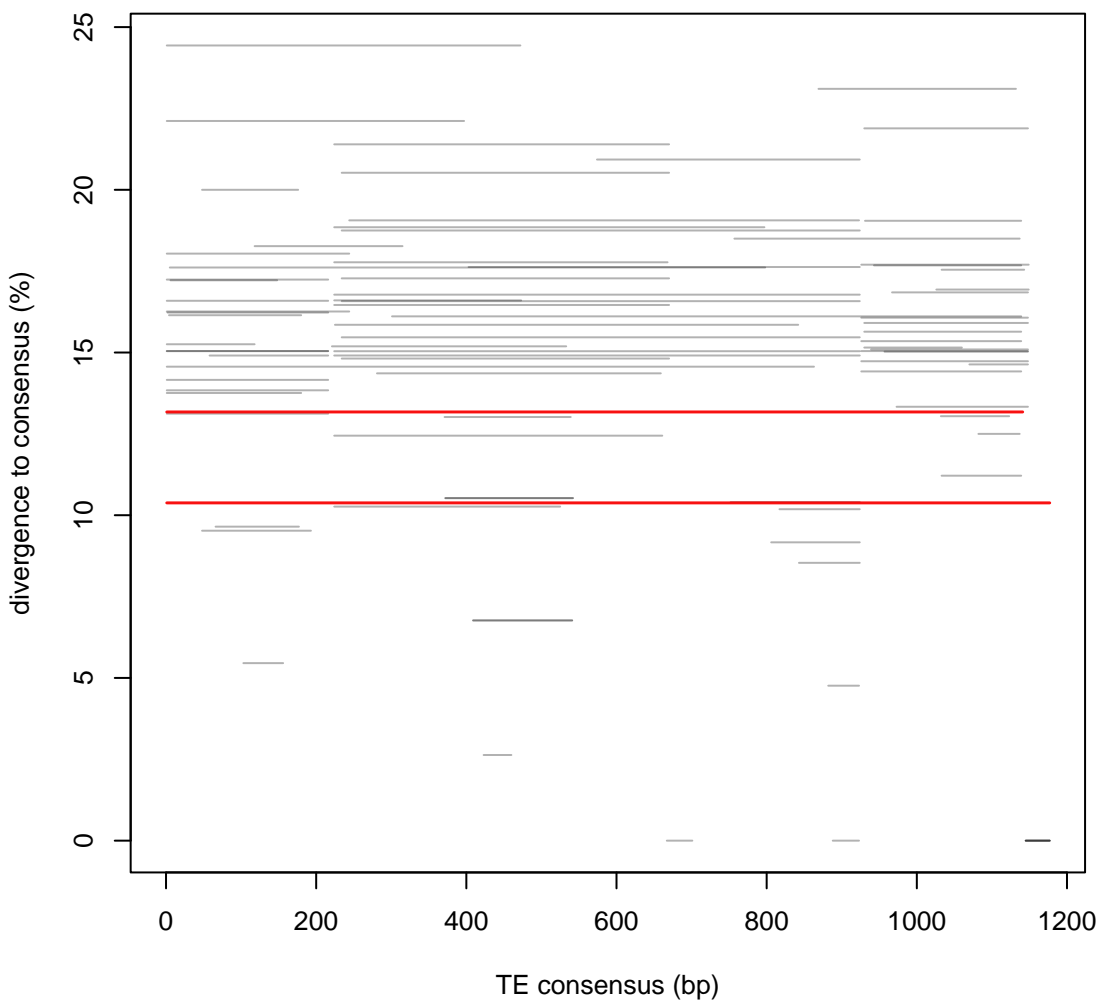
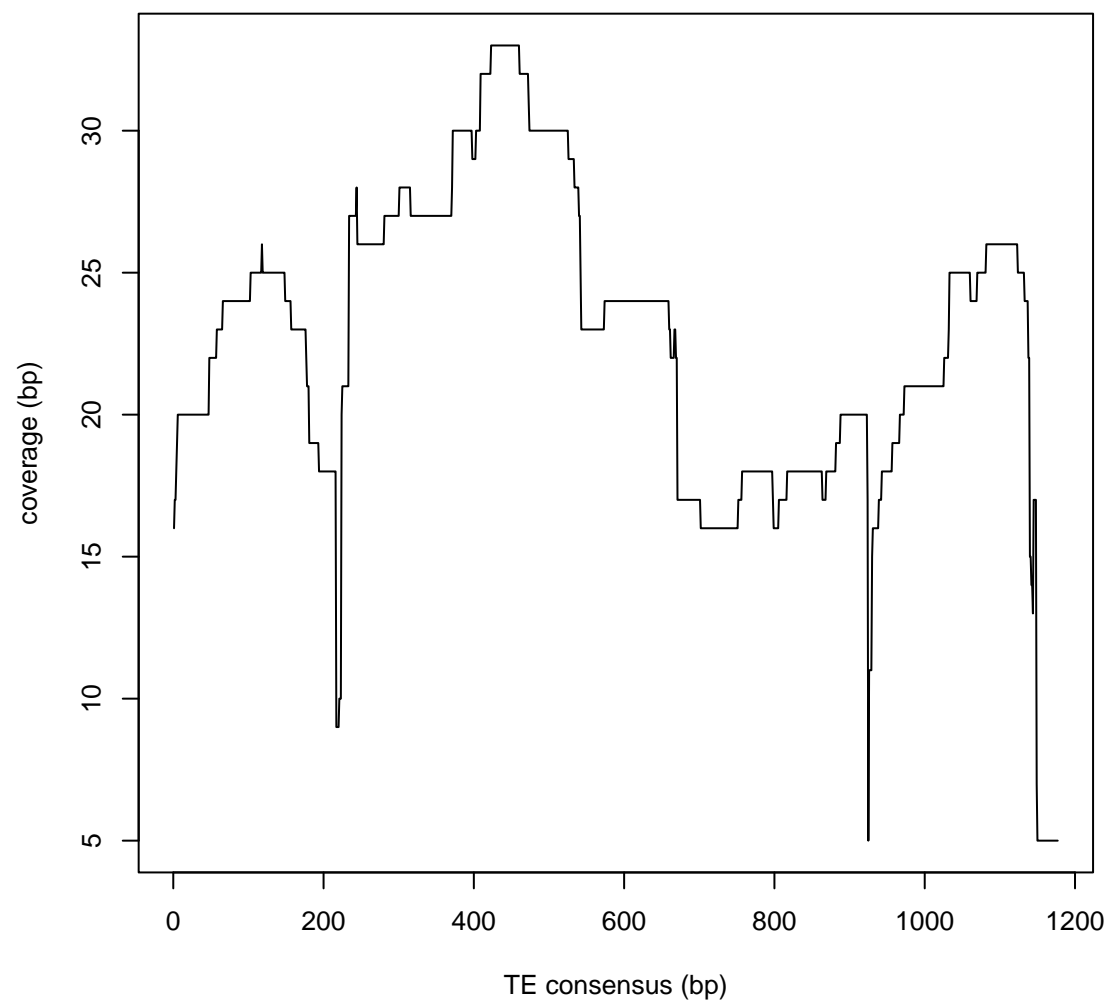


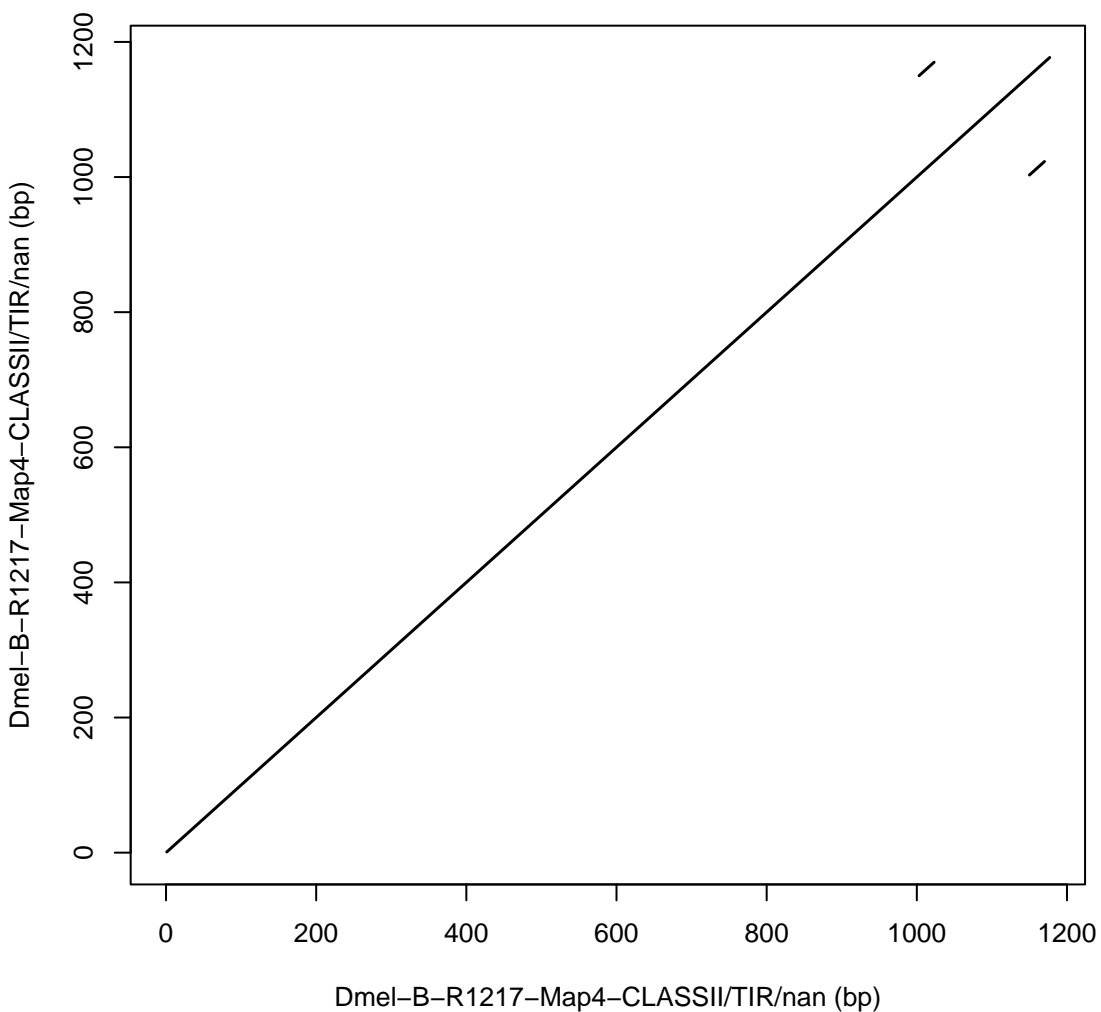
TE: Dmel-B-R1217-Map4-CLASSII/TIR/nan
consensus size: 1177bp; fragments: 90; full length: 2 (>=1059.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

