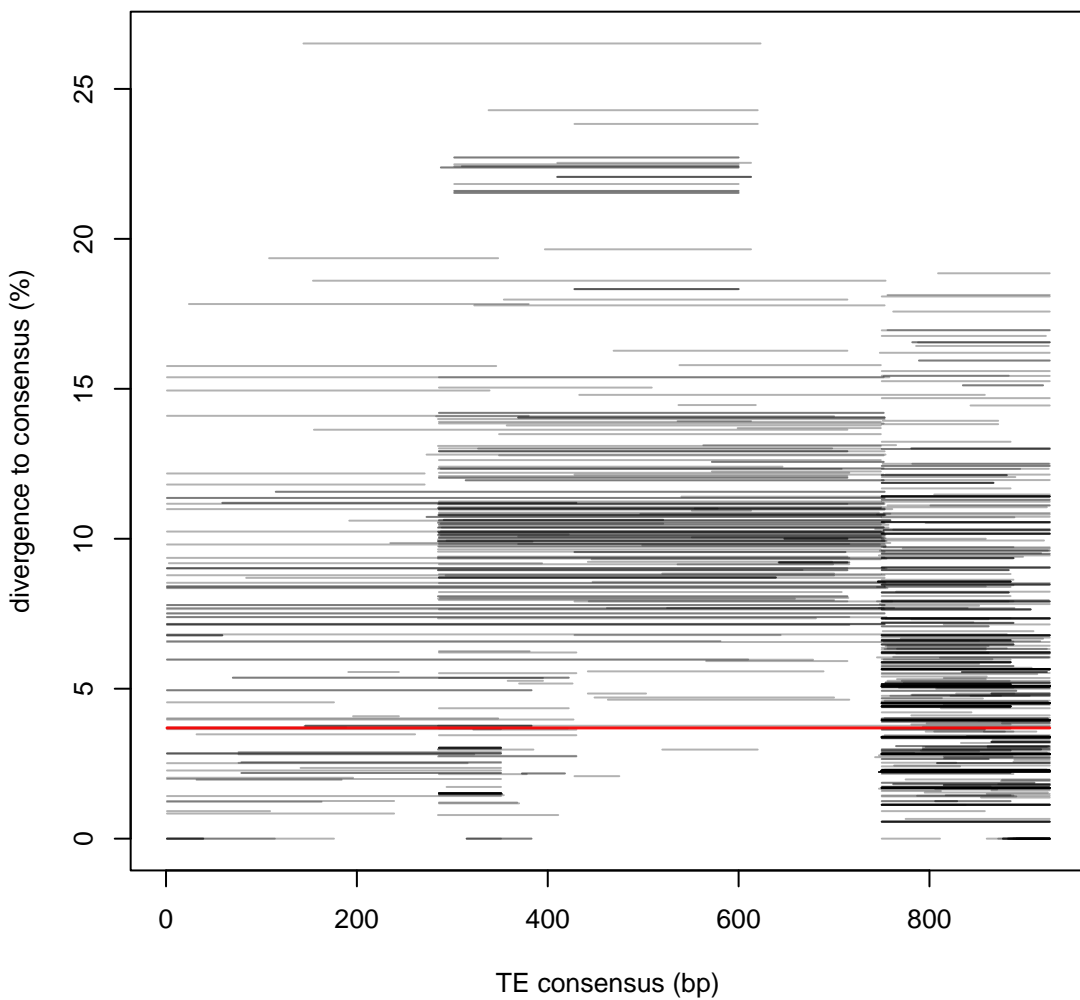
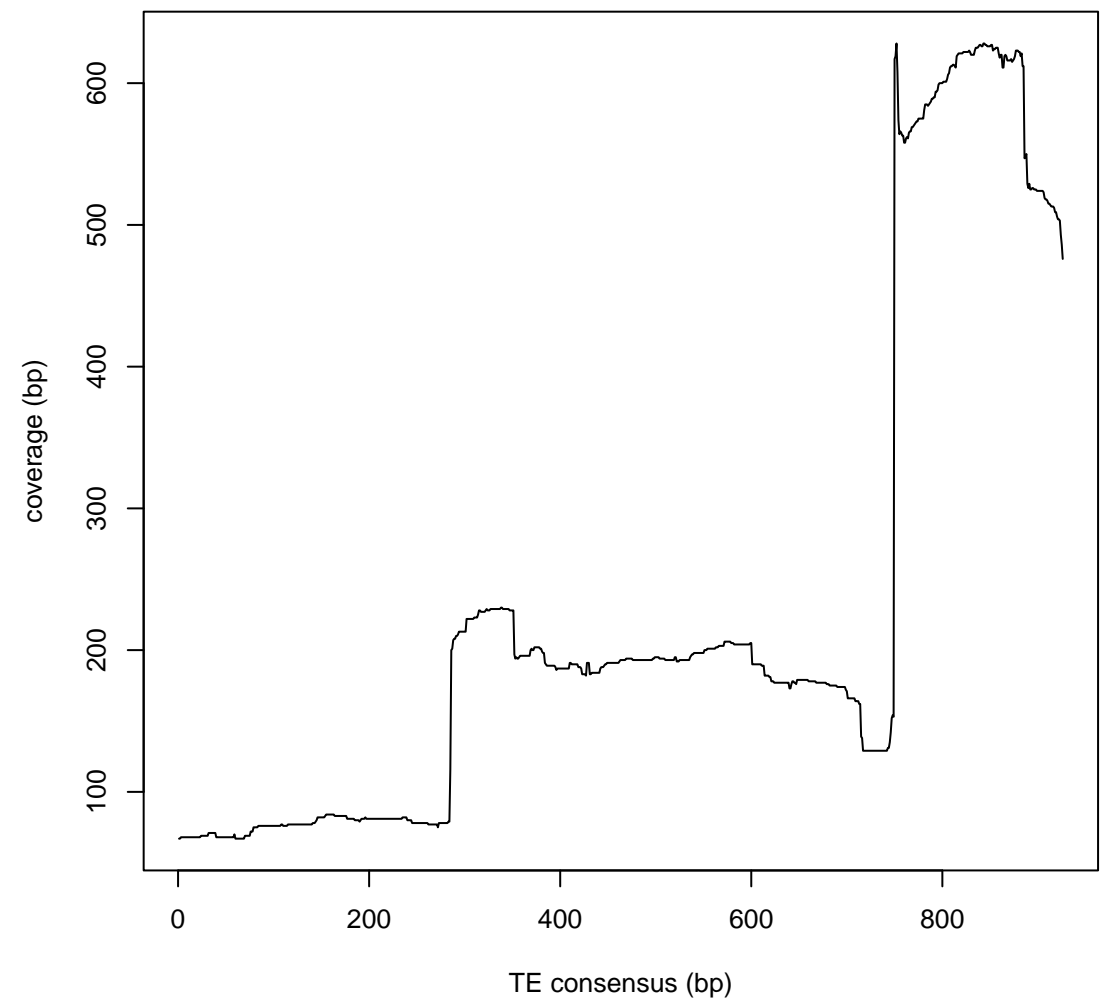


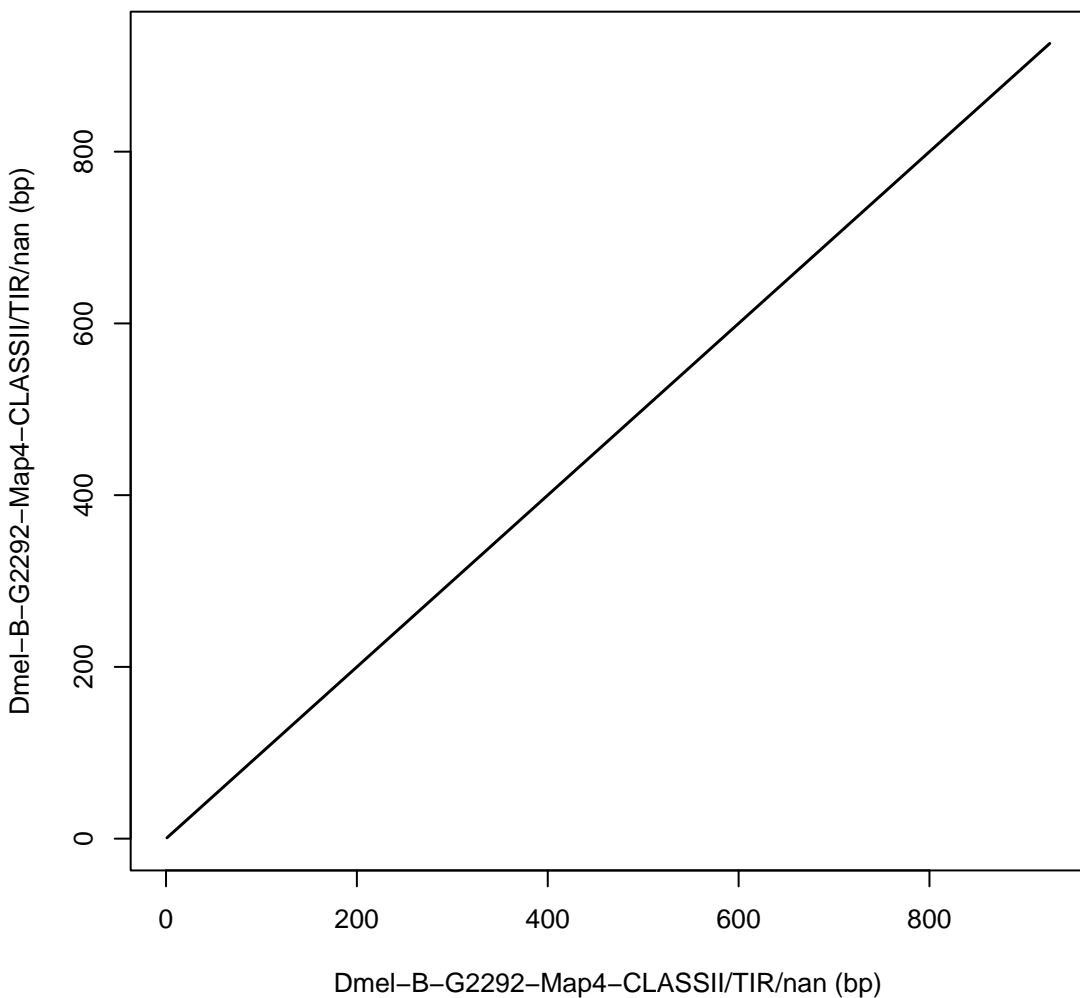
TE: Dmel-B-G2292-Map4-CLASSII/TIR/nan
consensus size: 926bp; fragments: 1000; full length: 1 (>=833.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

