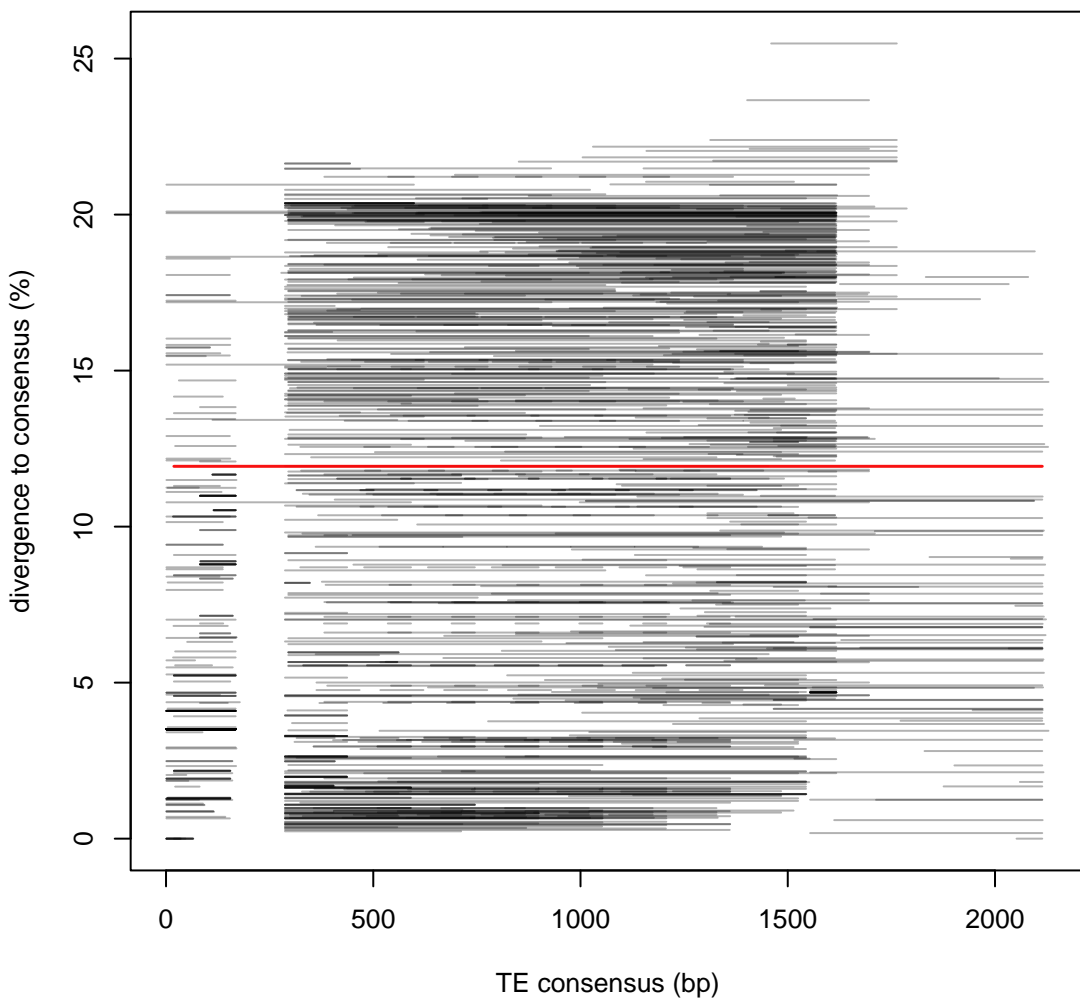
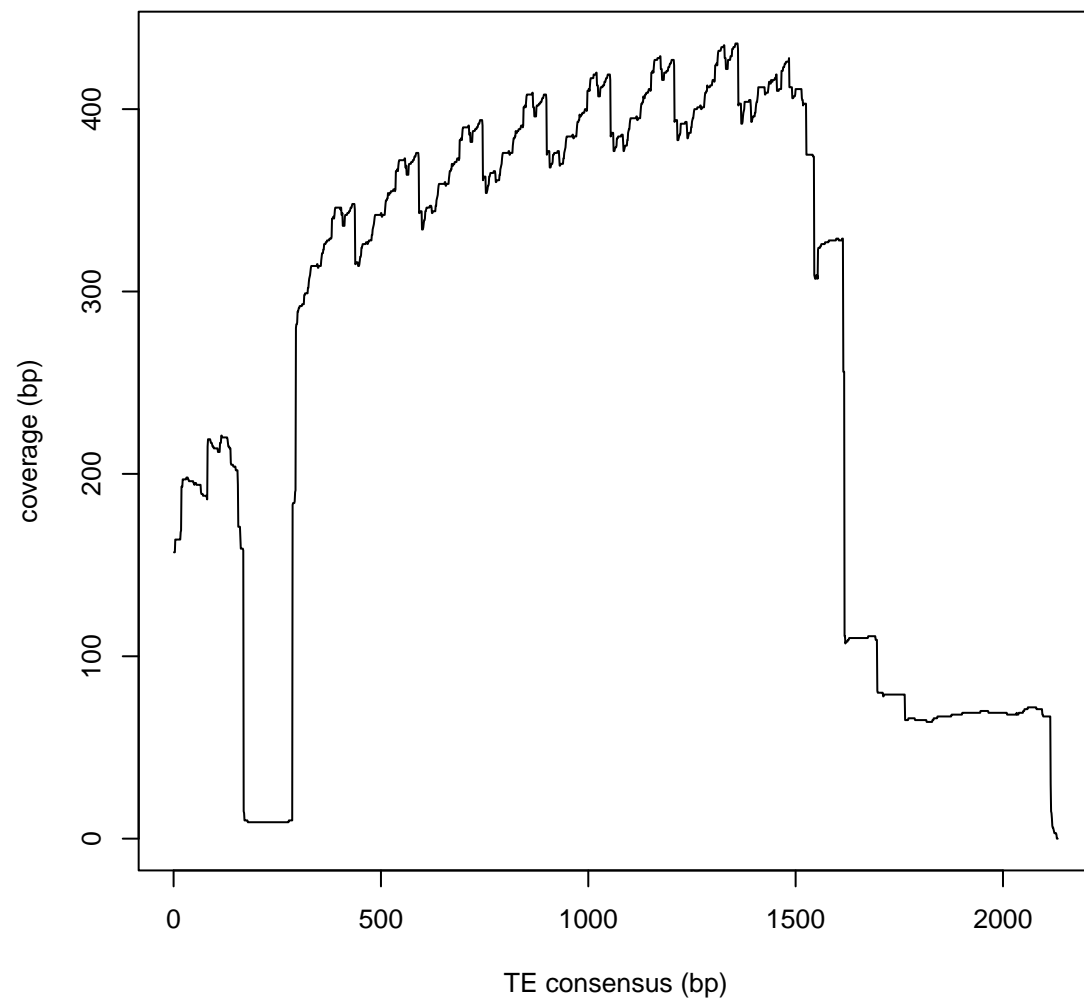


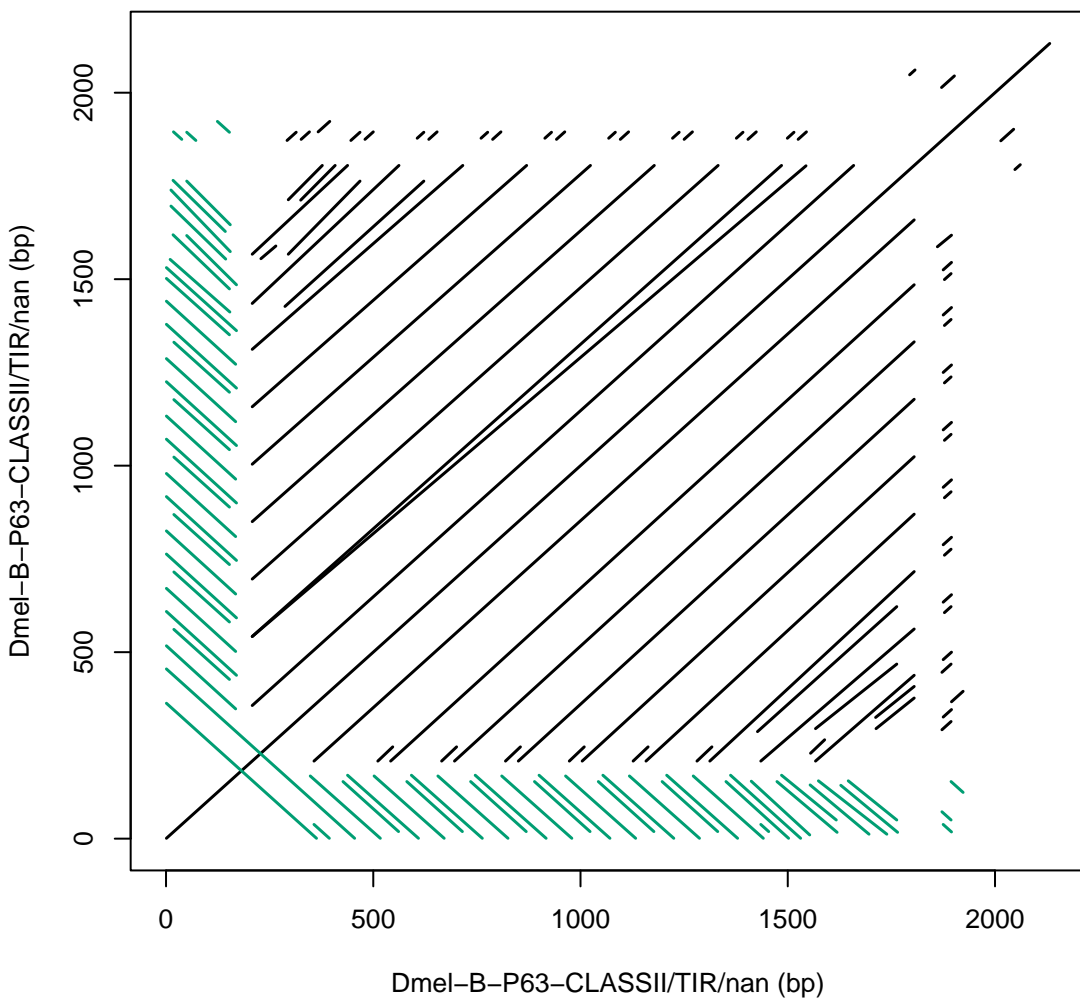
TE: Dmel-B-P63-CLASSII/TIR/nan
consensus size: 2132bp; fragments: 1274; full length: 1 (≥ 1918.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

