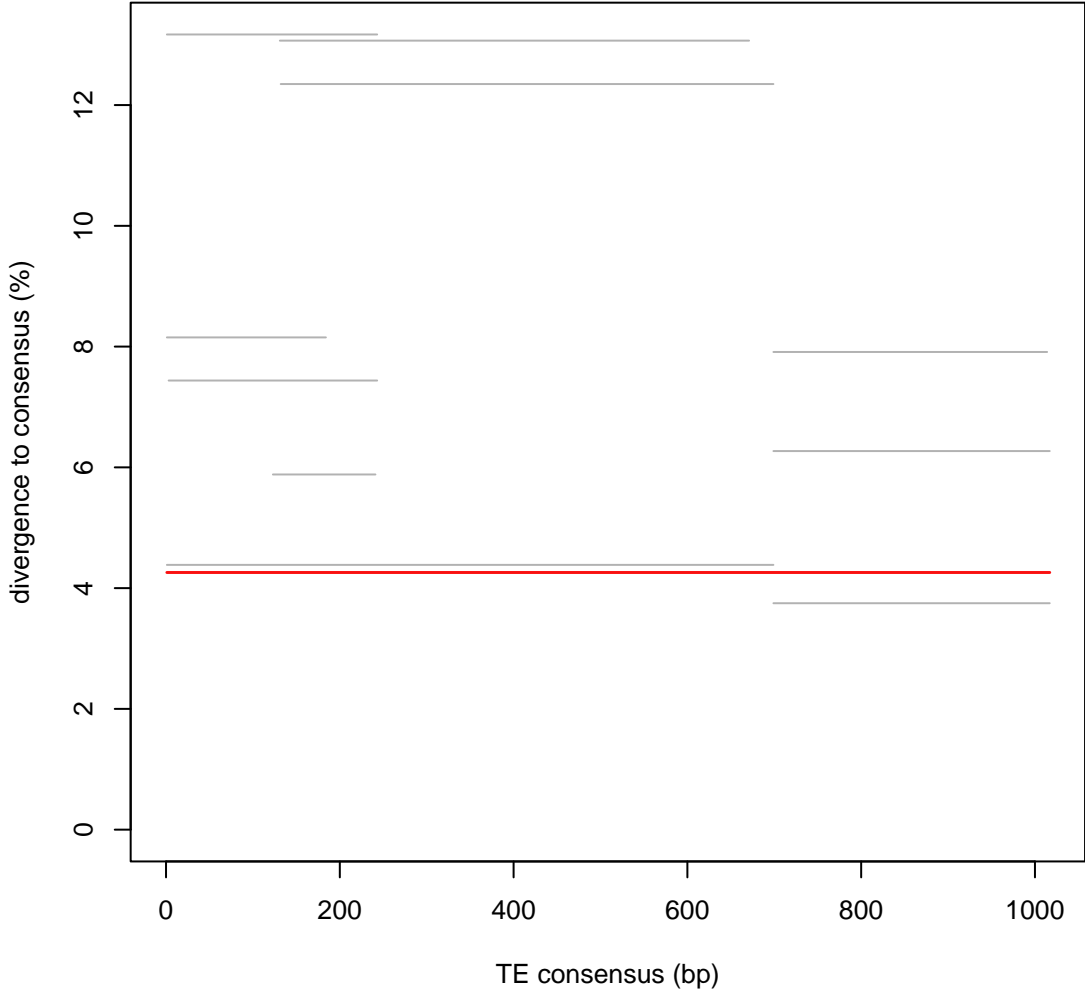
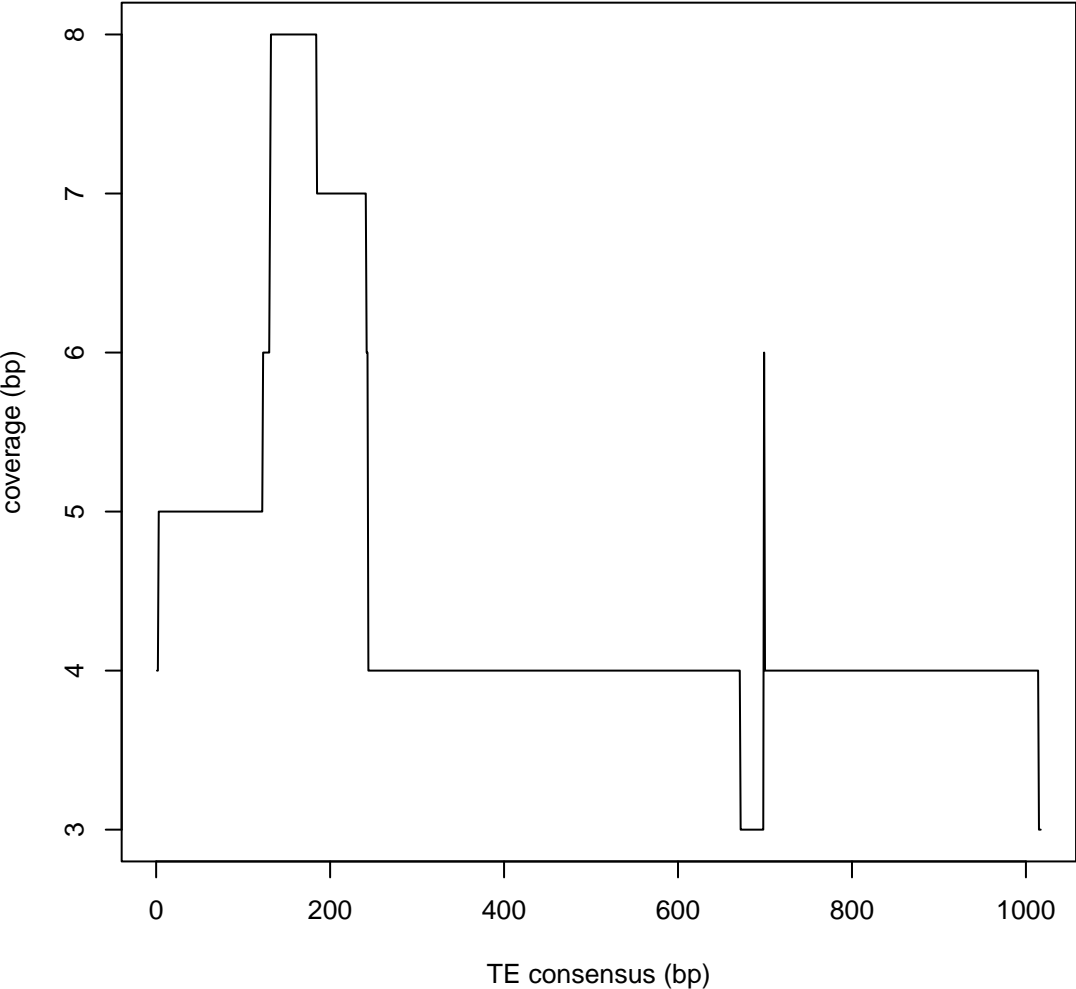


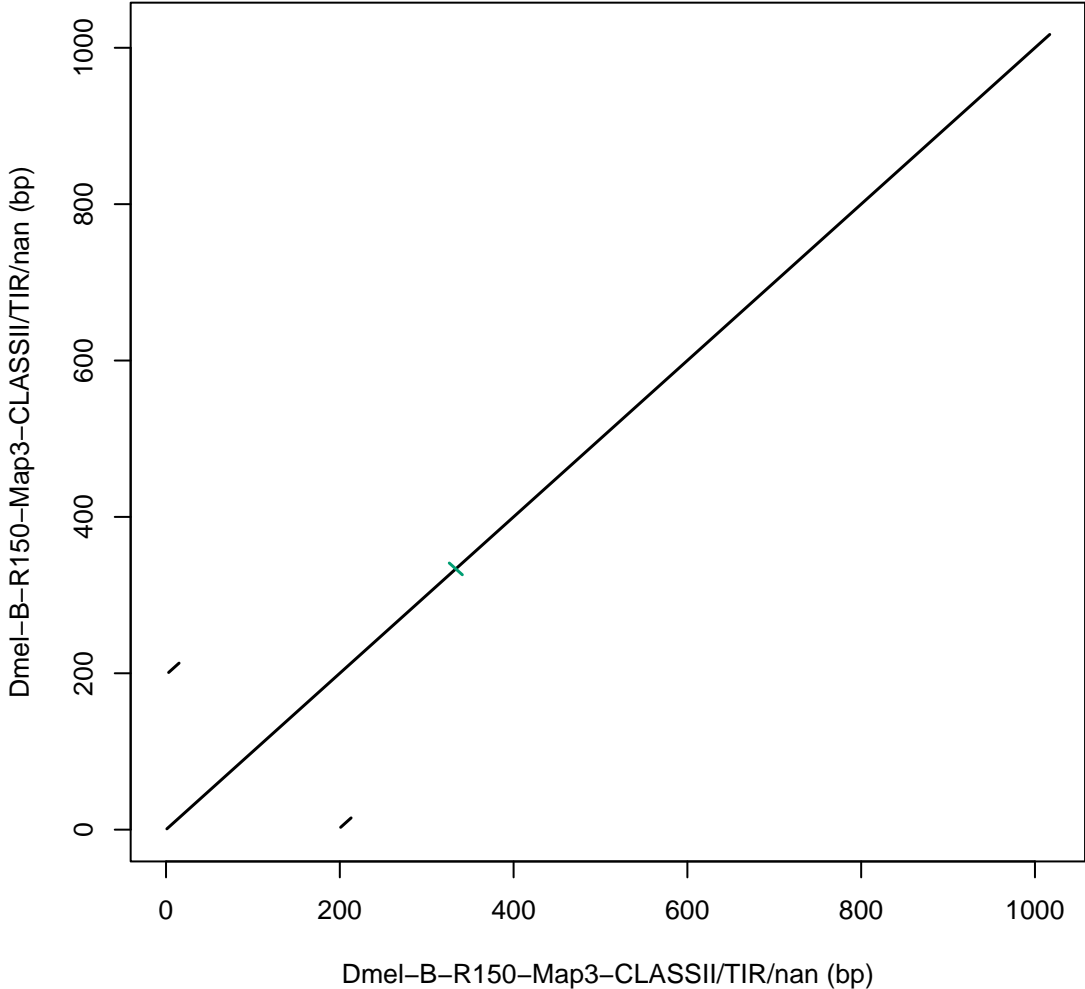
TE: Dmel-B-R150-Map3-CLASSII/TIR/nan  
consensus size: 1017bp; fragments: 11; full length: 1 (>=915.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

