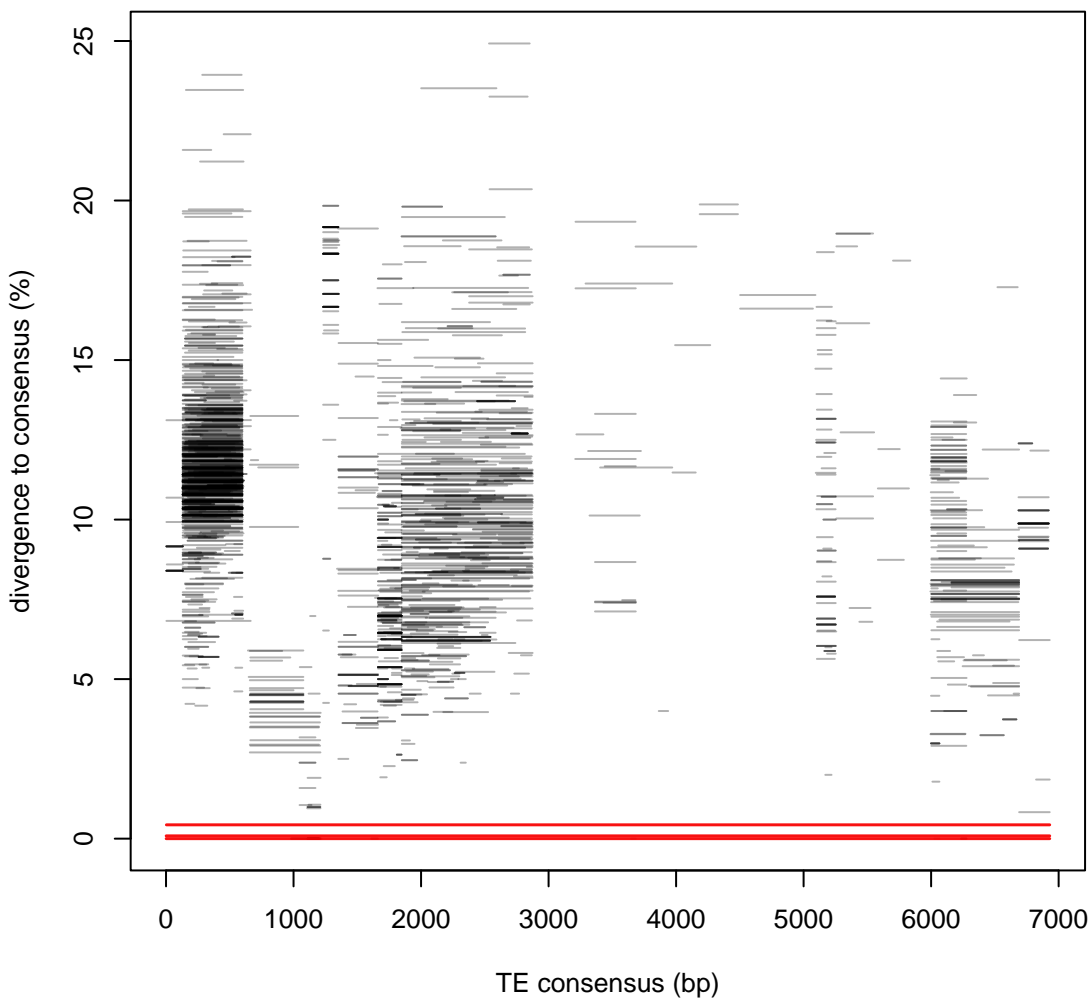
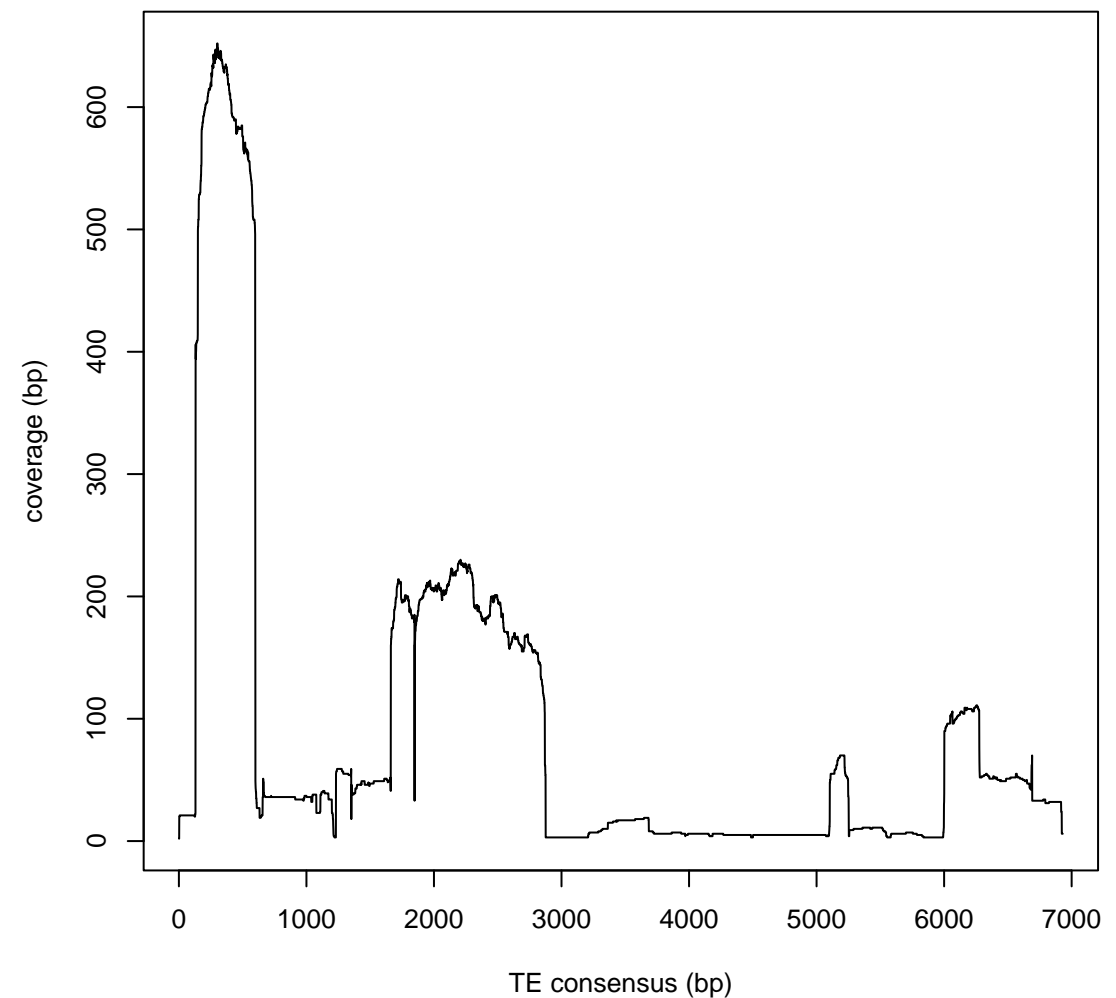


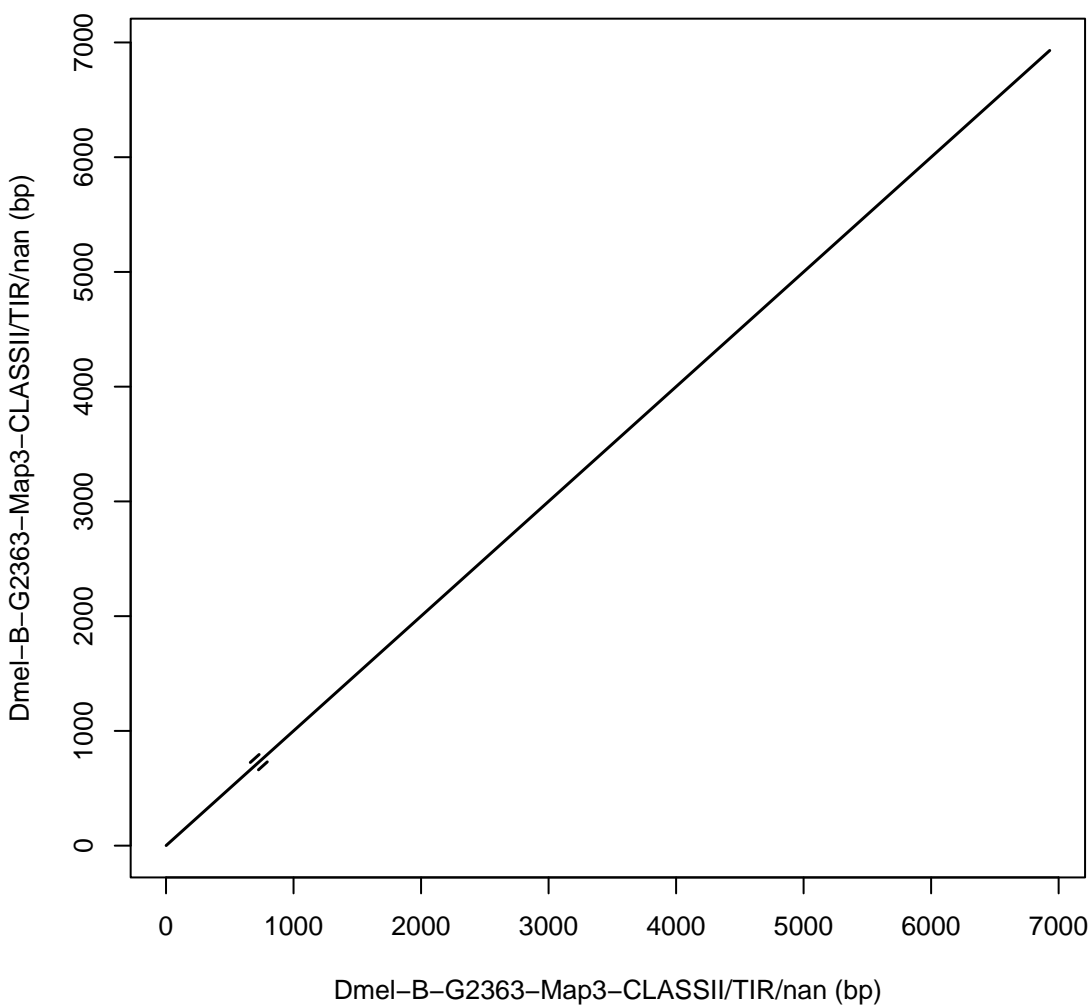
TE: Dmel-B-G2363-Map3-CLASSII/TIR/nan
consensus size: 6930bp; fragments: 1894; full length: 3 (>=6237bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

