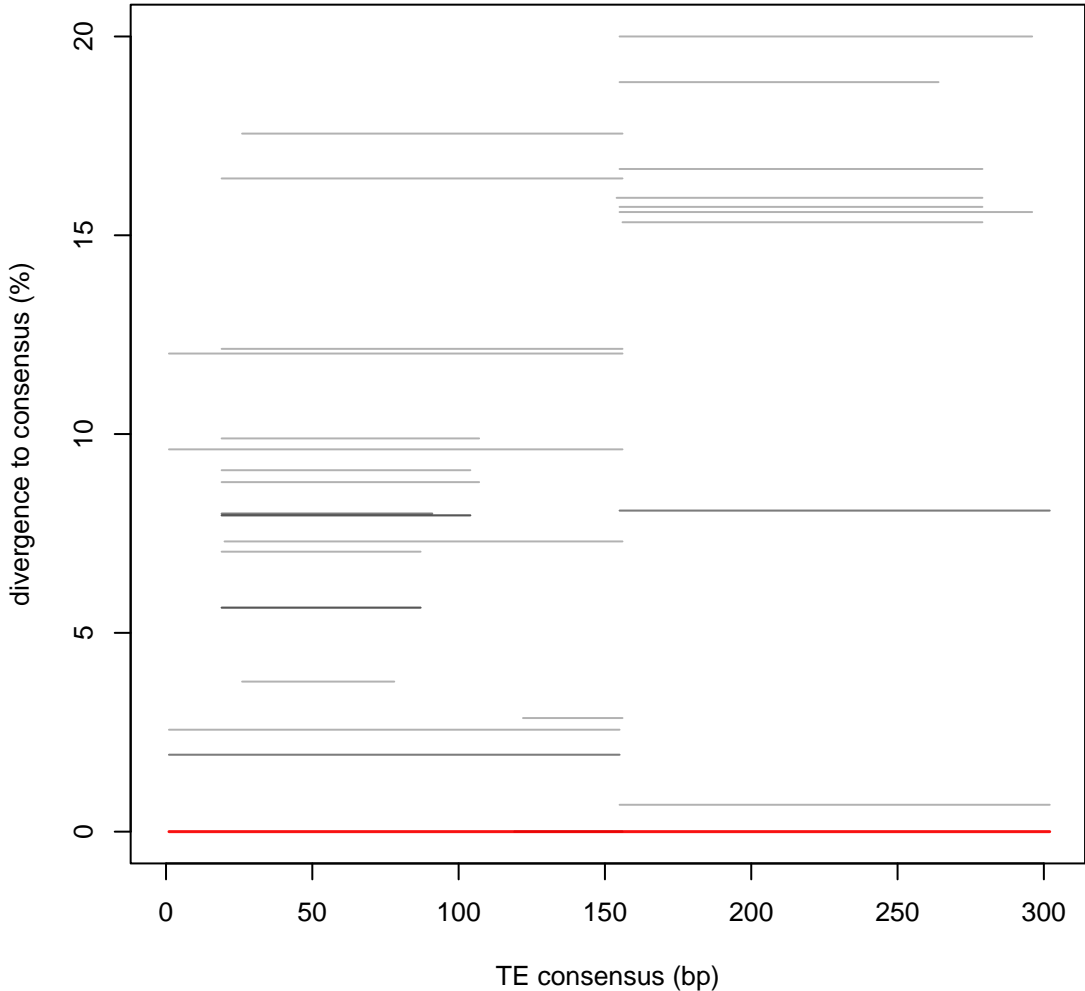
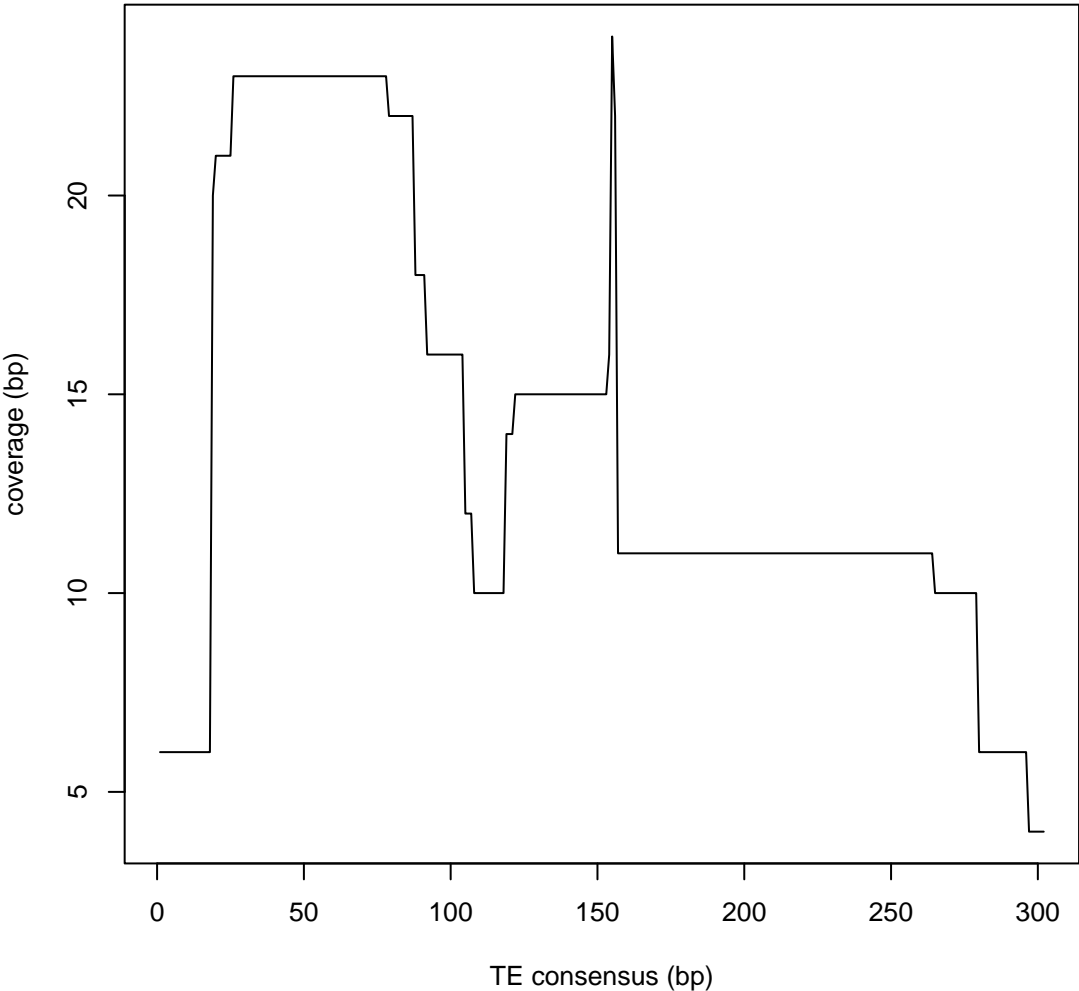


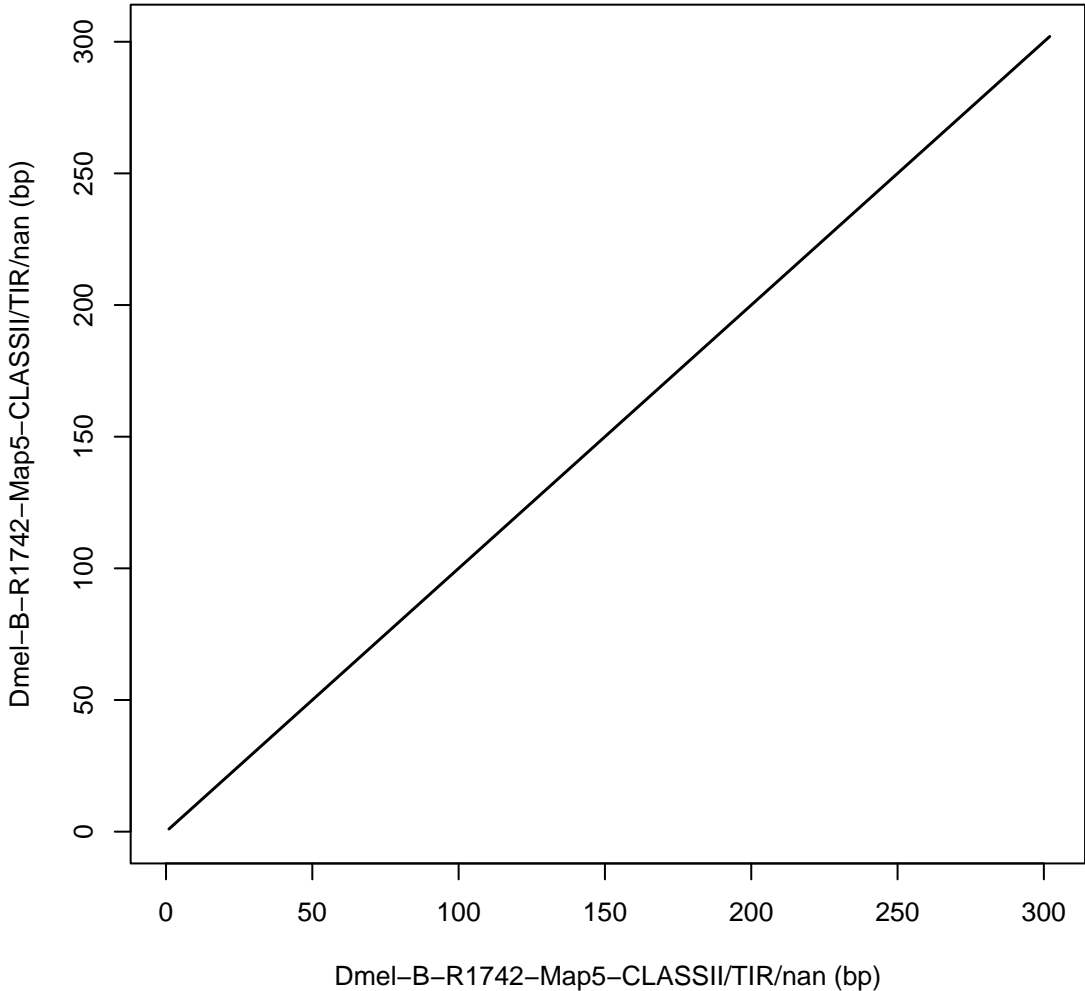
TE: Dmel-B-R1742-Map5-CLASSII/TIR/nan
consensus size: 302bp; fragments: 38; full length: 1 (≥ 271.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

