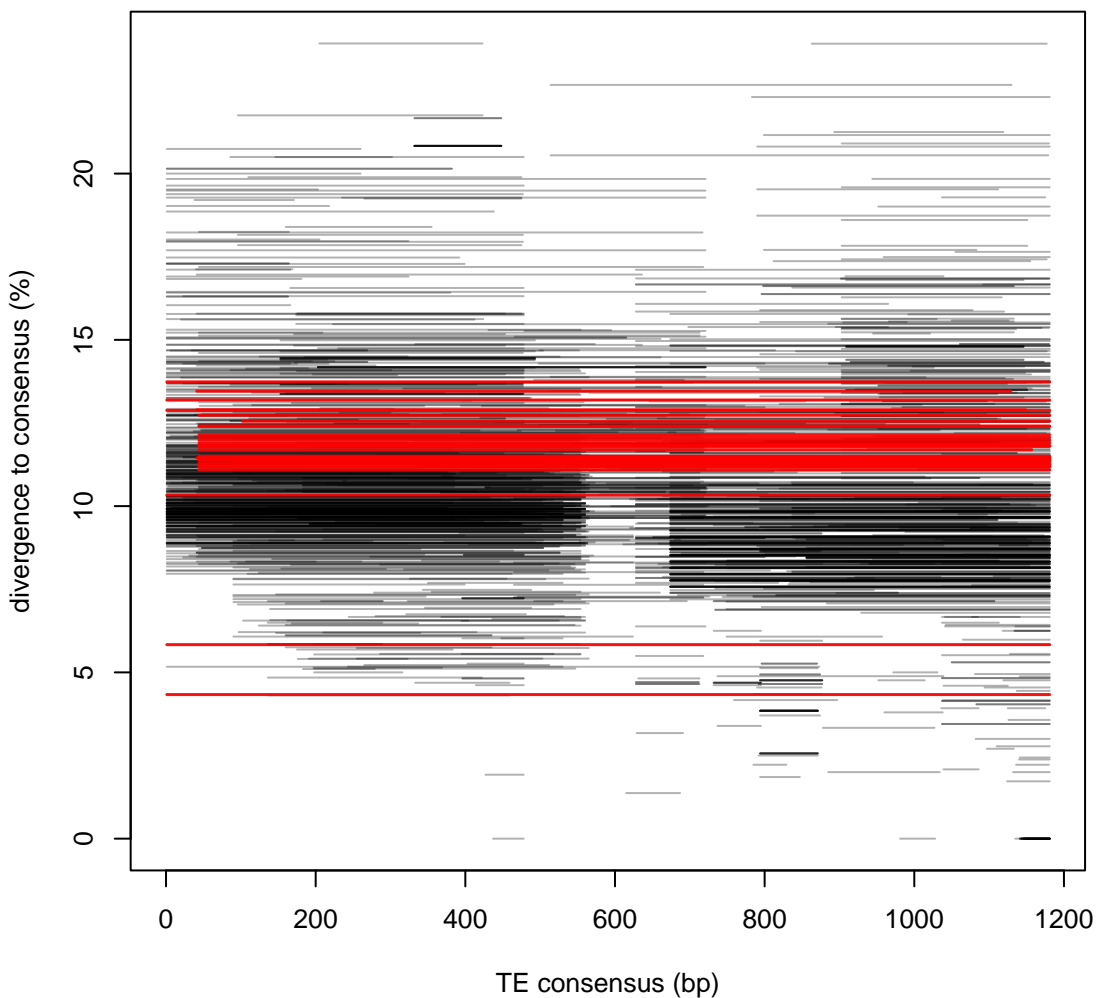
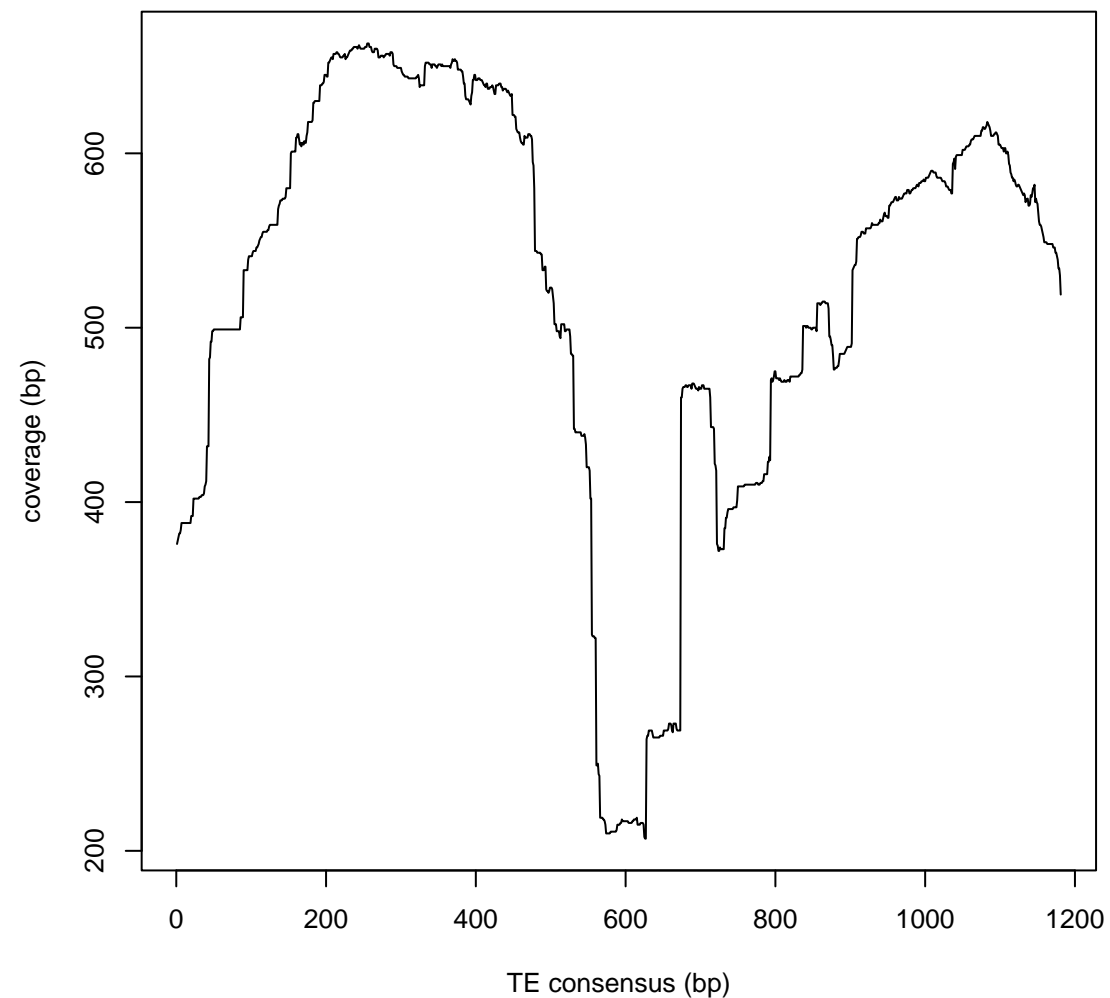


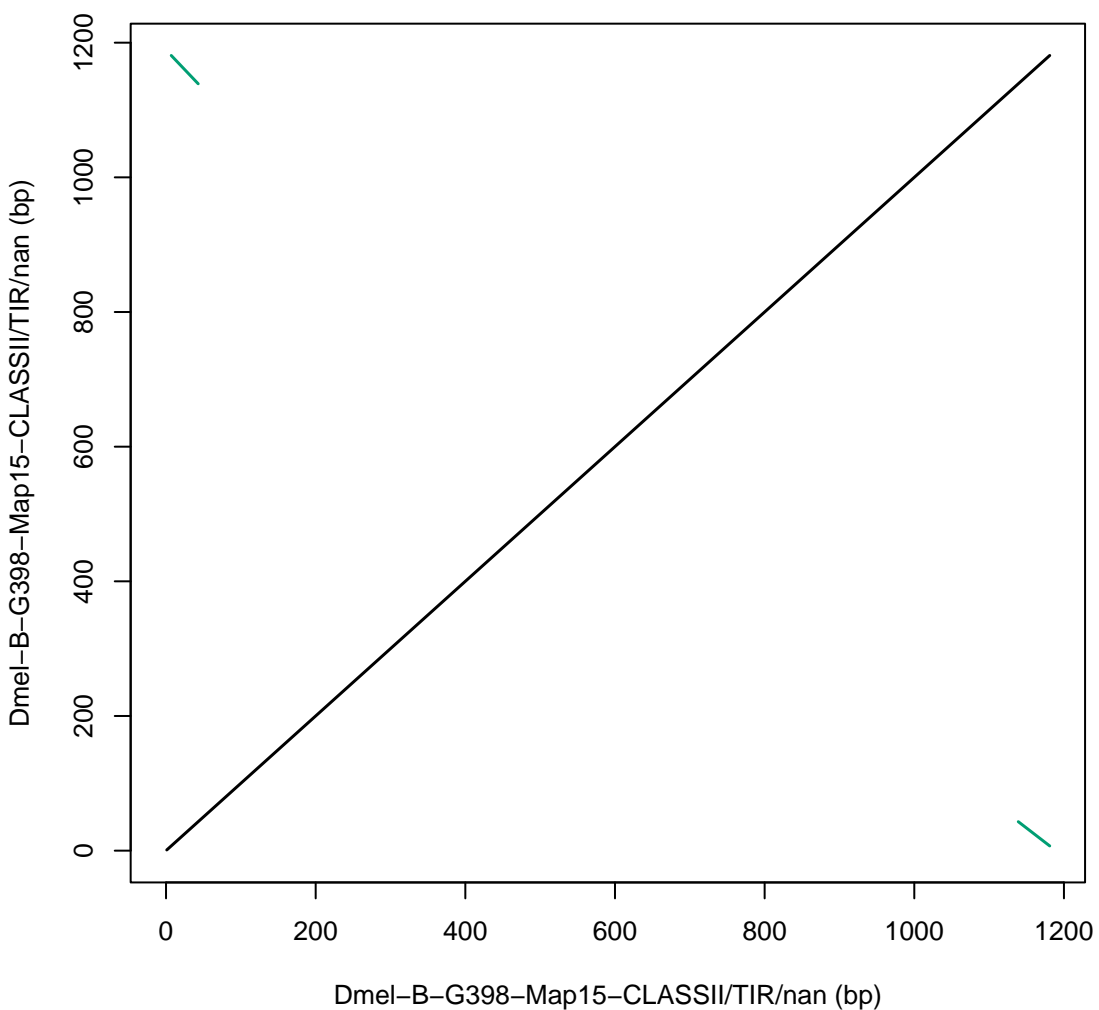
TE: Dmel-B-G398-Map15-CLASSII/TIR/nan  
consensus size: 1181bp; fragments: 1504; full length: 36 ( $\geq 1062.9$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

