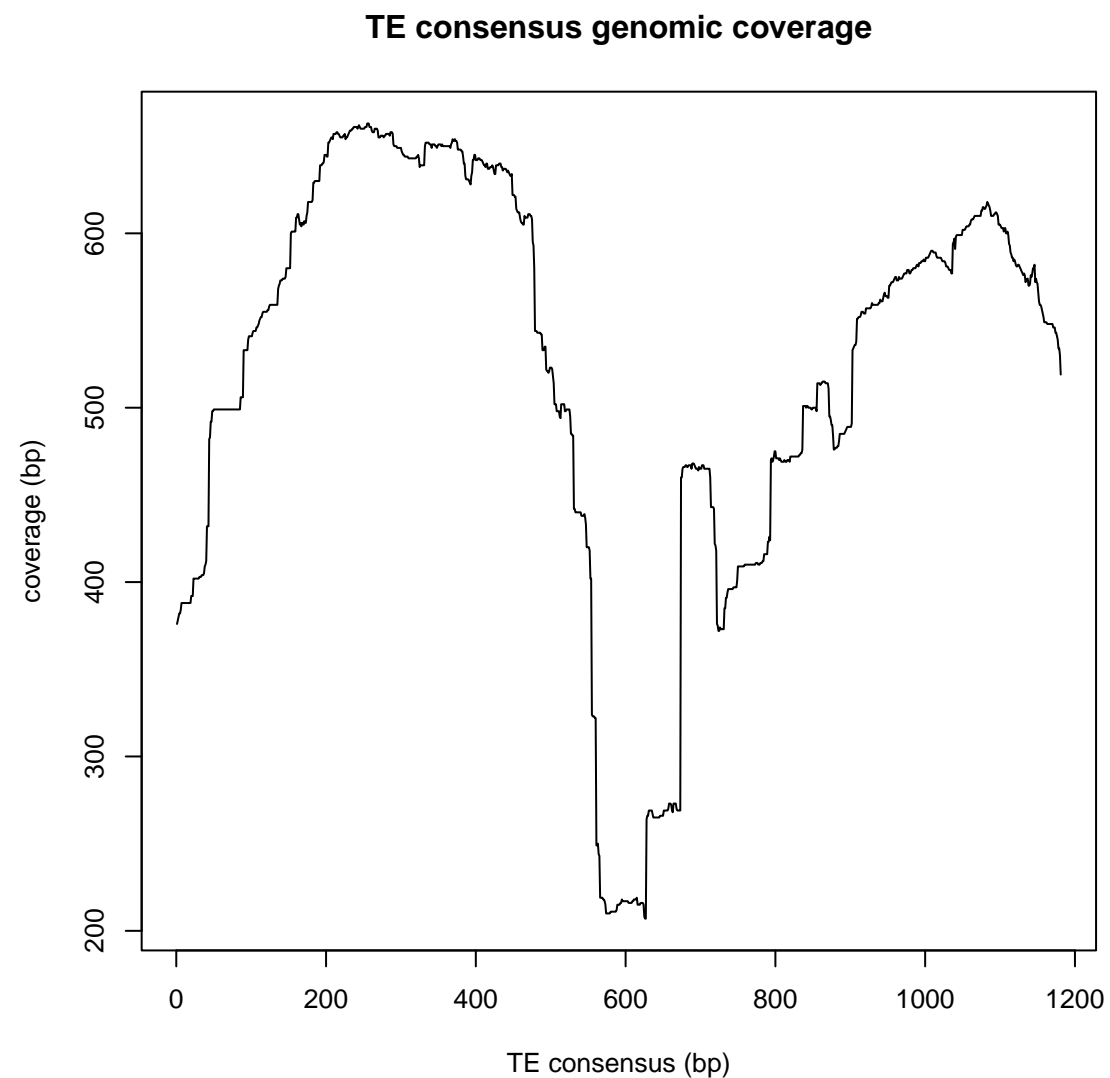
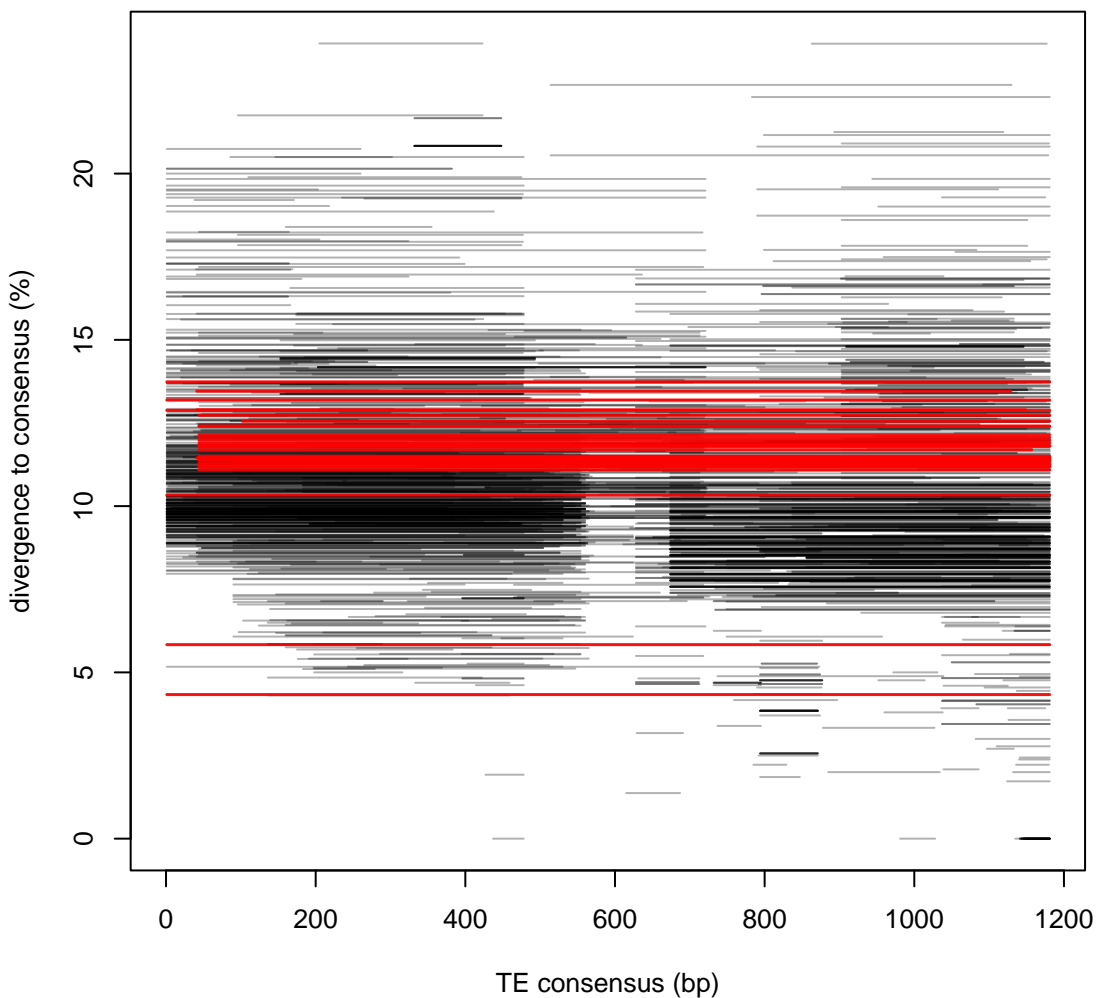
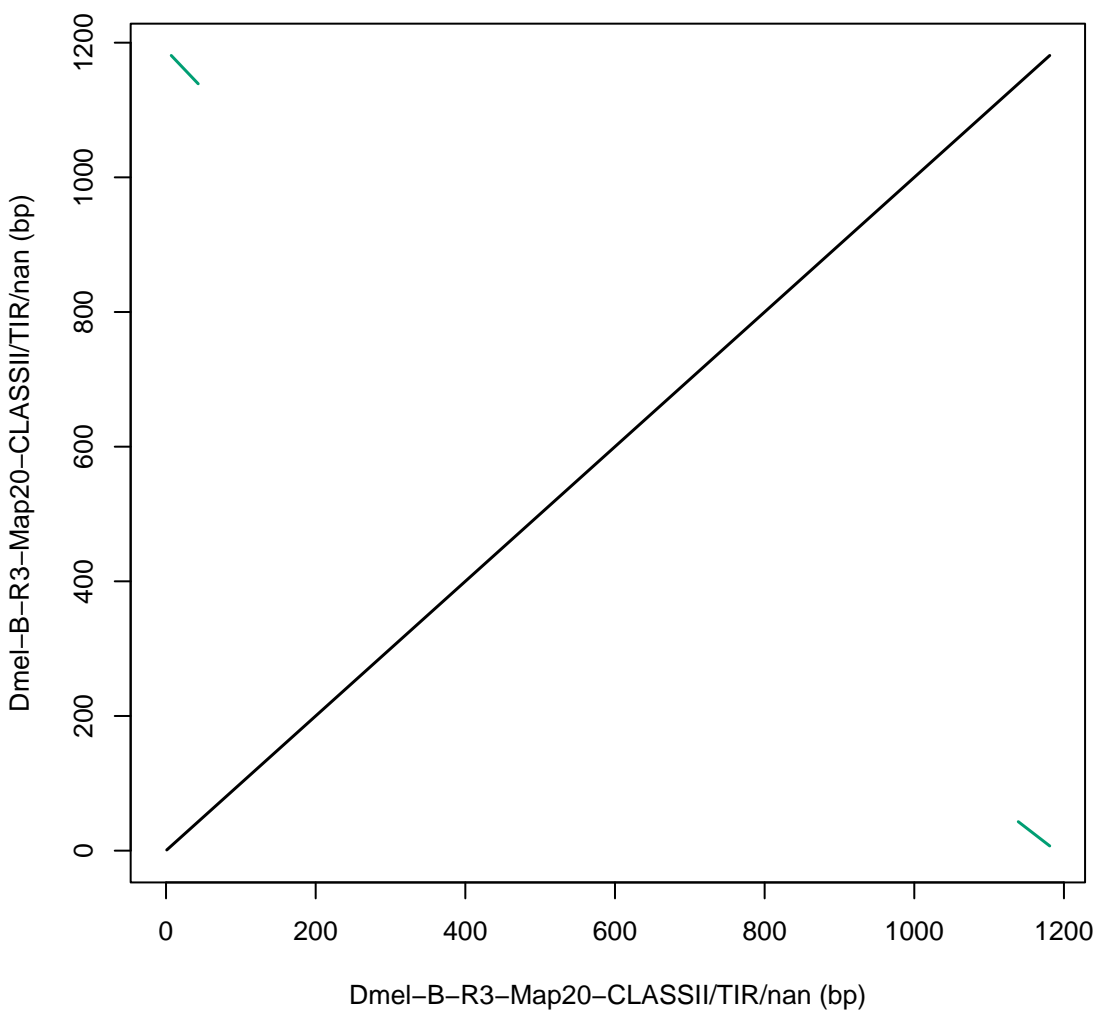


TE: Dmel-B-R3-Map20-CLASSII/TIR/nan  
consensus size: 1181bp; fragments: 1504; full length: 36 (>=1062.9bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

