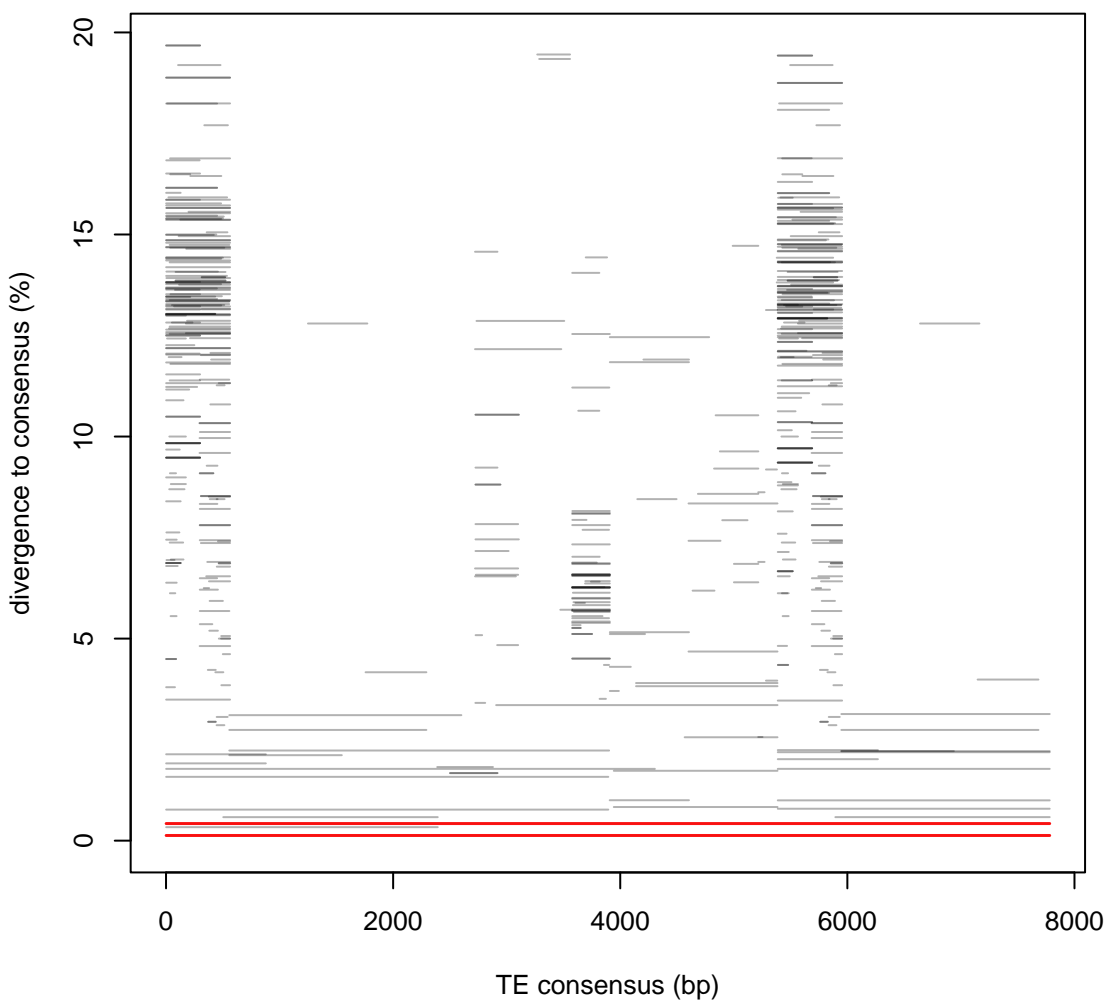
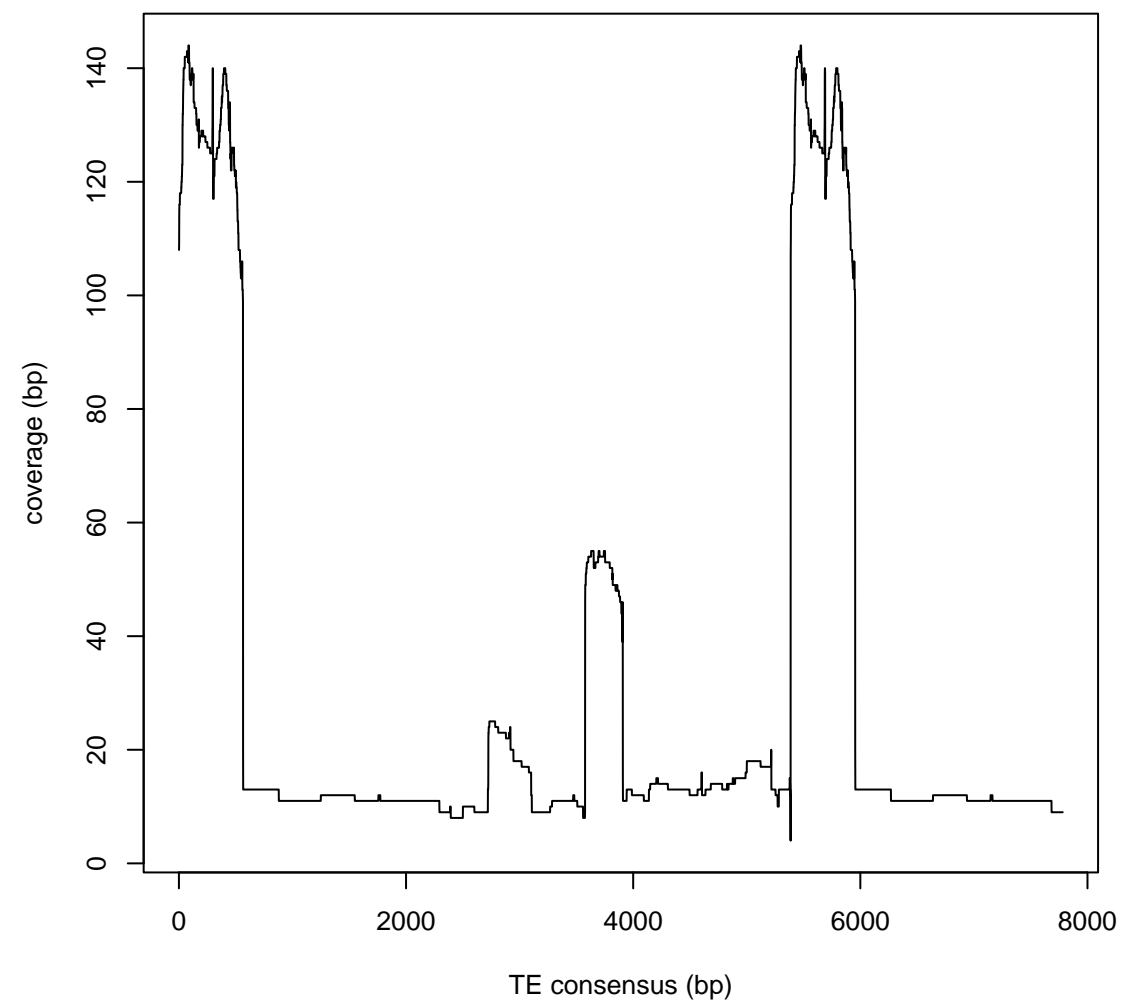


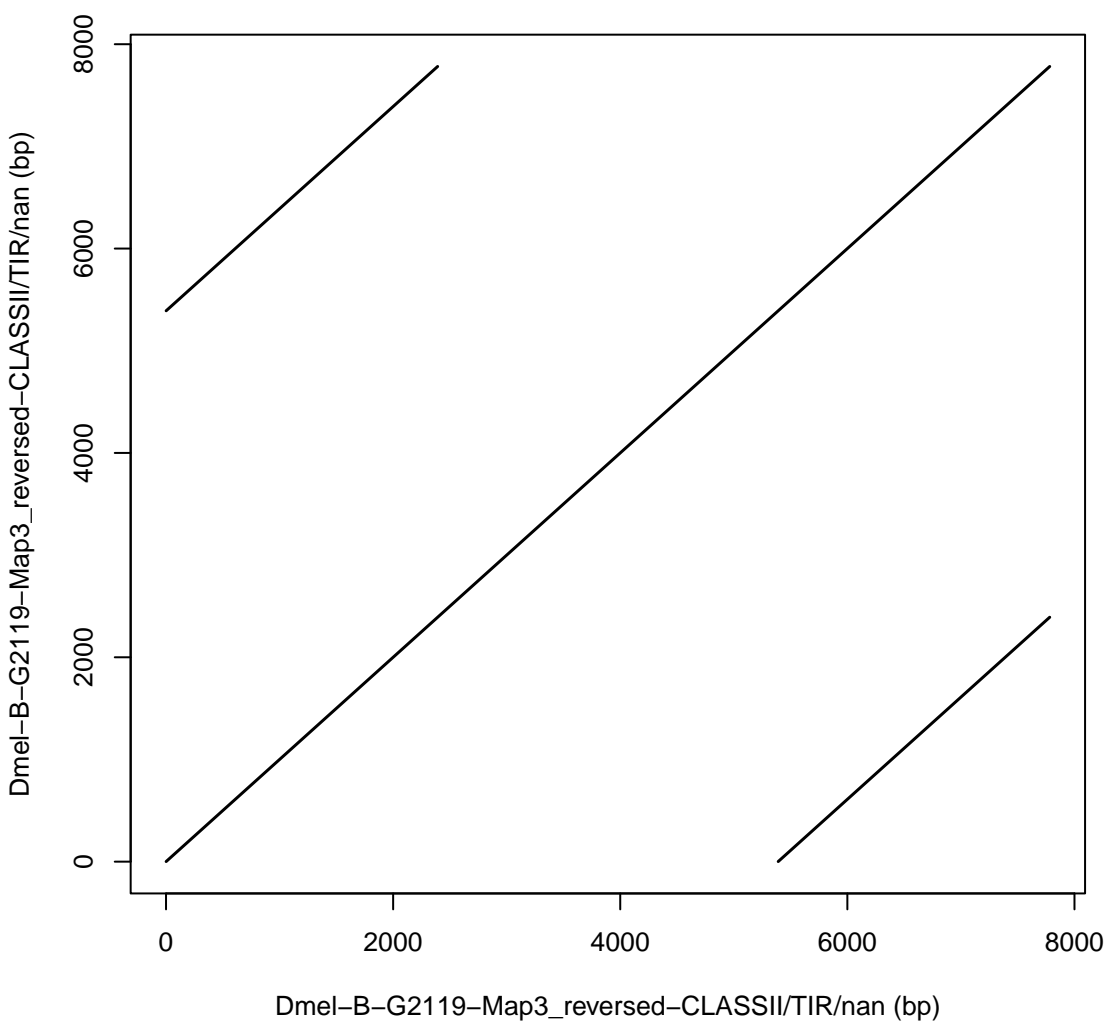
TE: Dmel-B-G2119-Map3_reversed-CLASSII/TIR/nan
consensus size: 7782bp; fragments: 553; full length: 2 (≥ 7003.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

