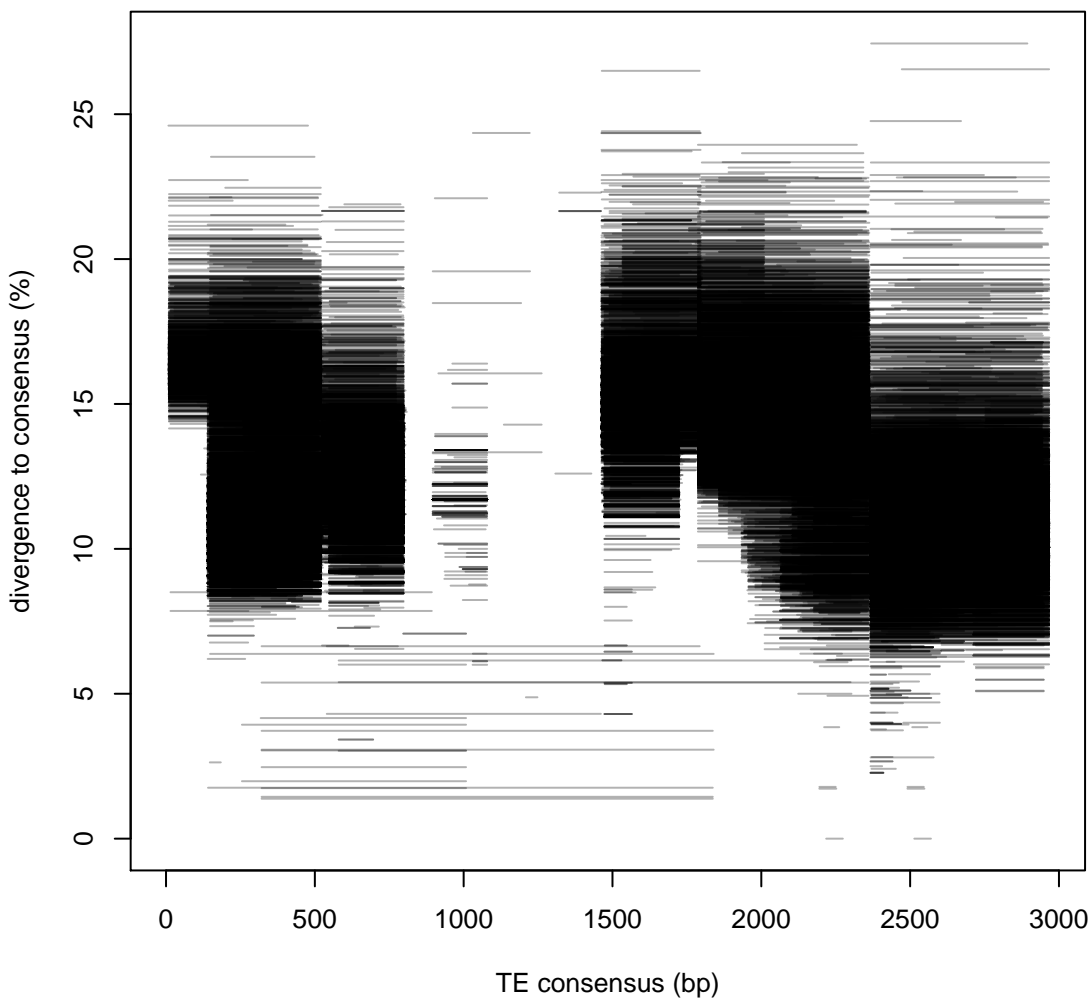
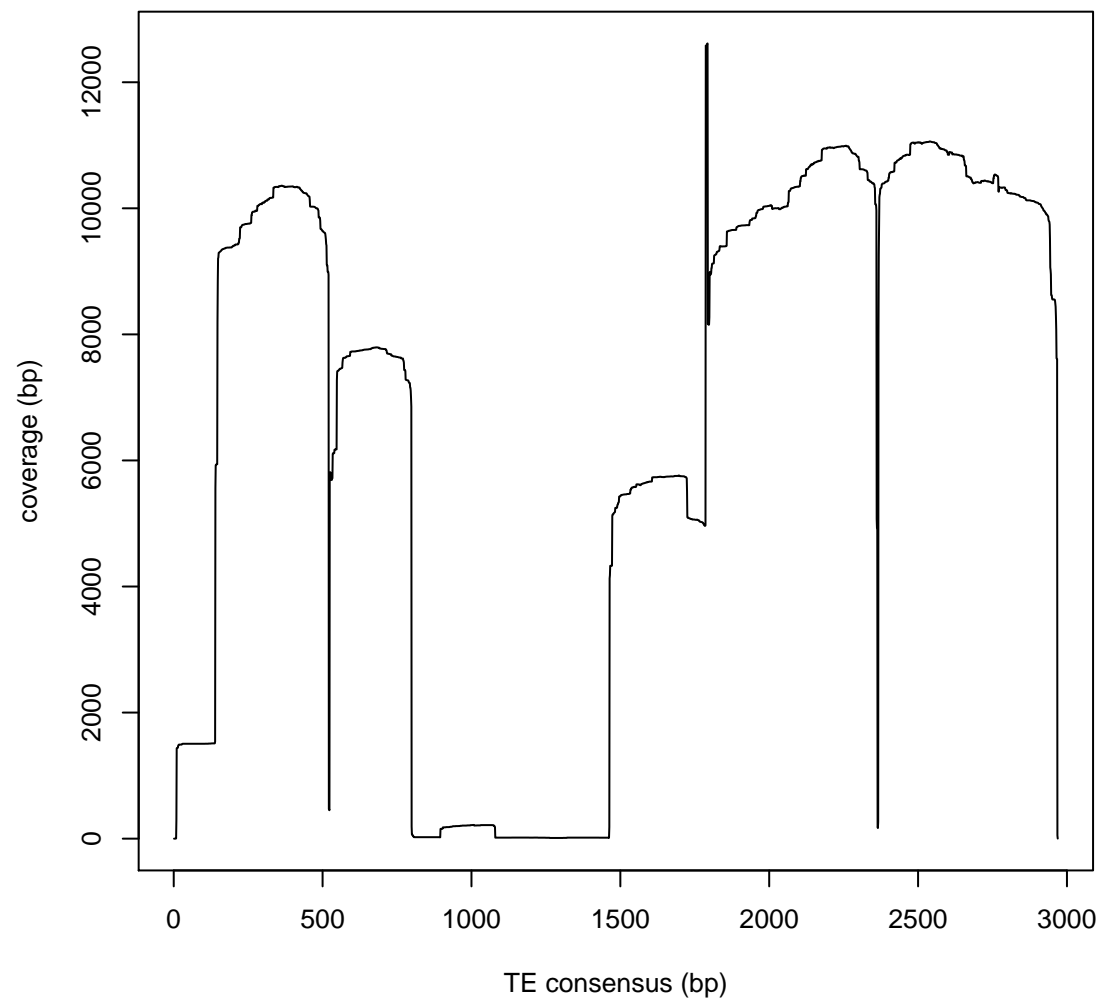


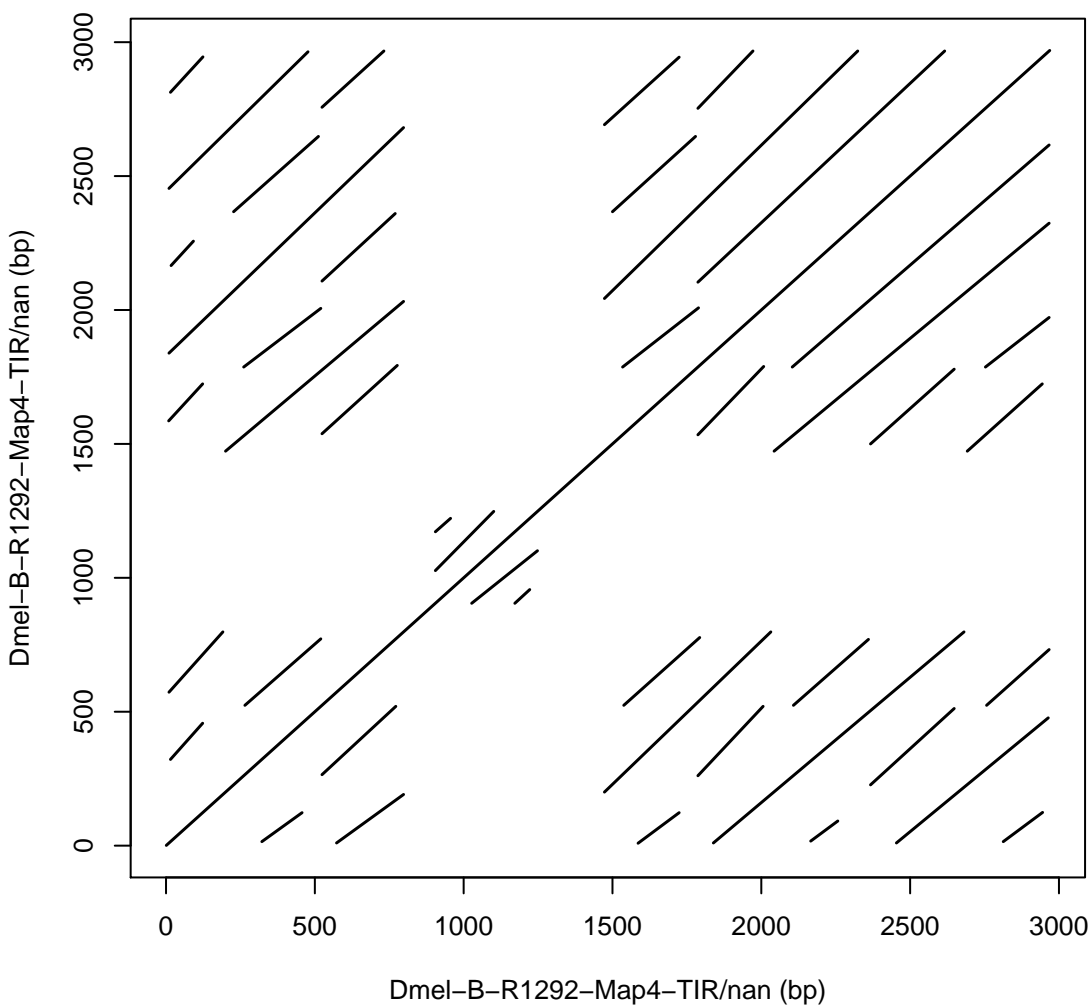
TE: Dmel-B-R1292-Map4-TIR/nan
consensus size: 2969bp; fragments: 46714; full length: 0 (>=2672.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

