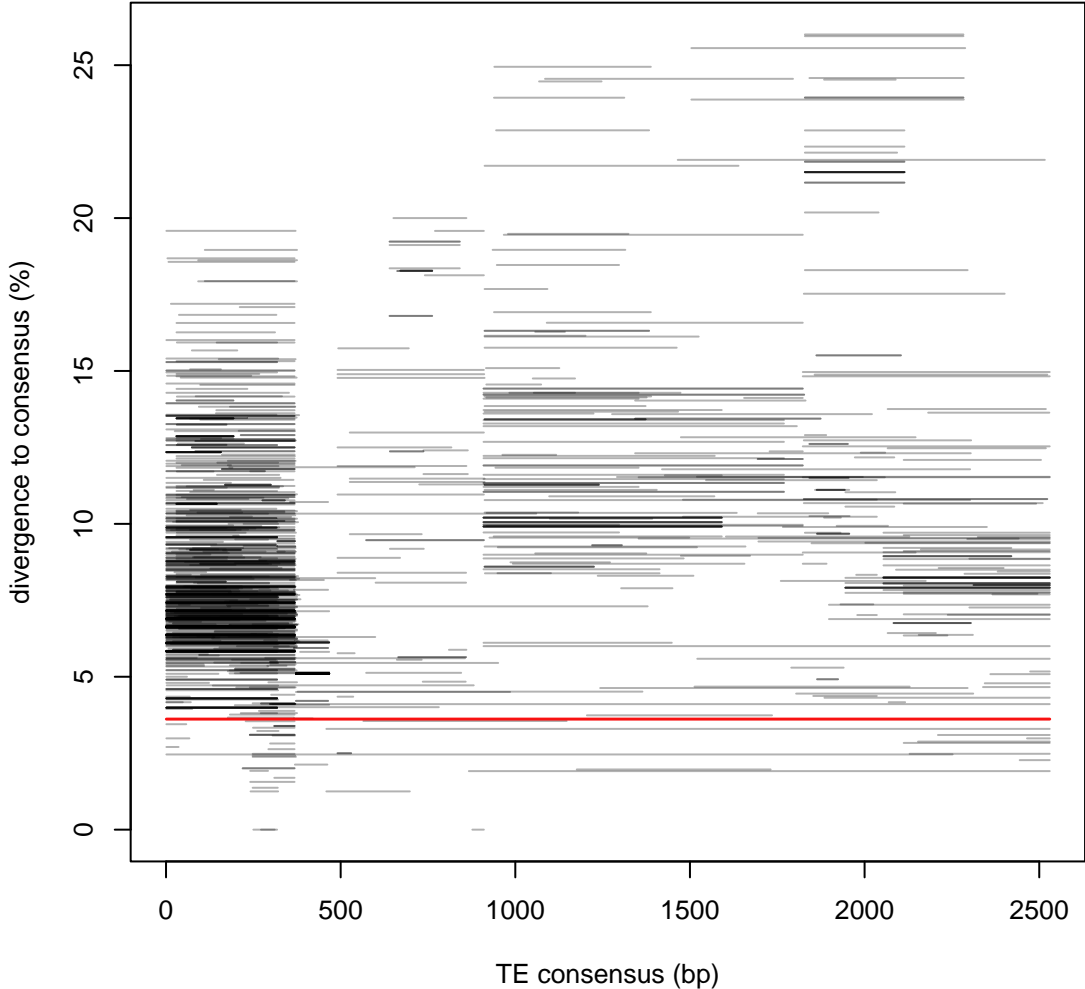
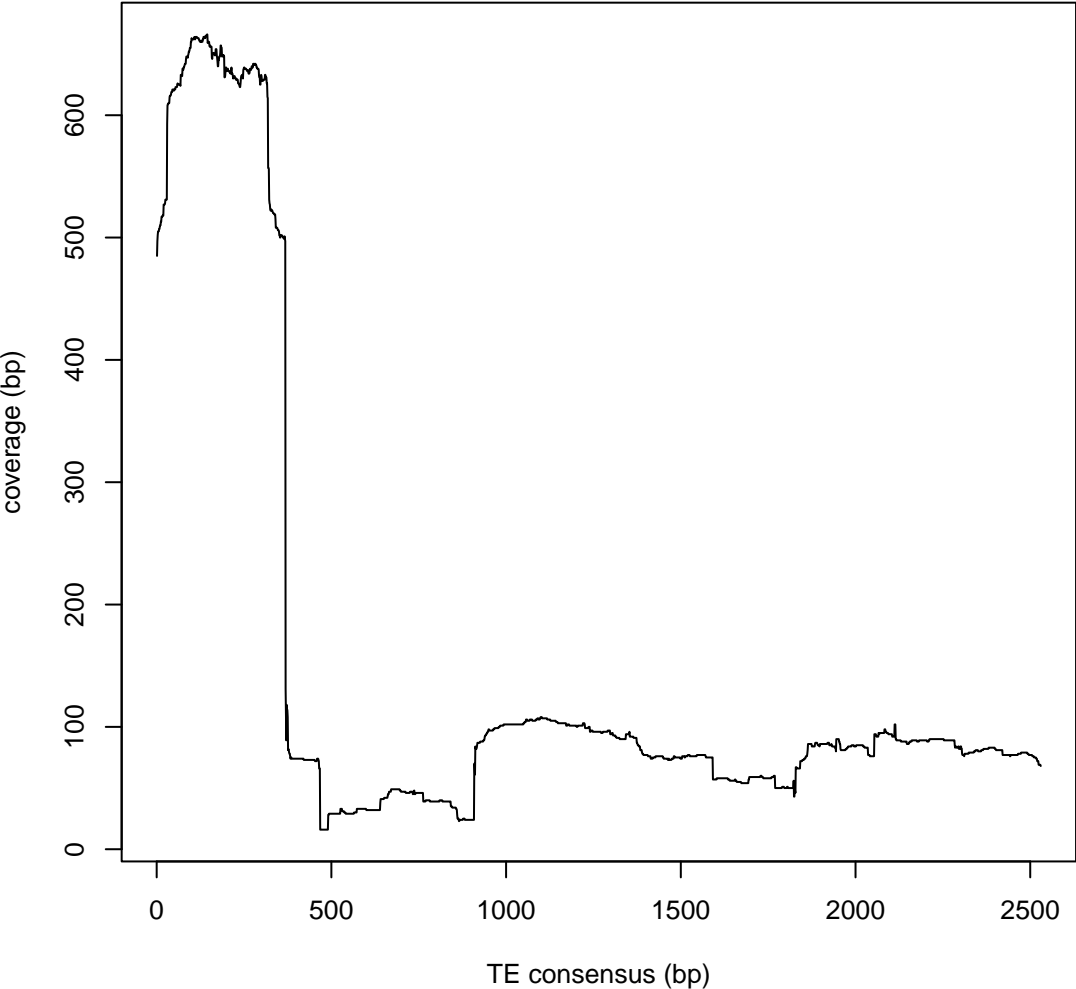


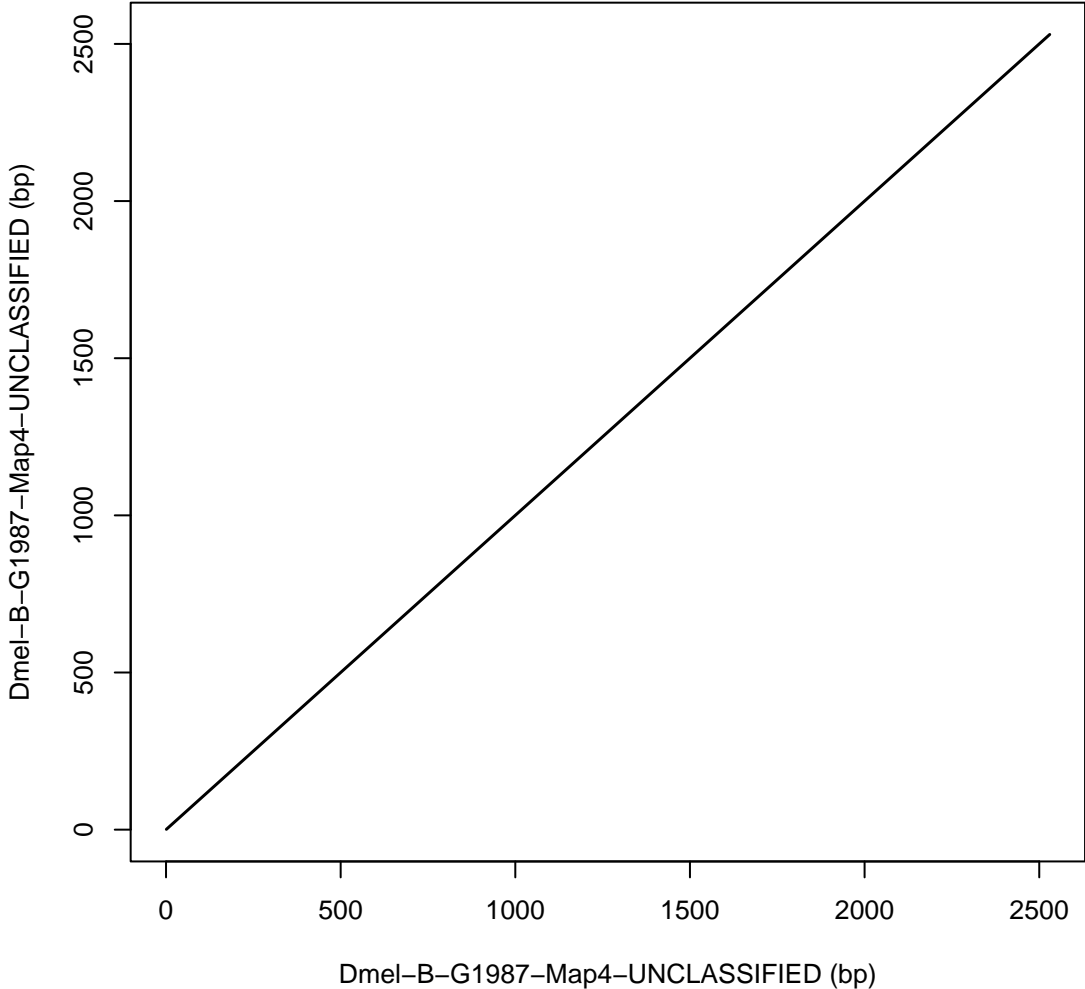
TE: Dmel-B-G1987-Map4-UNCLASSIFIED
consensus size: 2530bp; fragments: 1208; full length: 1 (>=2277bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

