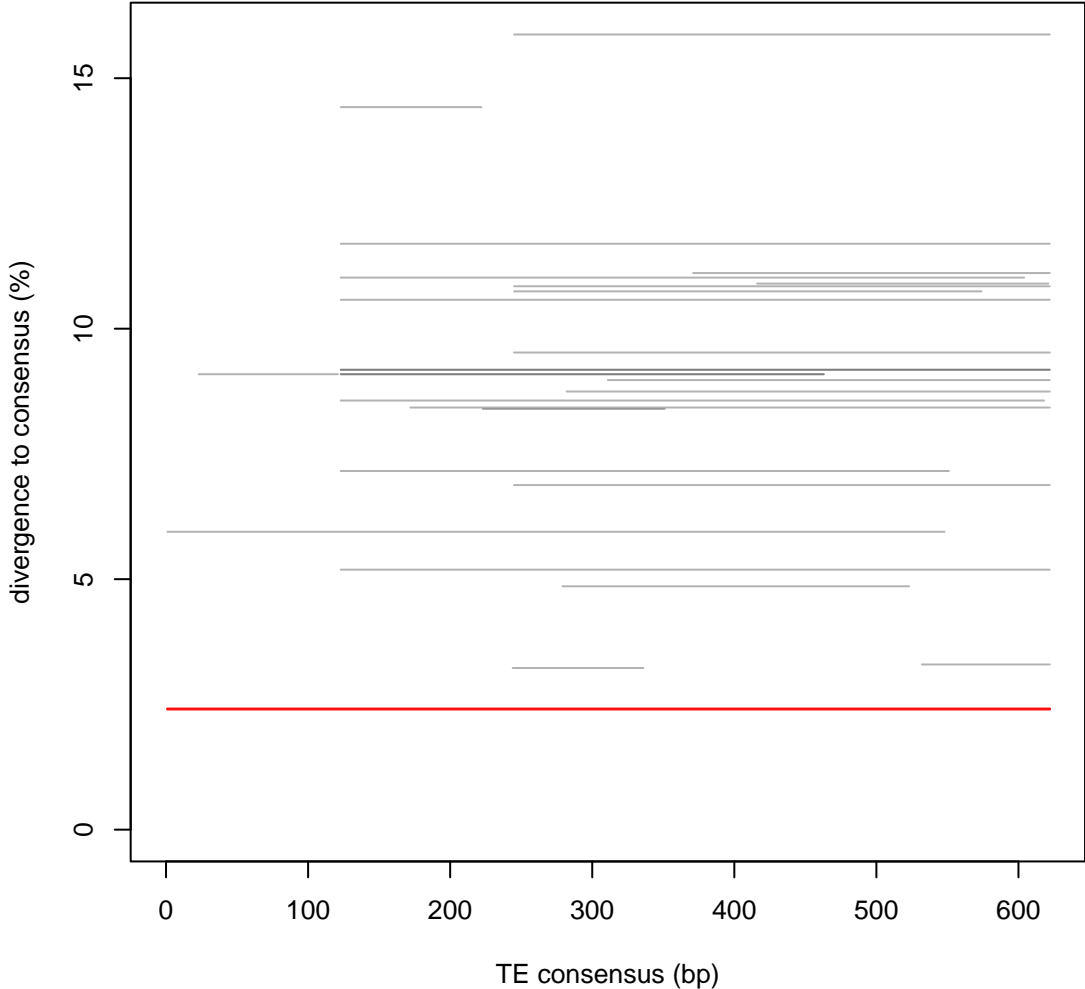
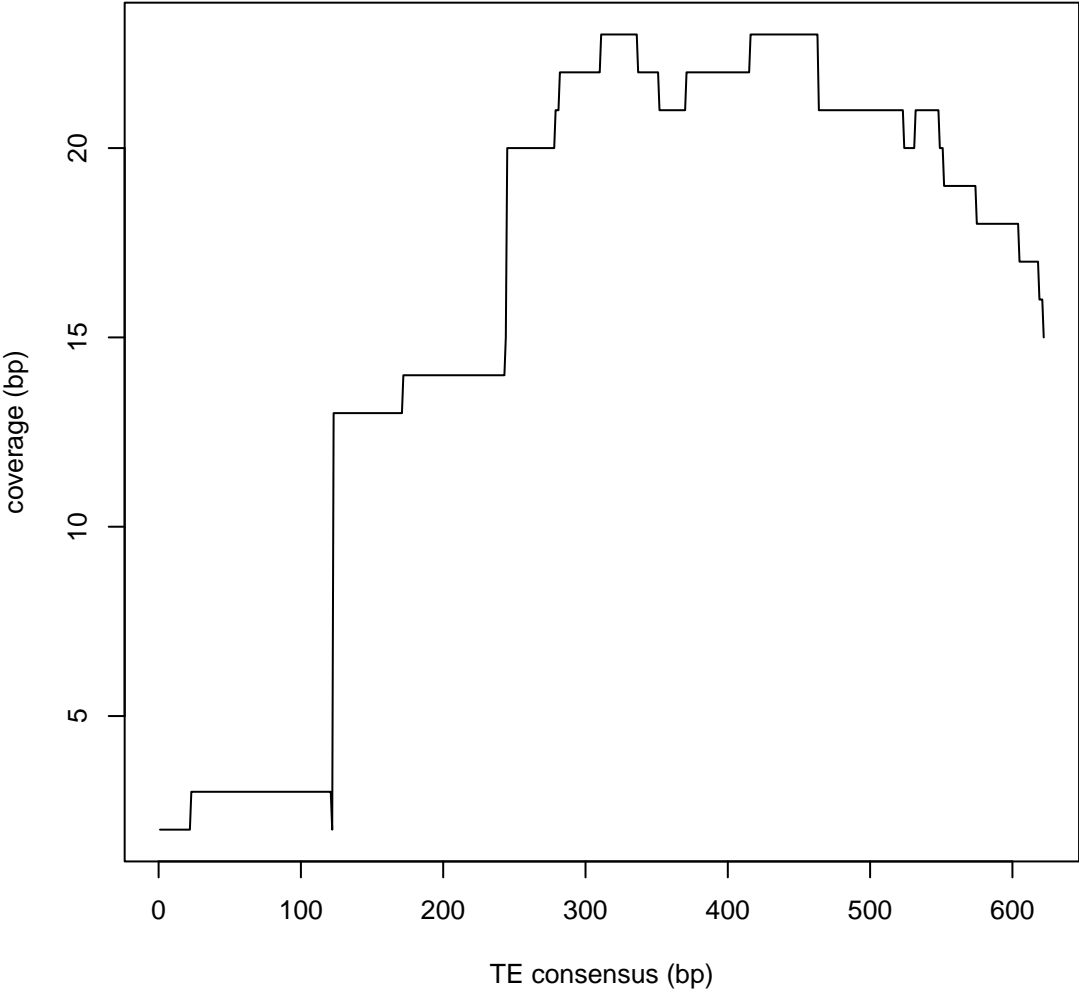


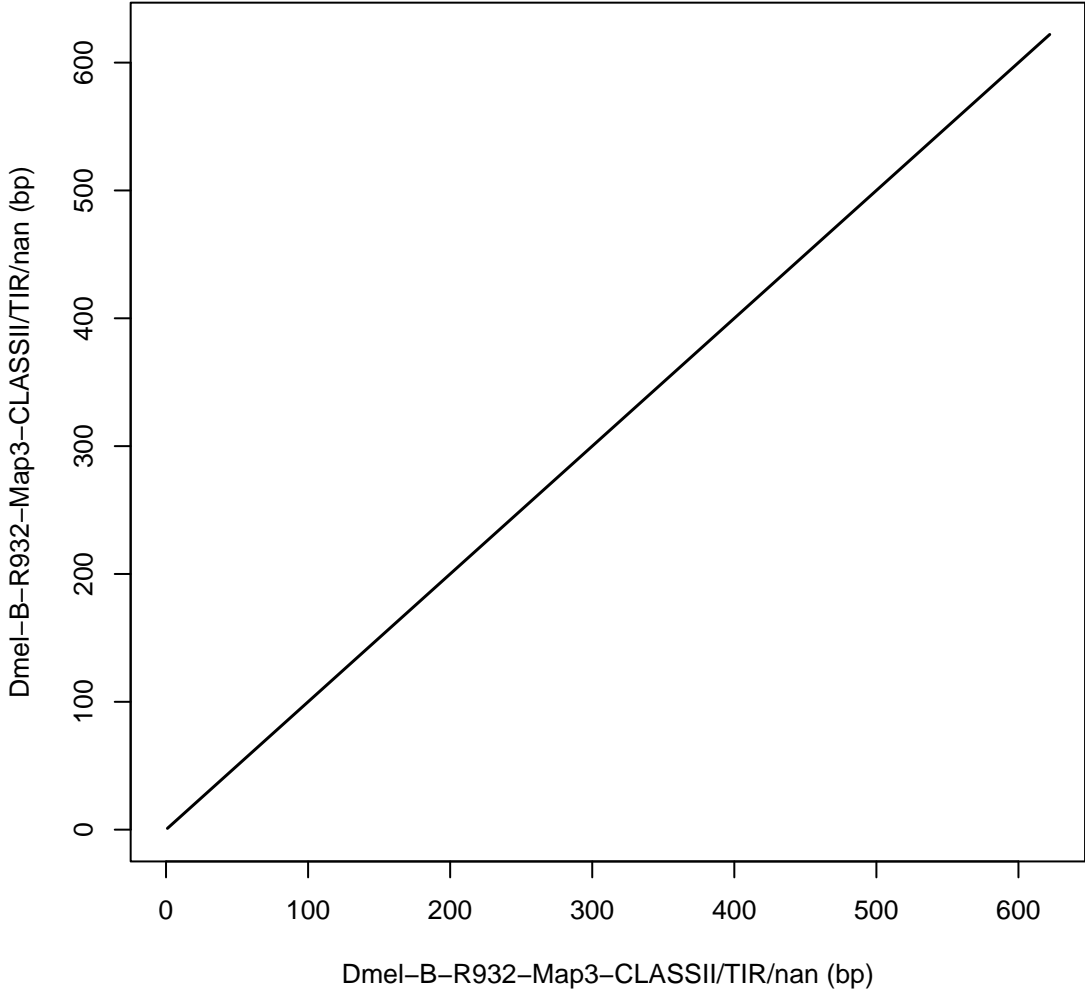
TE: Dmel-B-R932-Map3-CLASSII/TIR/nan
consensus size: 622bp; fragments: 28; full length: 1 (>=559.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

