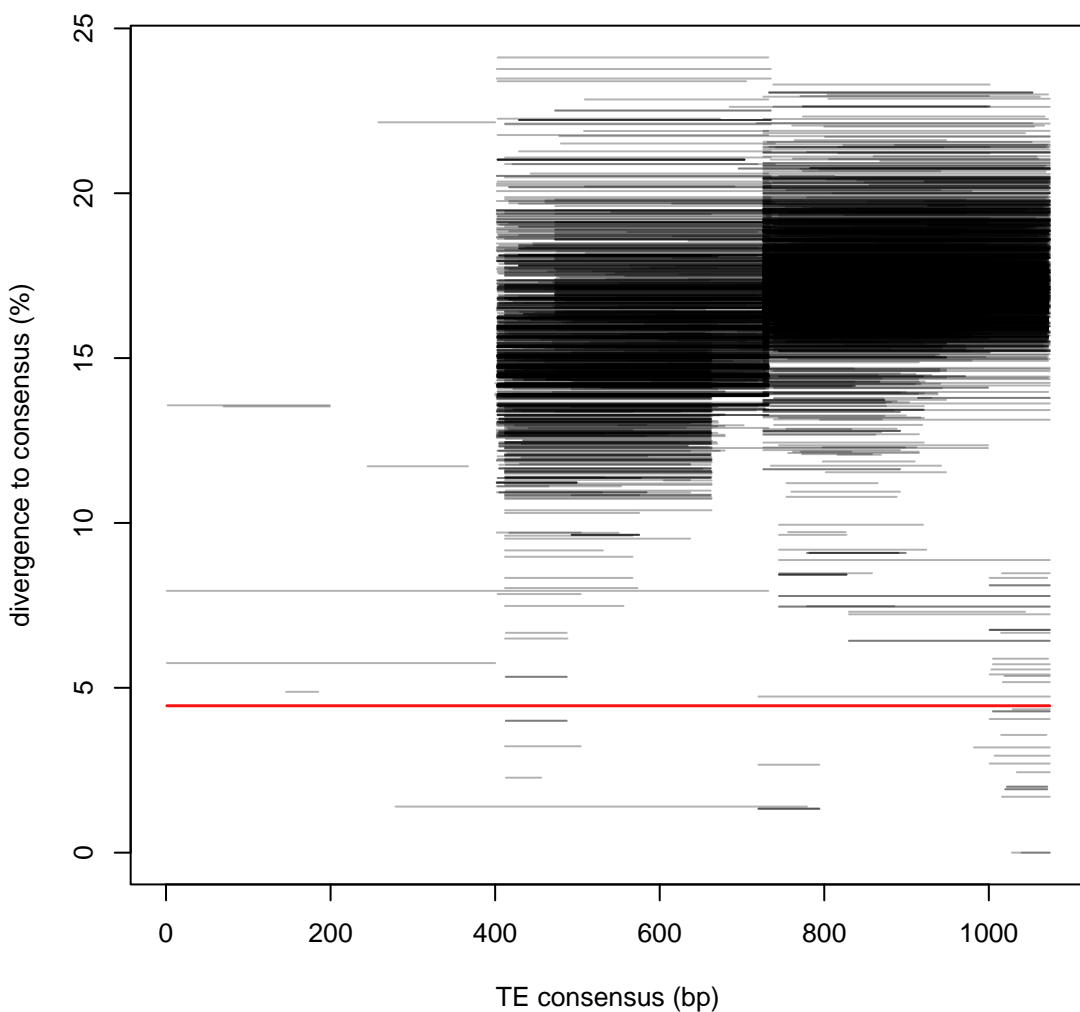
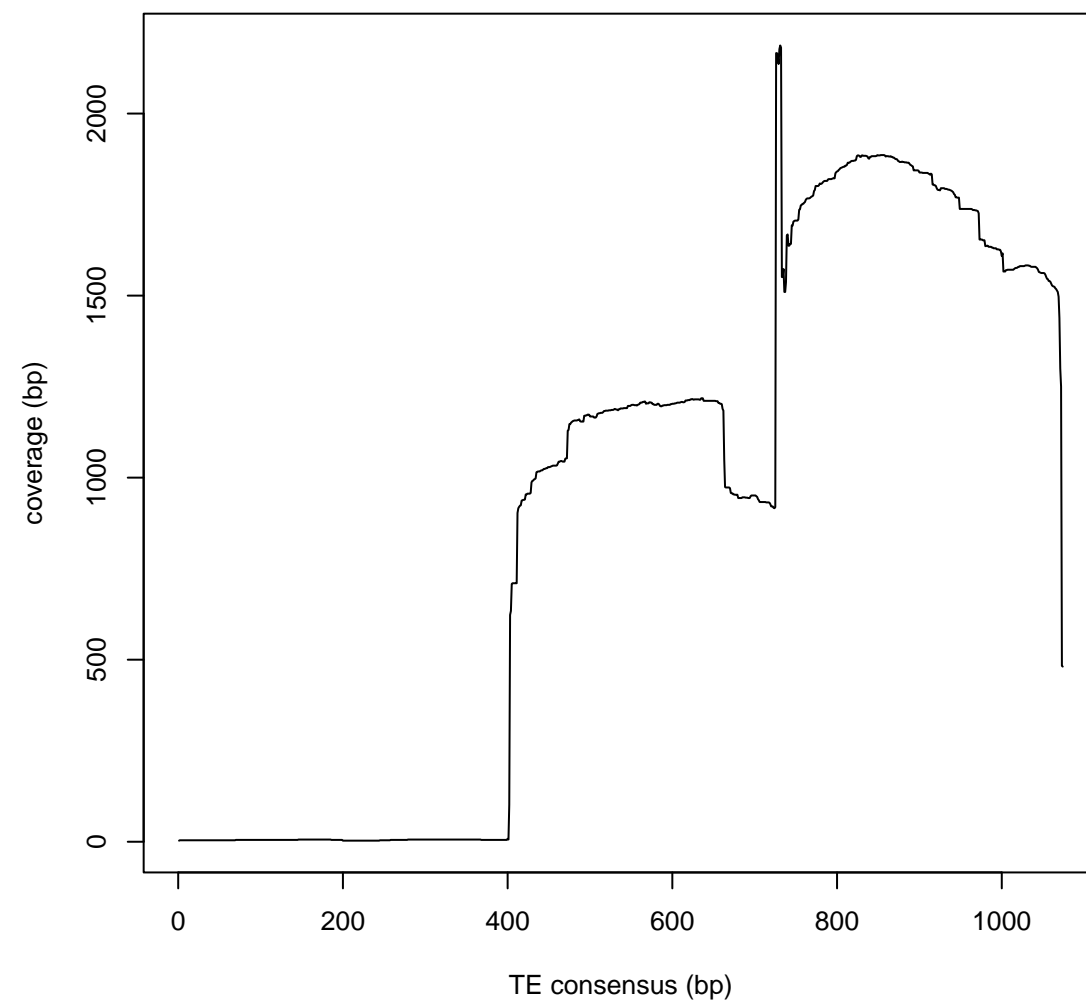


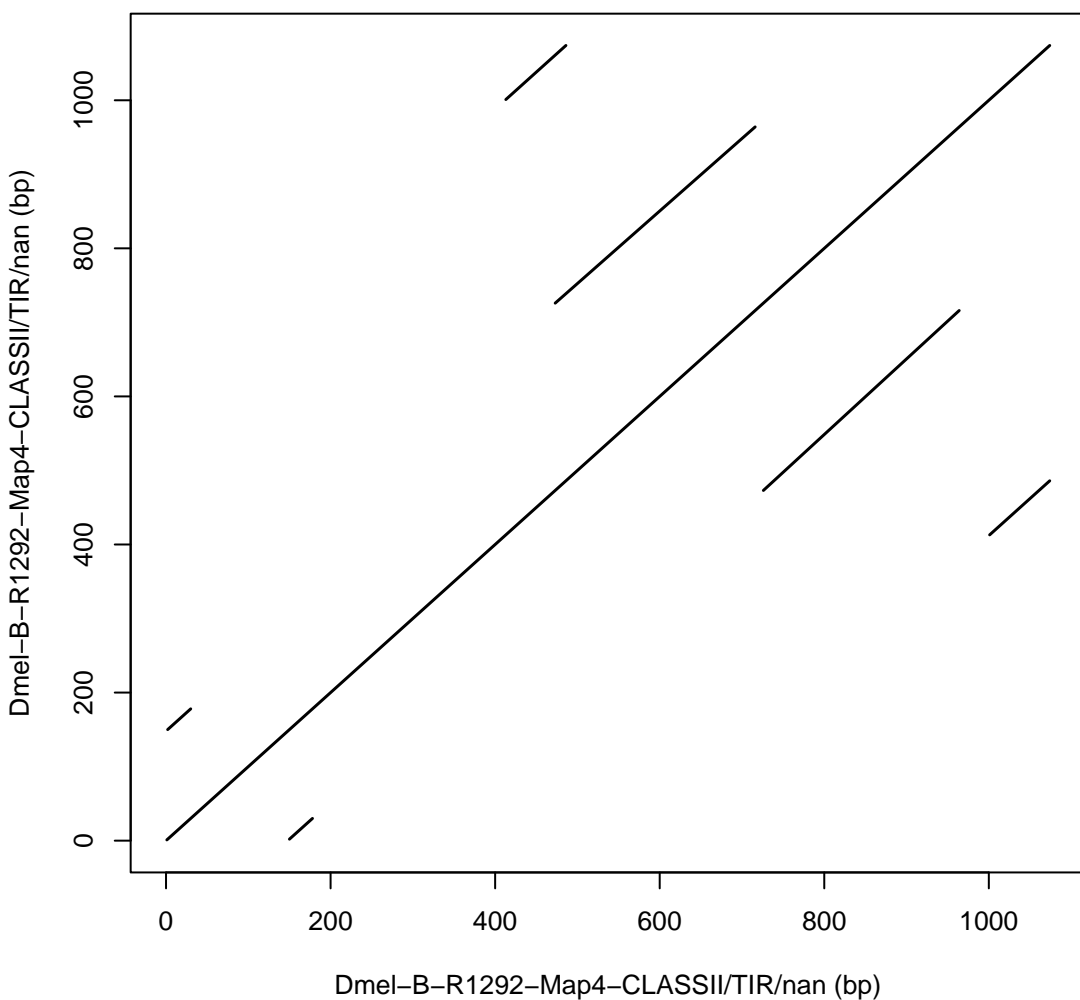
TE: Dmel-B-R1292-Map4-CLASSII/TIR/nan
consensus size: 1074bp; fragments: 3145; full length: 1 (>=966.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

