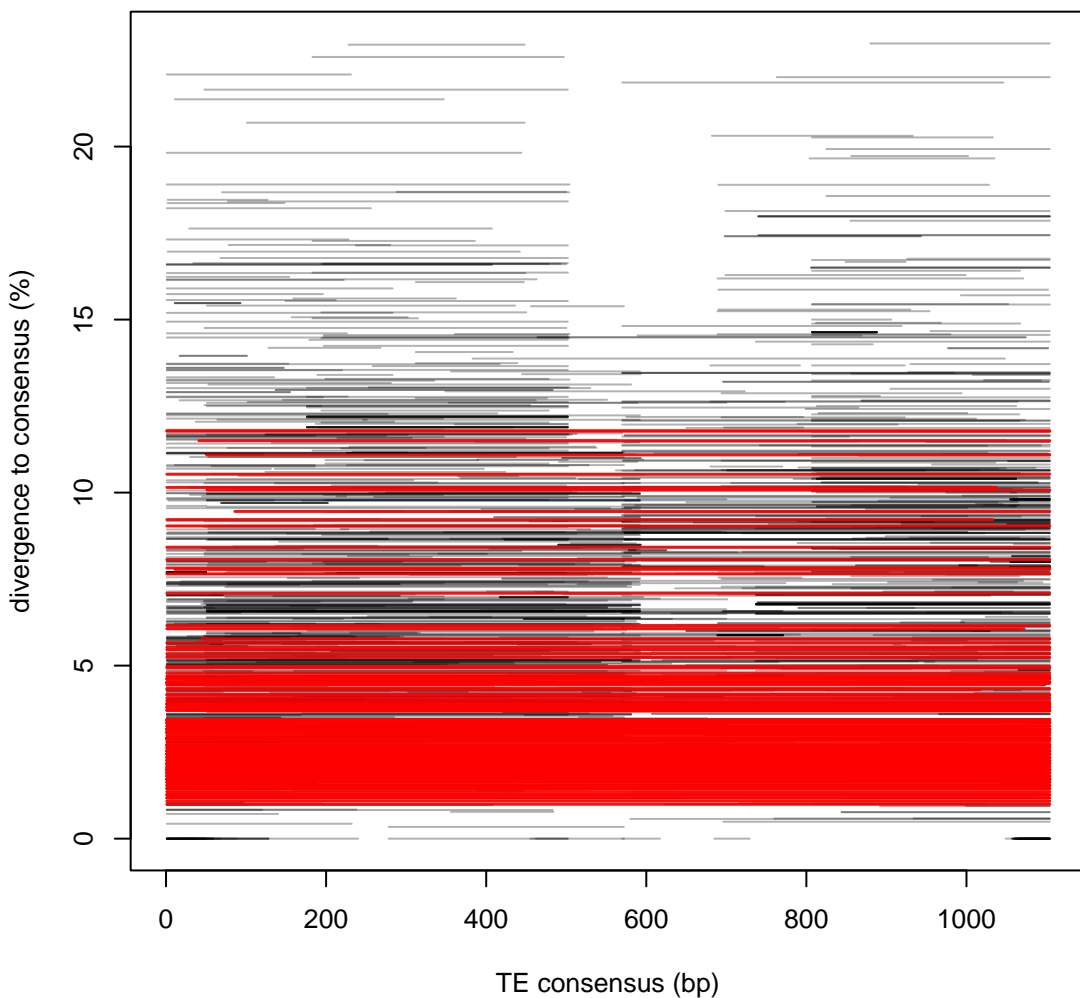
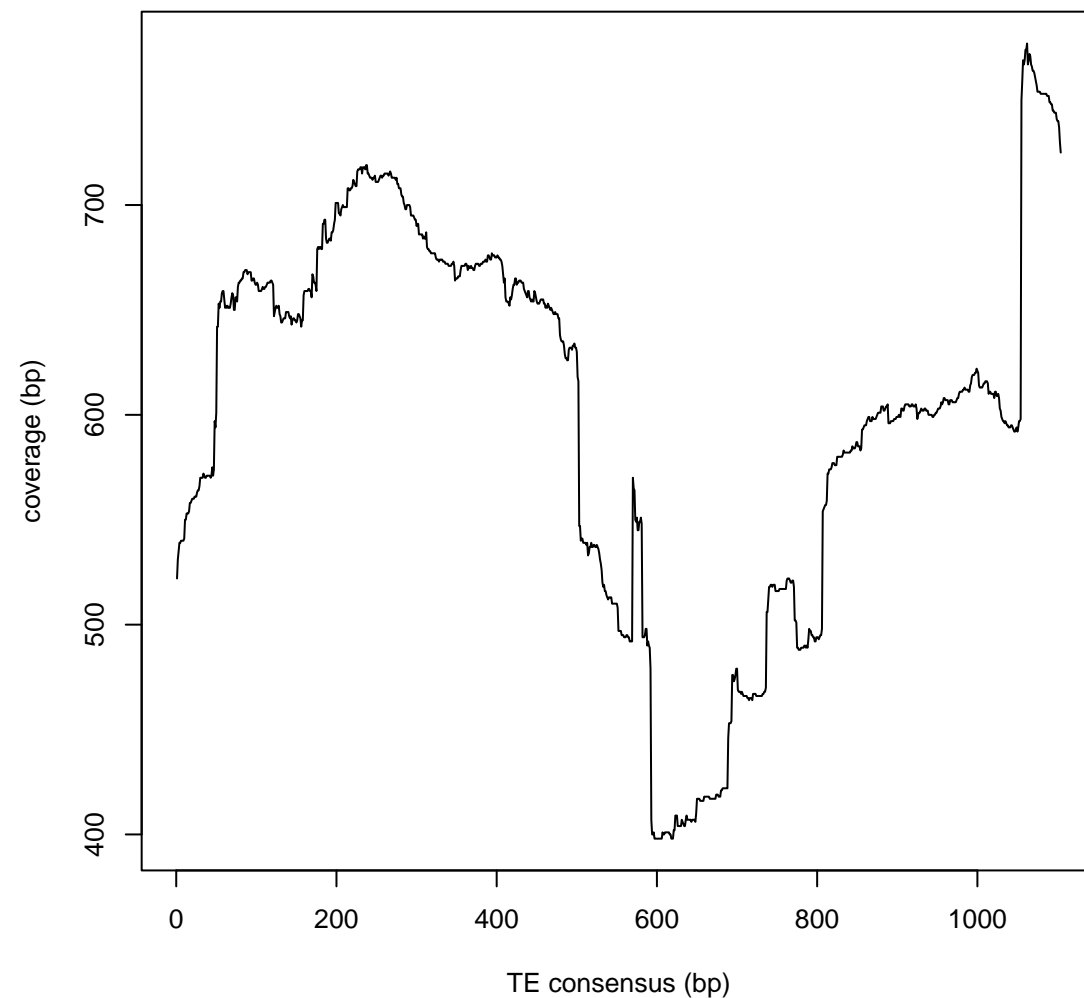


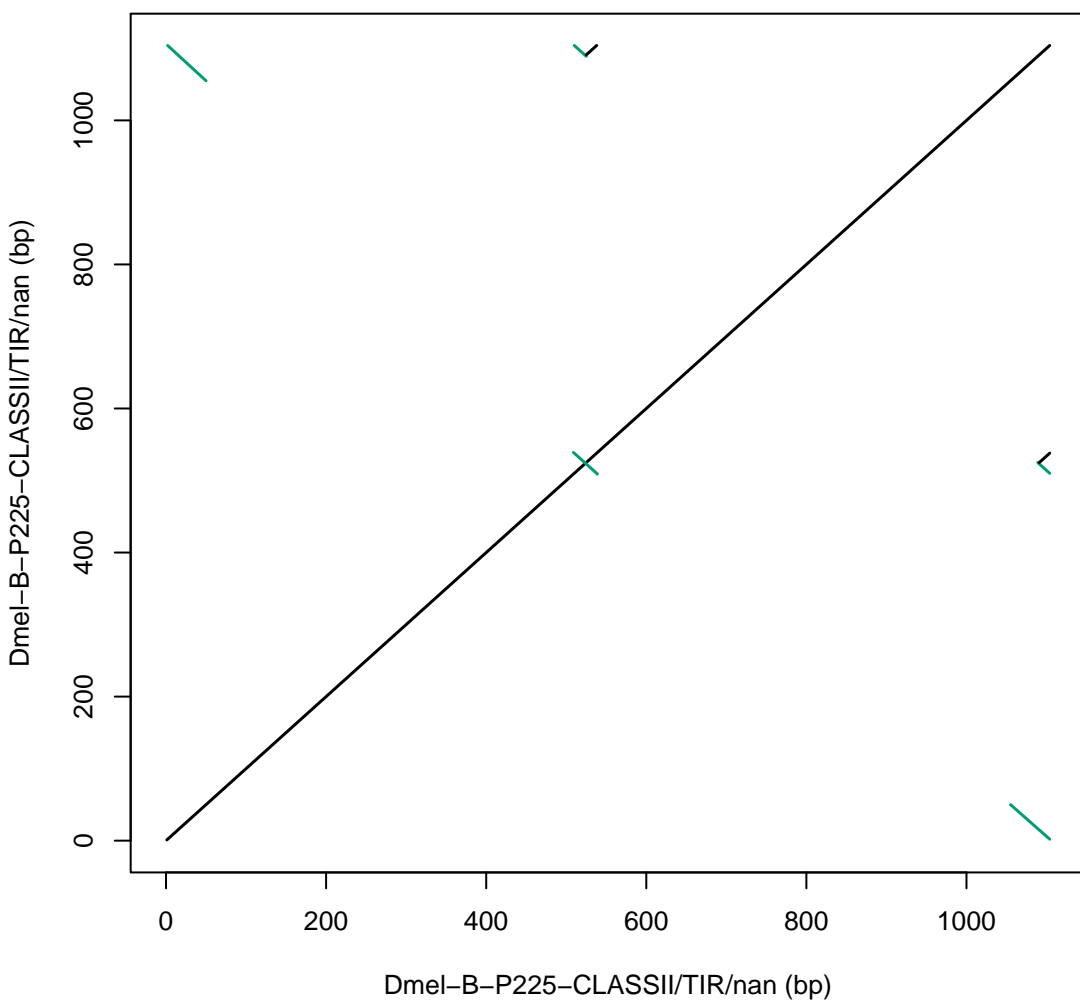
TE: Dmel-B-P225-CLASSII/TIR/nan
consensus size: 1104bp; fragments: 1827; full length: 165 (≥ 993.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

