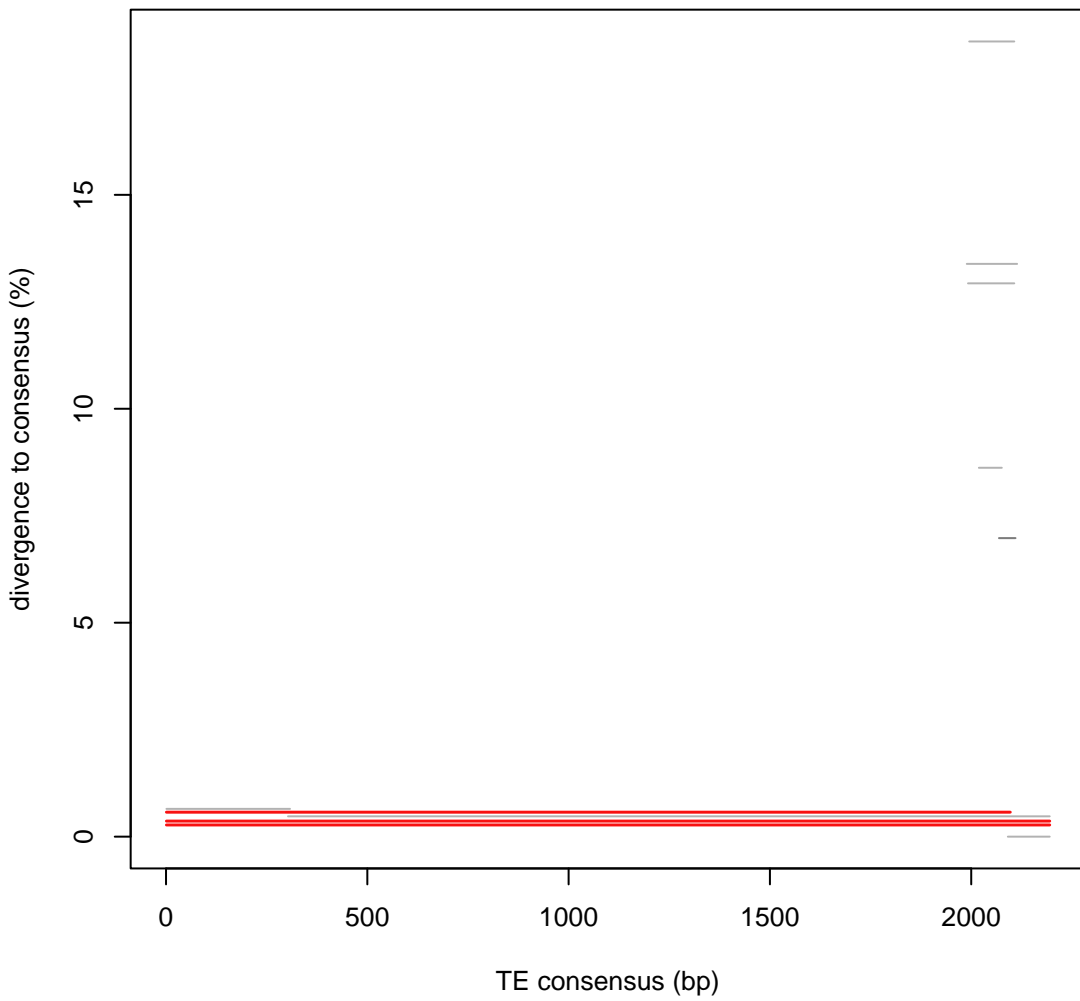
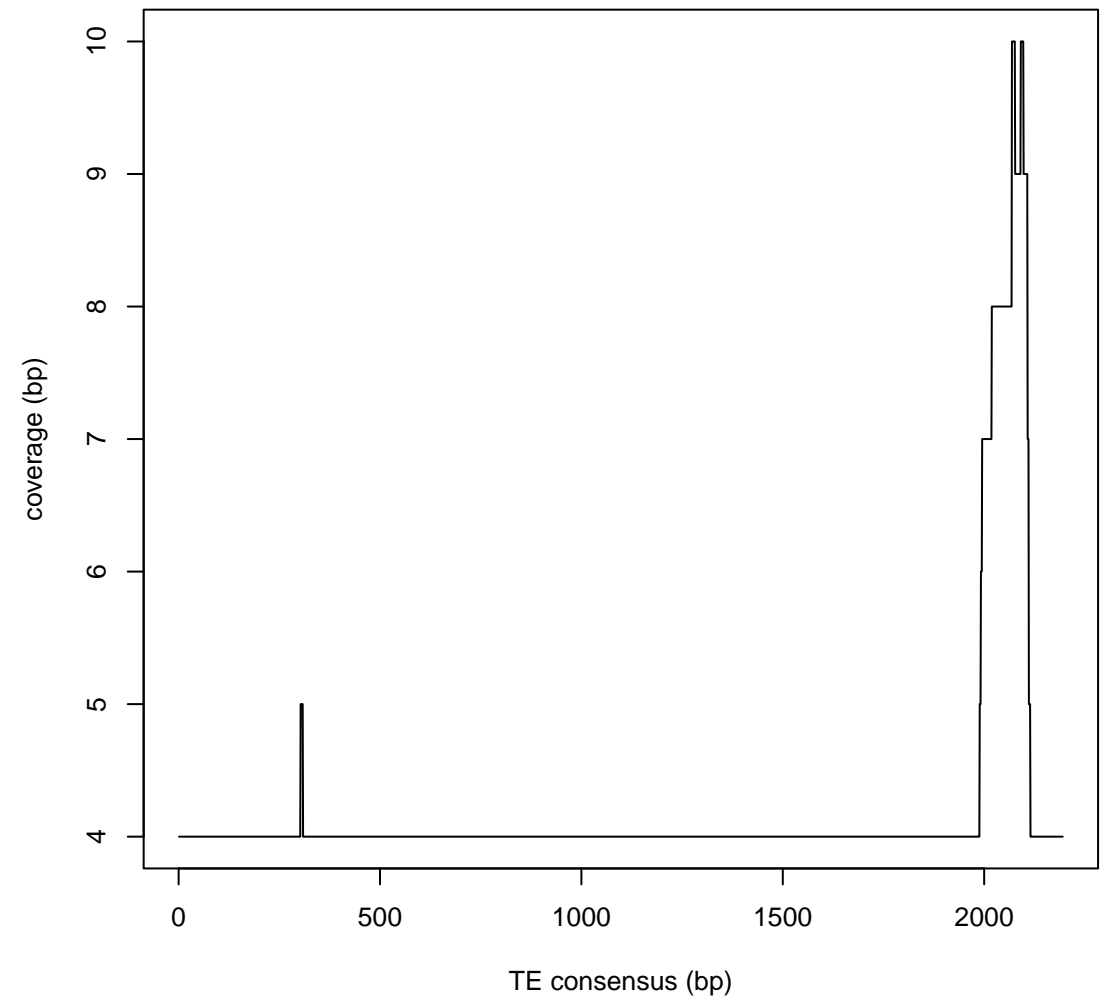


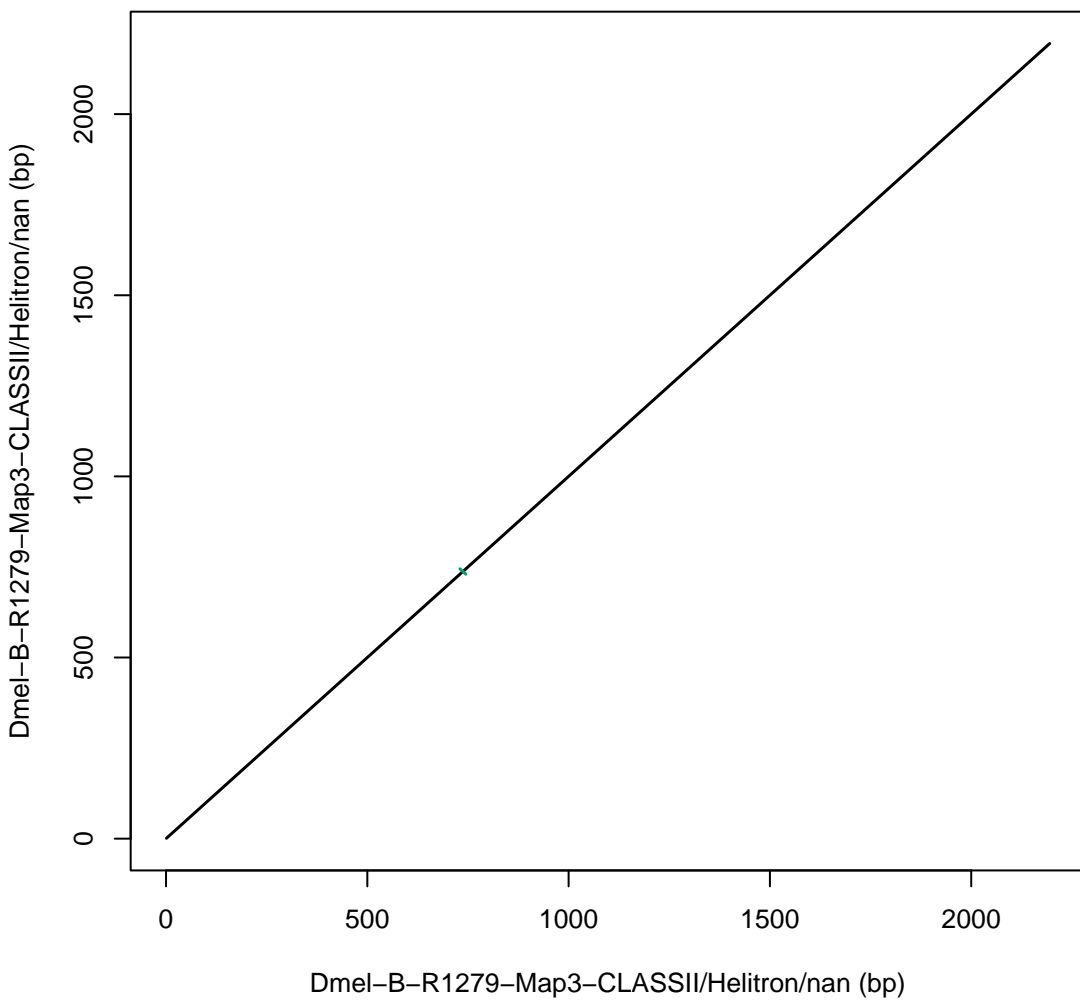
TE: Dmel-B-R1279-Map3-CLASSII/Helitron/nan
consensus size: 2195bp; fragments: 12; full length: 3 (≥ 1975.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

