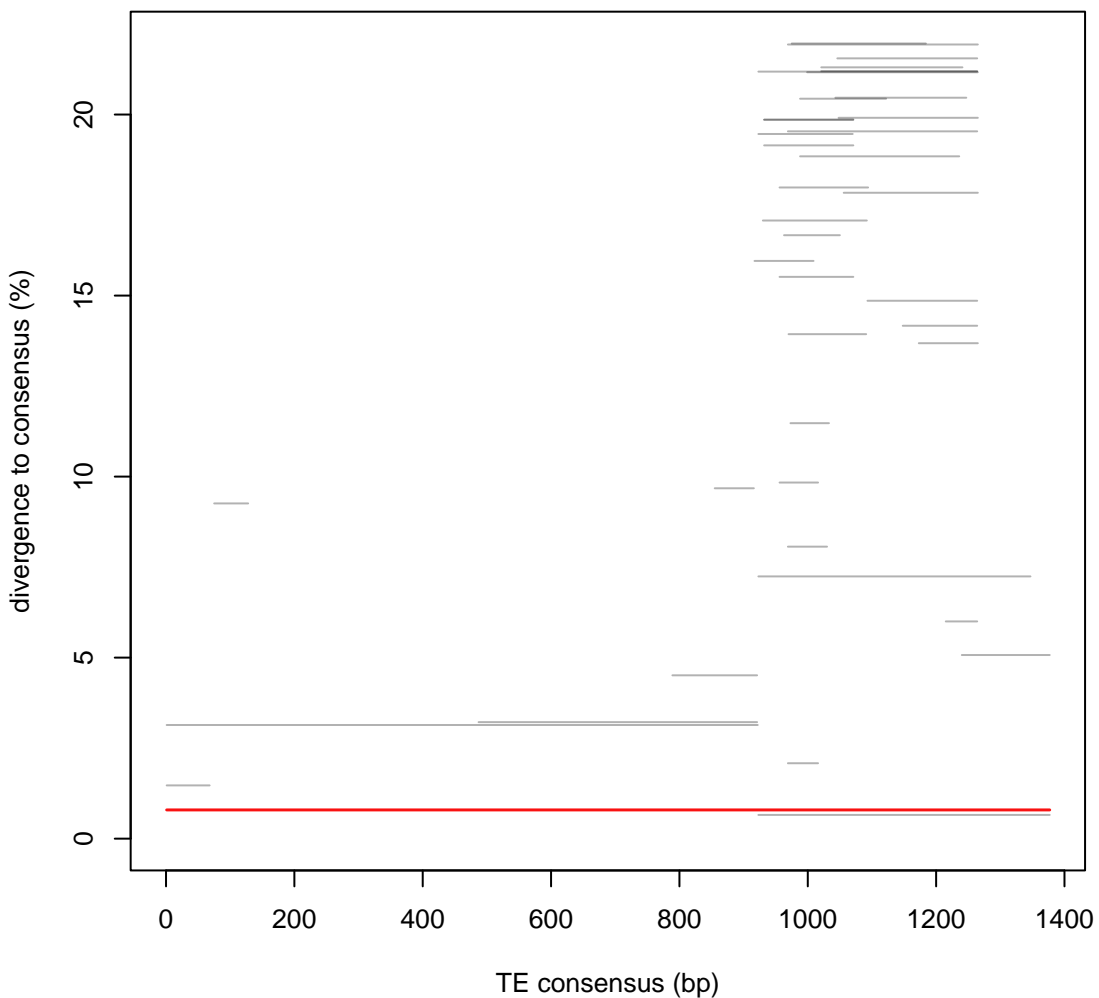
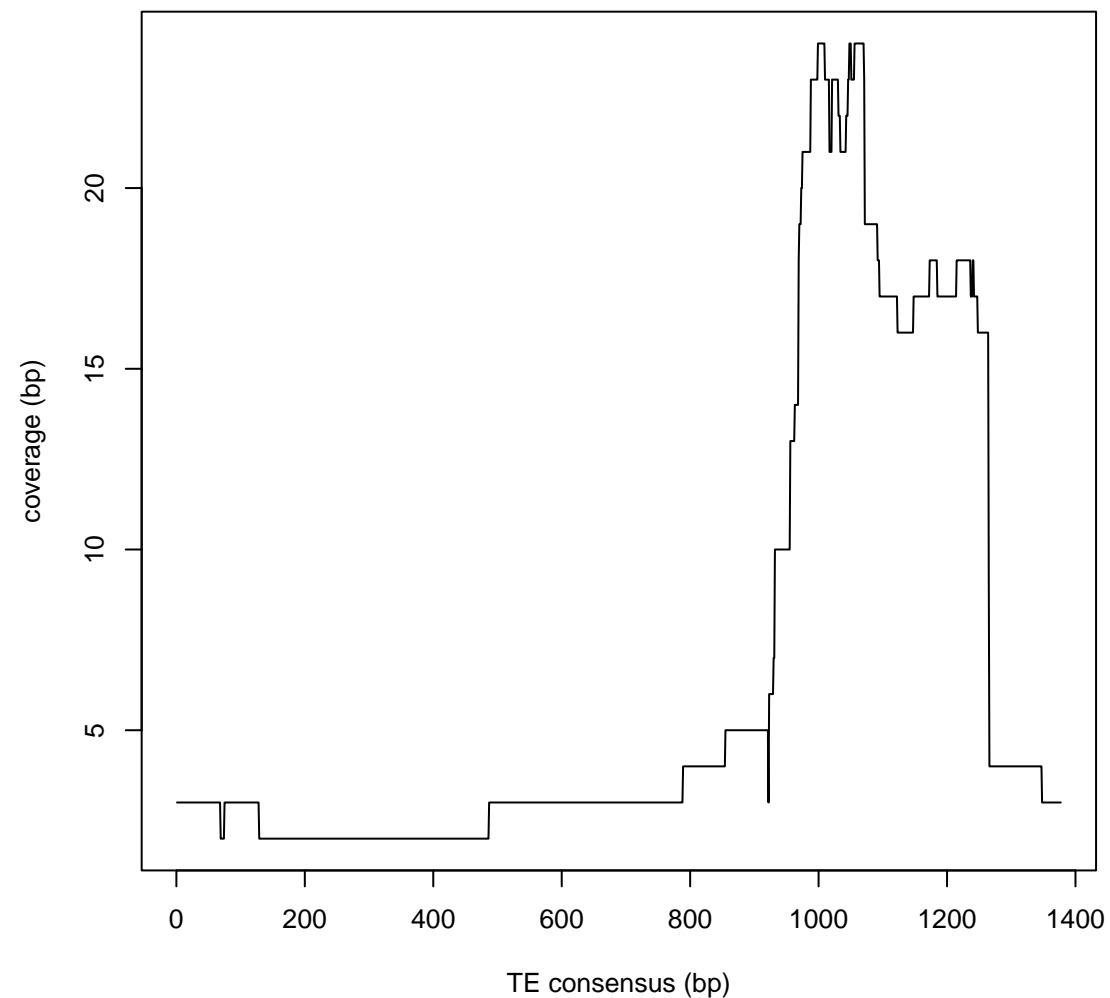


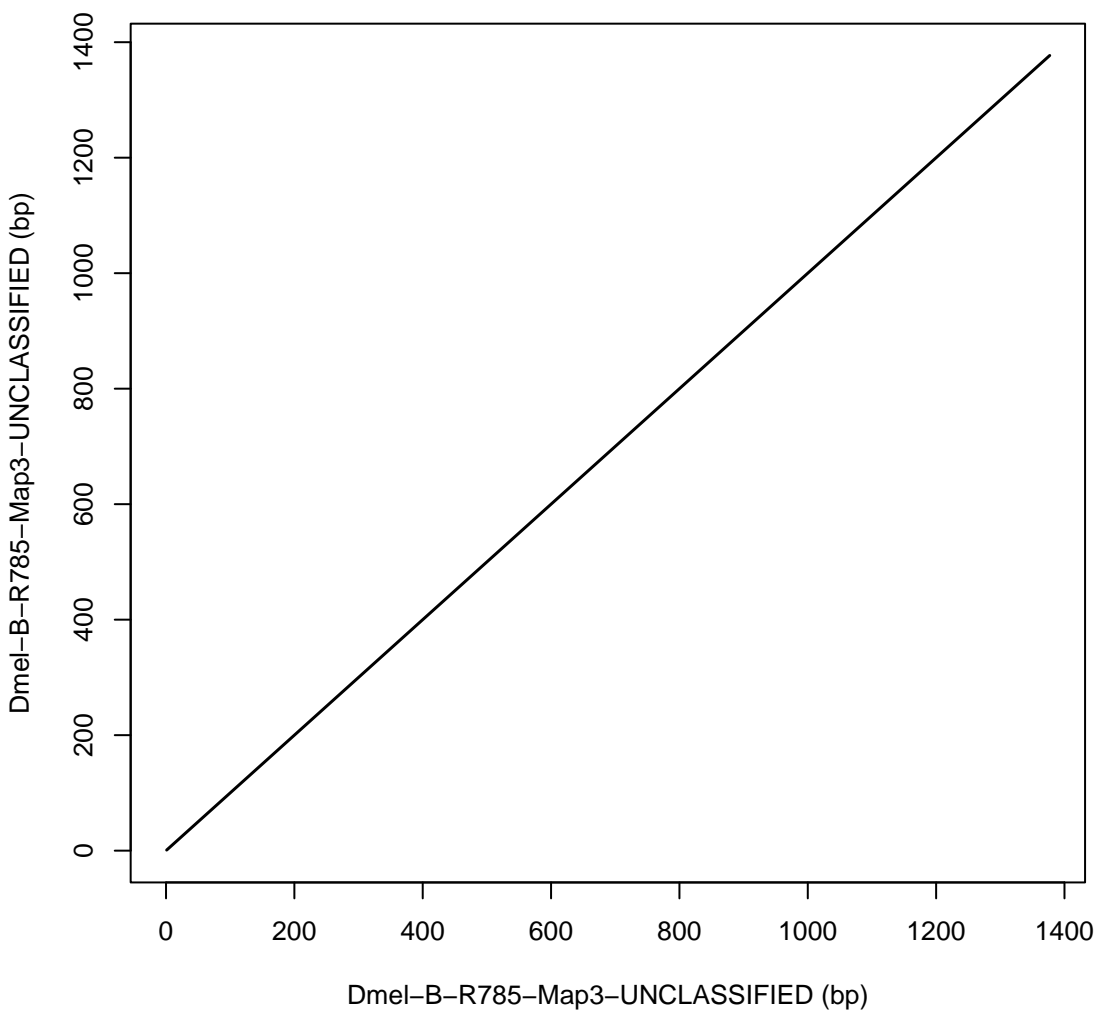
TE: Dmel-B-R785-Map3-UNCLASSIFIED
consensus size: 1377bp; fragments: 41; full length: 1 (≥ 1239.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

