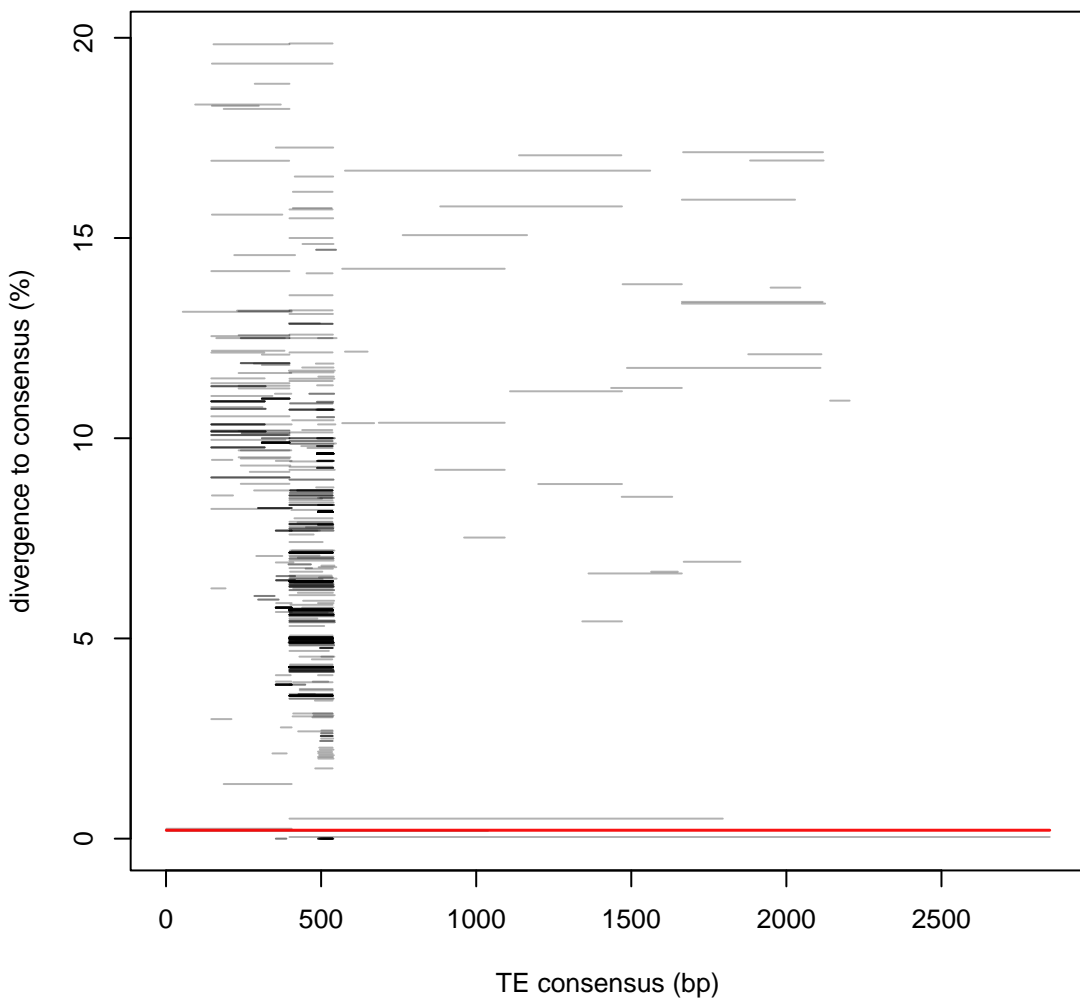
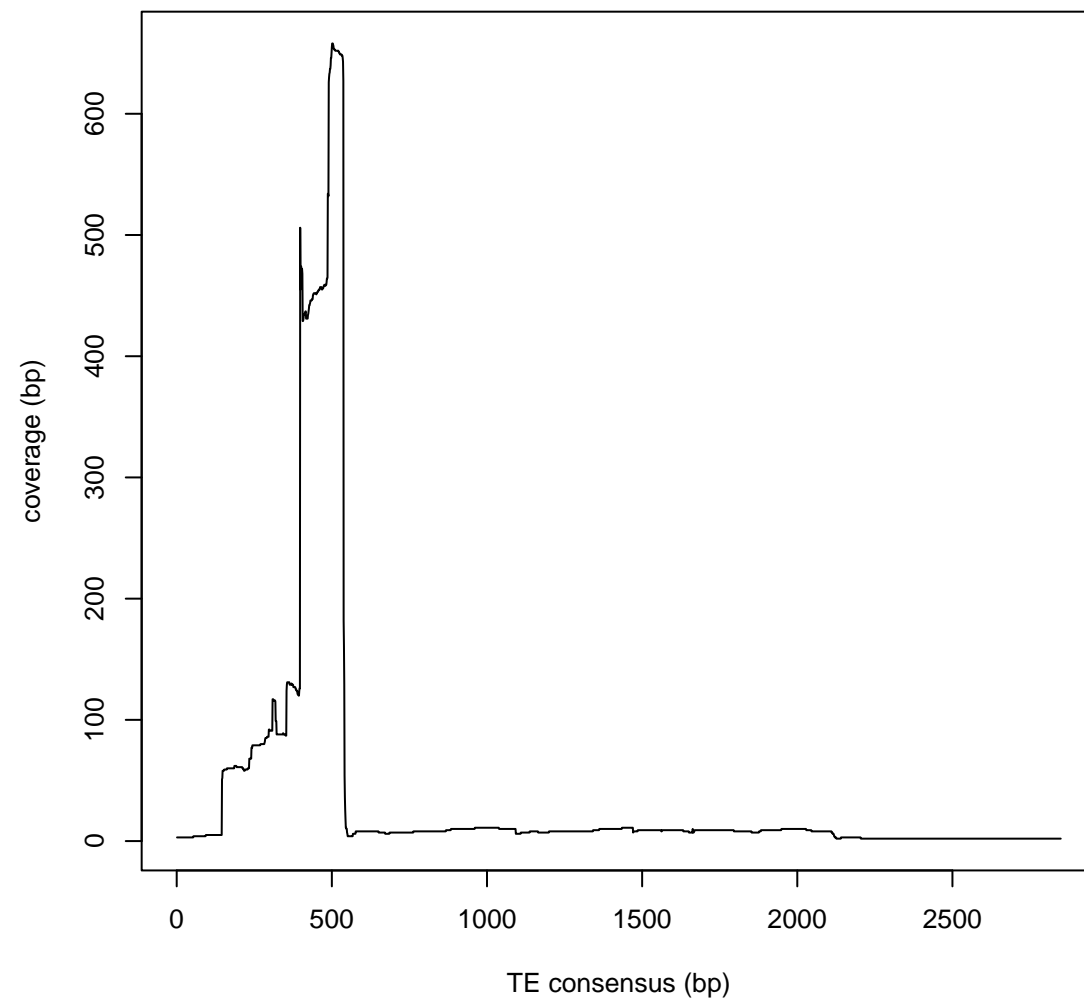


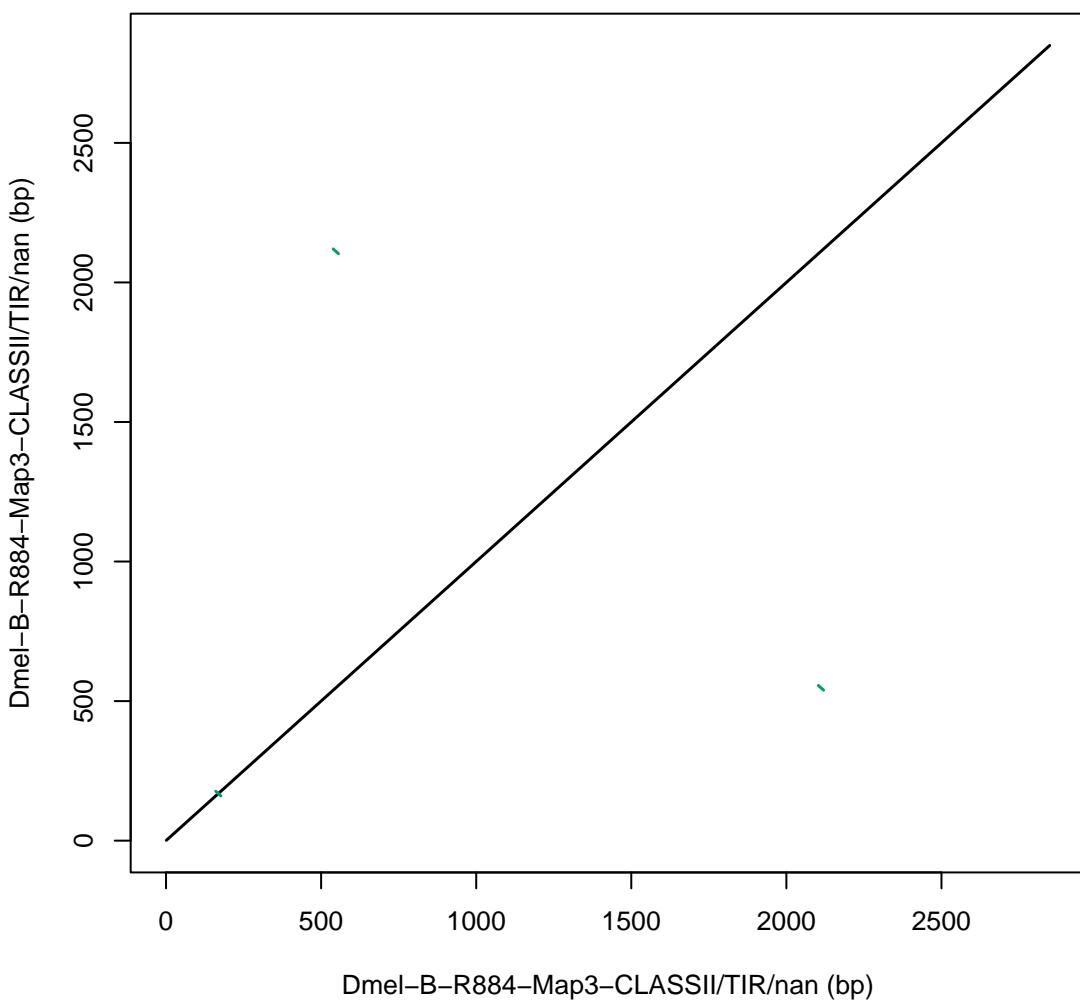
TE: Dmel-B-R884-Map3-CLASSII/TIR/nan
consensus size: 2849bp; fragments: 864; full length: 1 (>=2564.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

