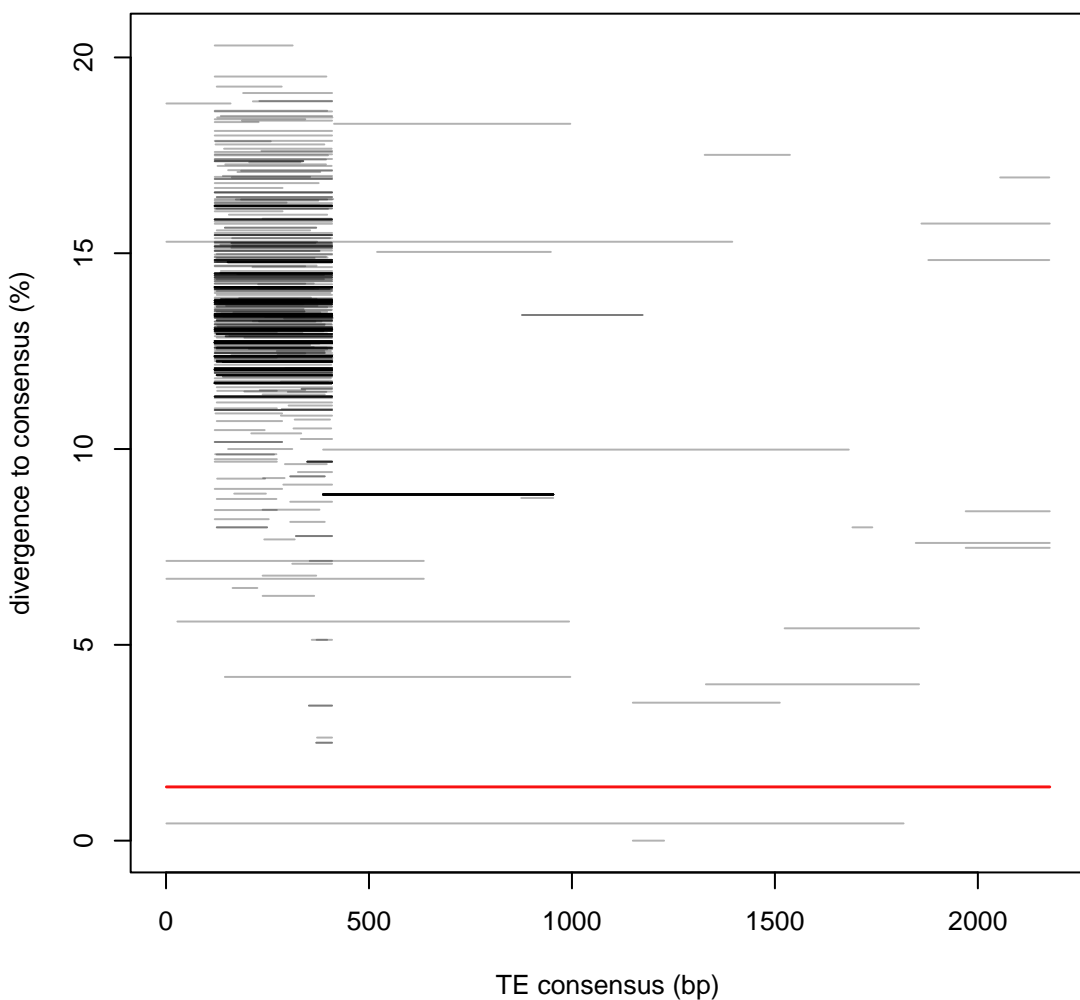
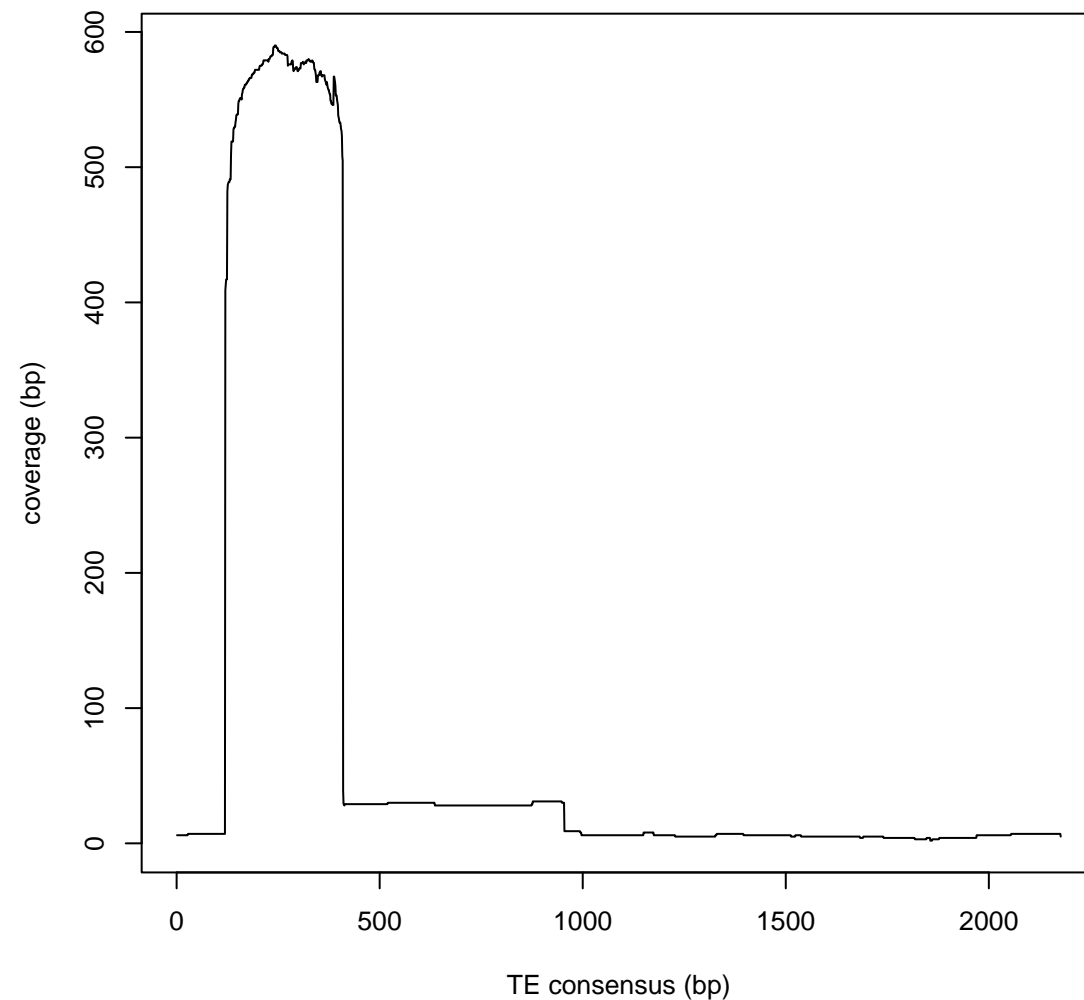


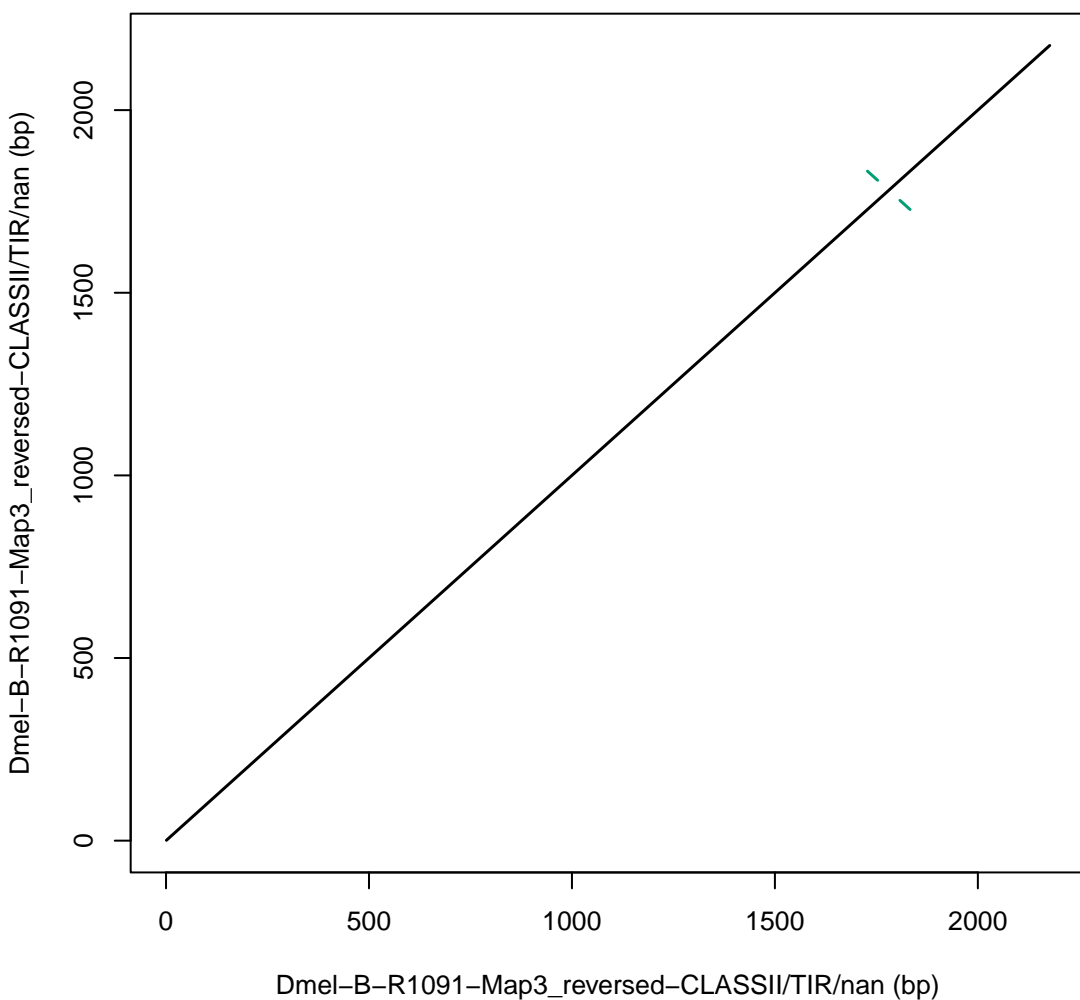
TE: Dmel-B-R1091-Map3\_reversed-CLASSII/TIR/nan  
consensus size: 2177bp; fragments: 669; full length: 1 (>=1959.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

