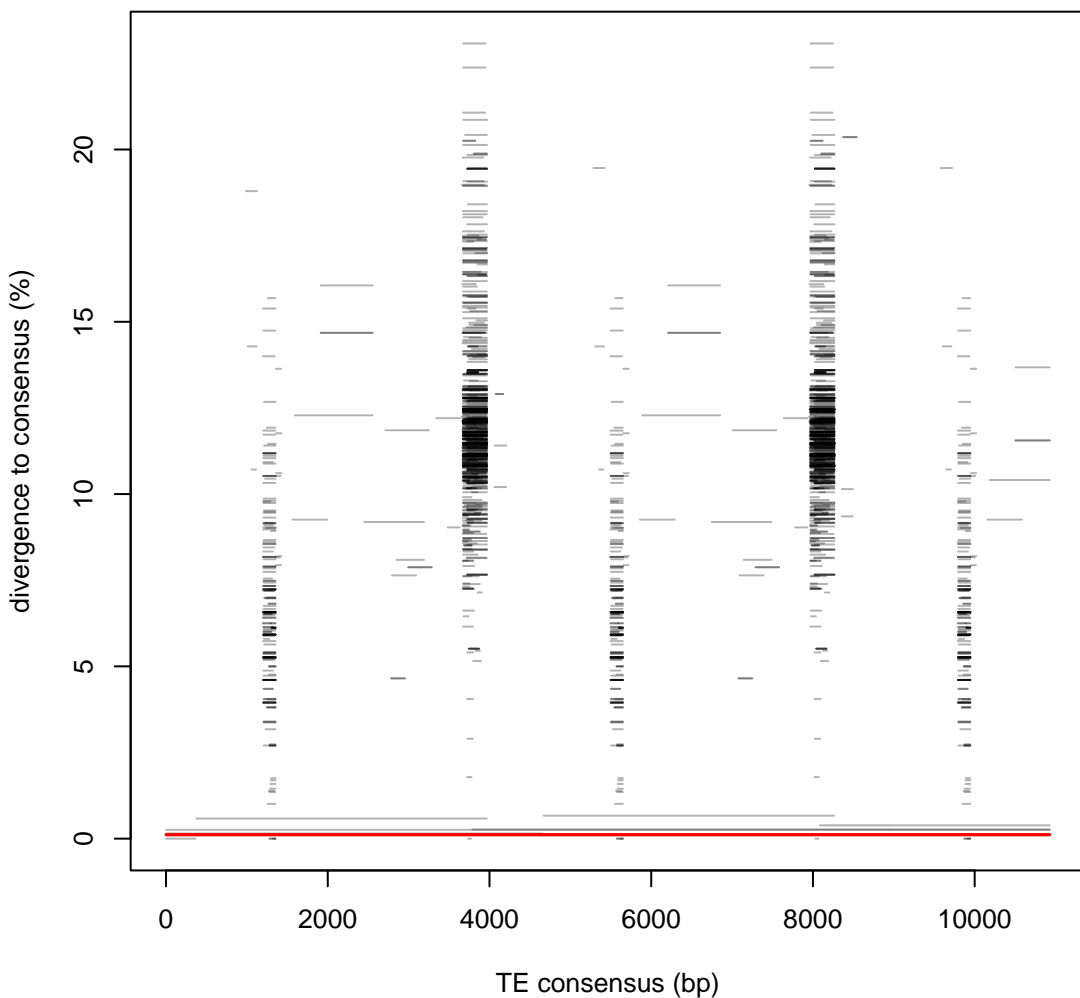
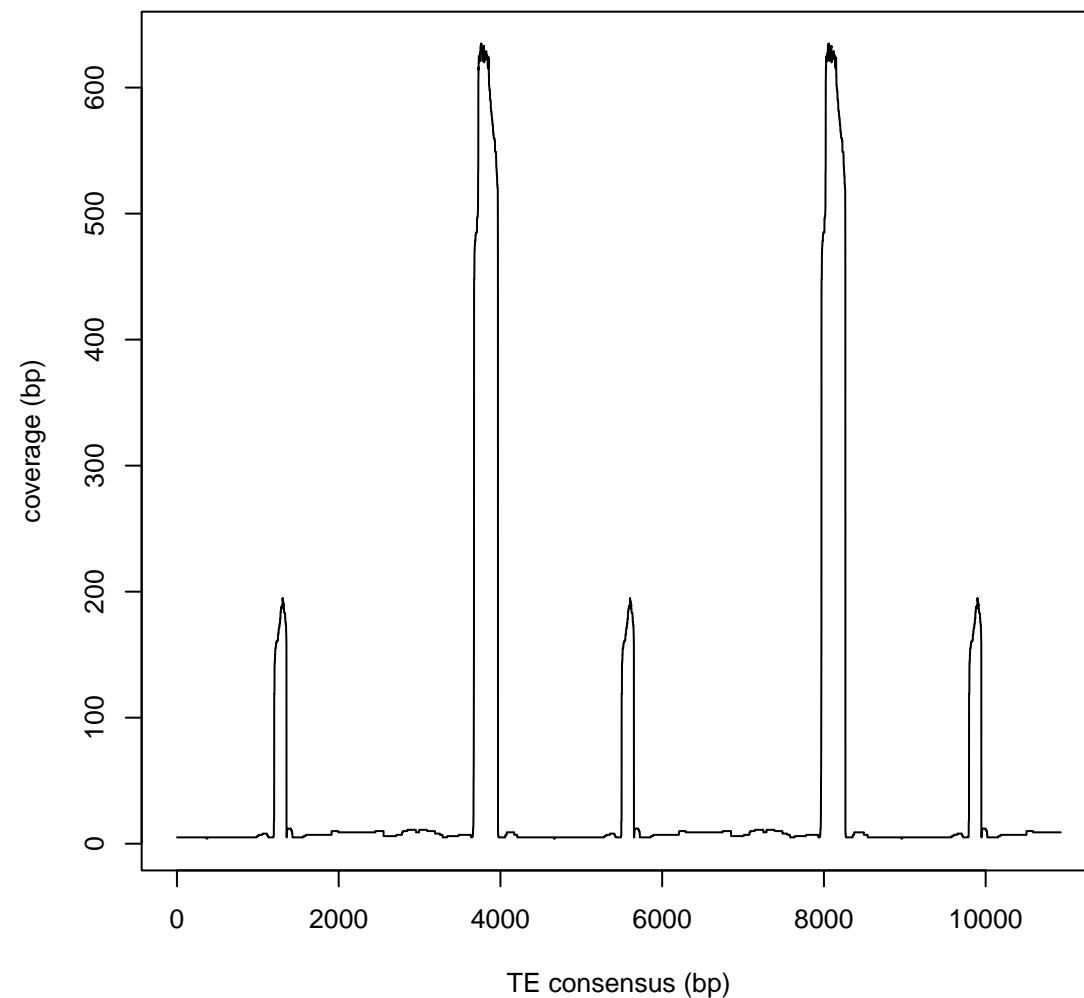


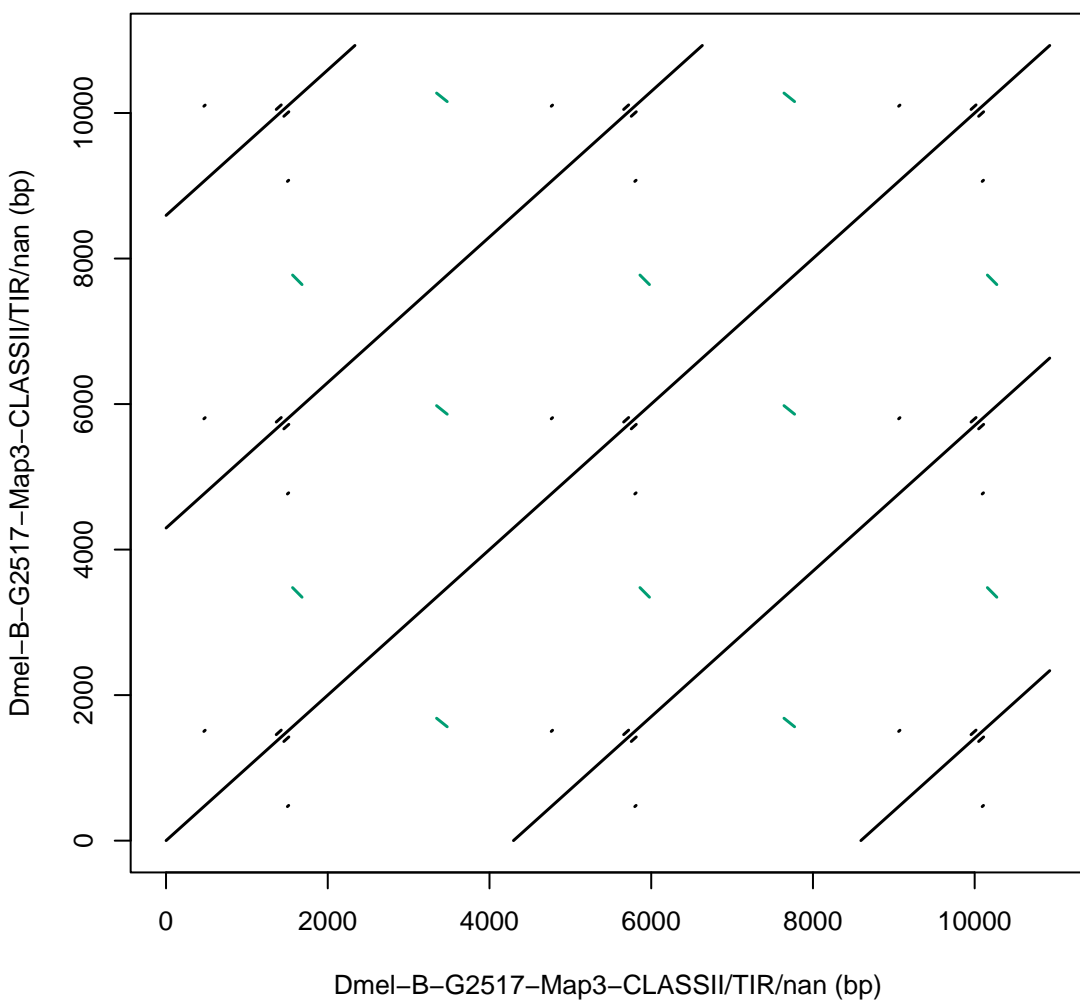
TE: Dmel-B-G2517-Map3-CLASSII/TIR/nan
consensus size: 10928bp; fragments: 2073; full length: 2 (>=9835.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

