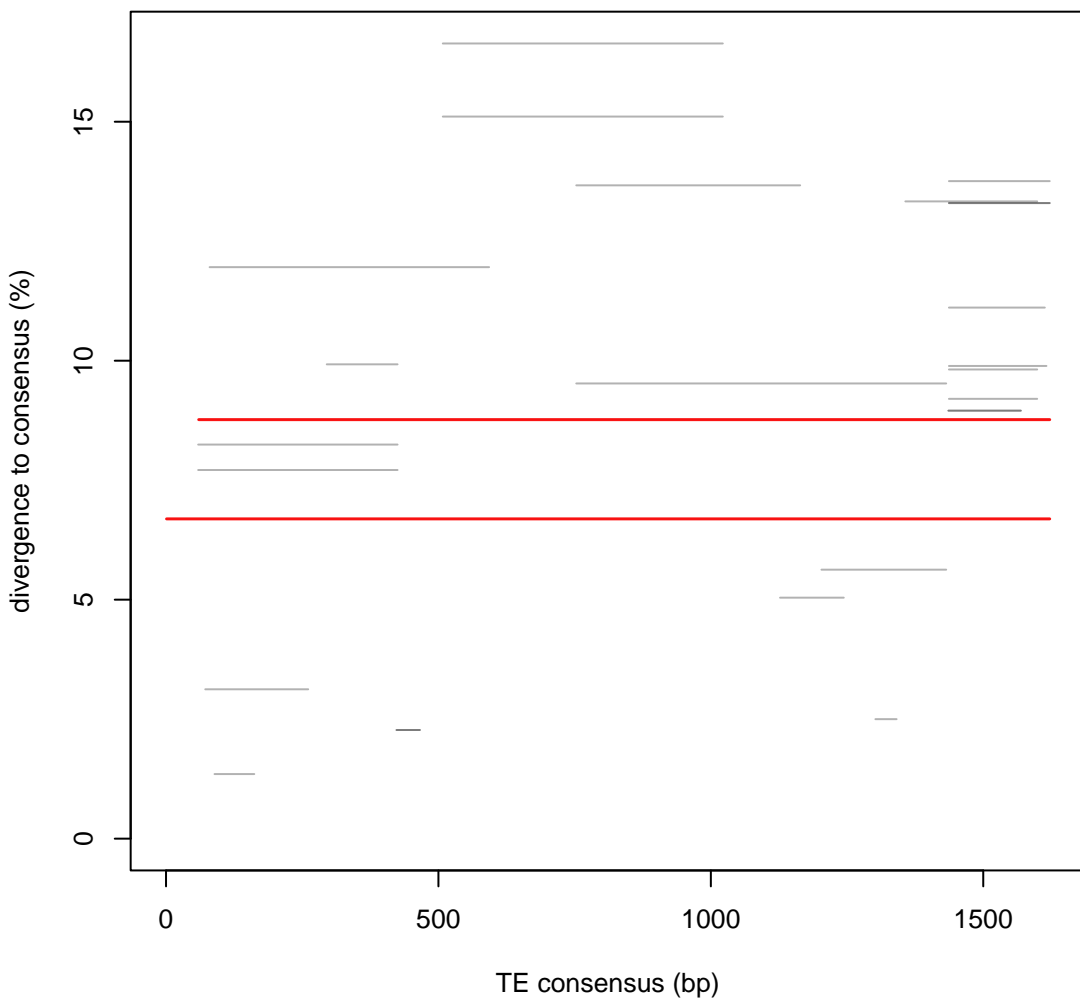
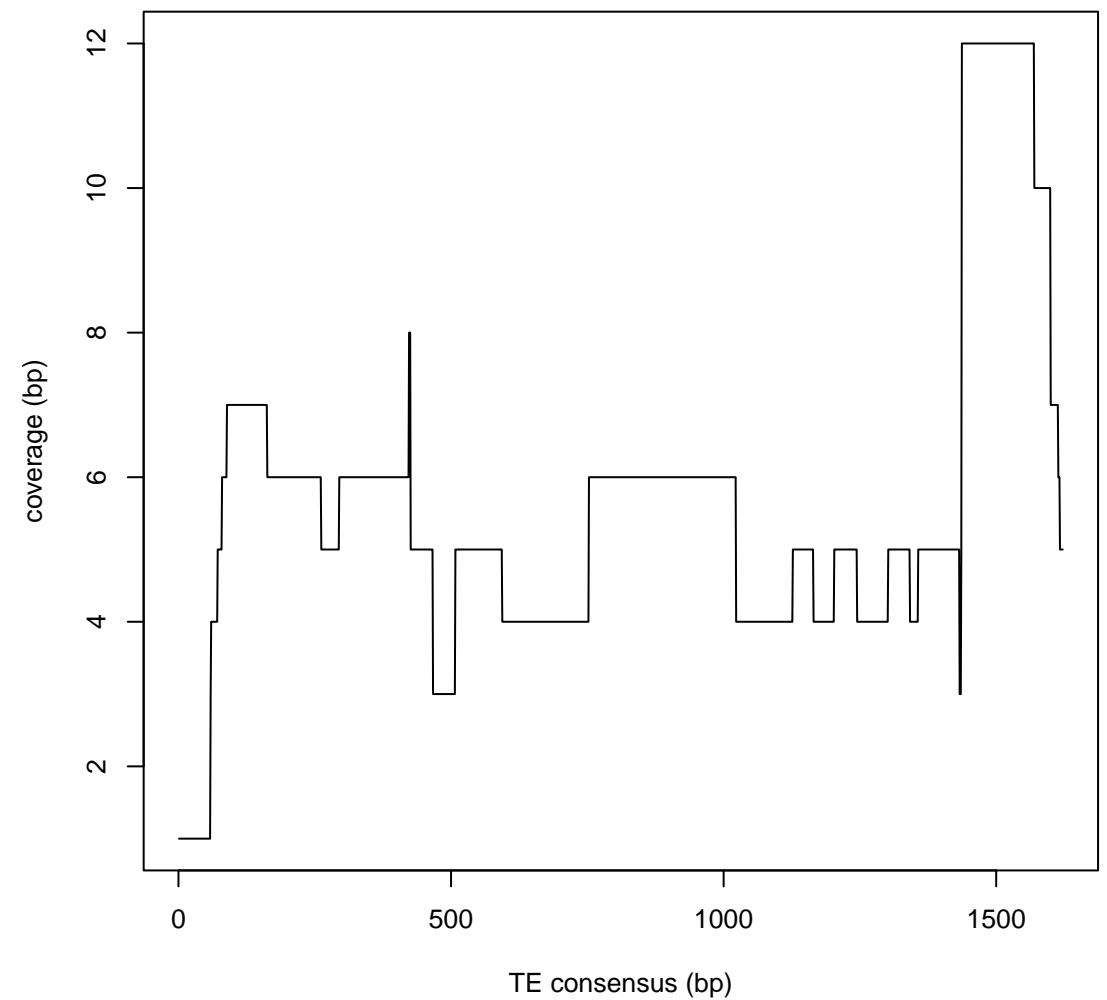


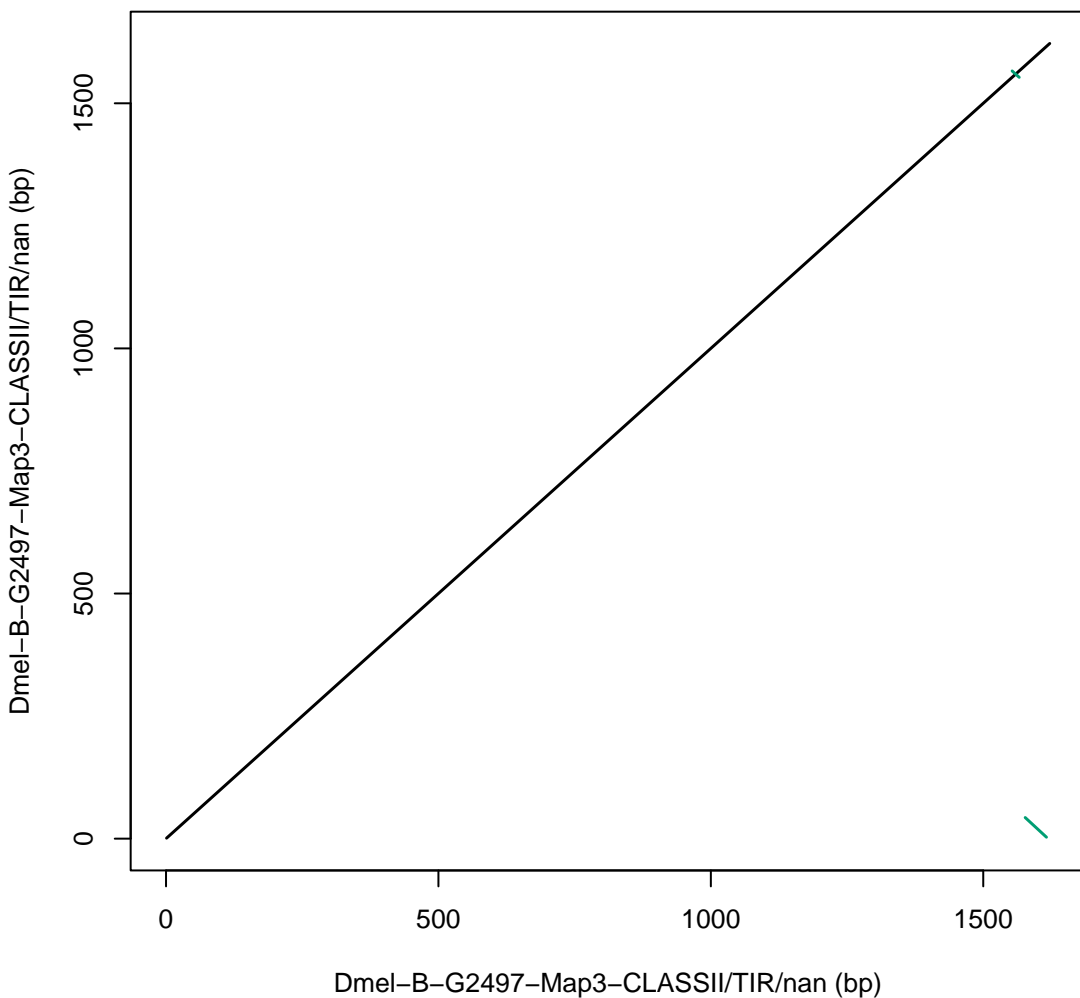
TE: Dmel-B-G2497-Map3-CLASSII/TIR/nan  
consensus size: 1622bp; fragments: 27; full length: 2 ( $\geq 1459.8$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

