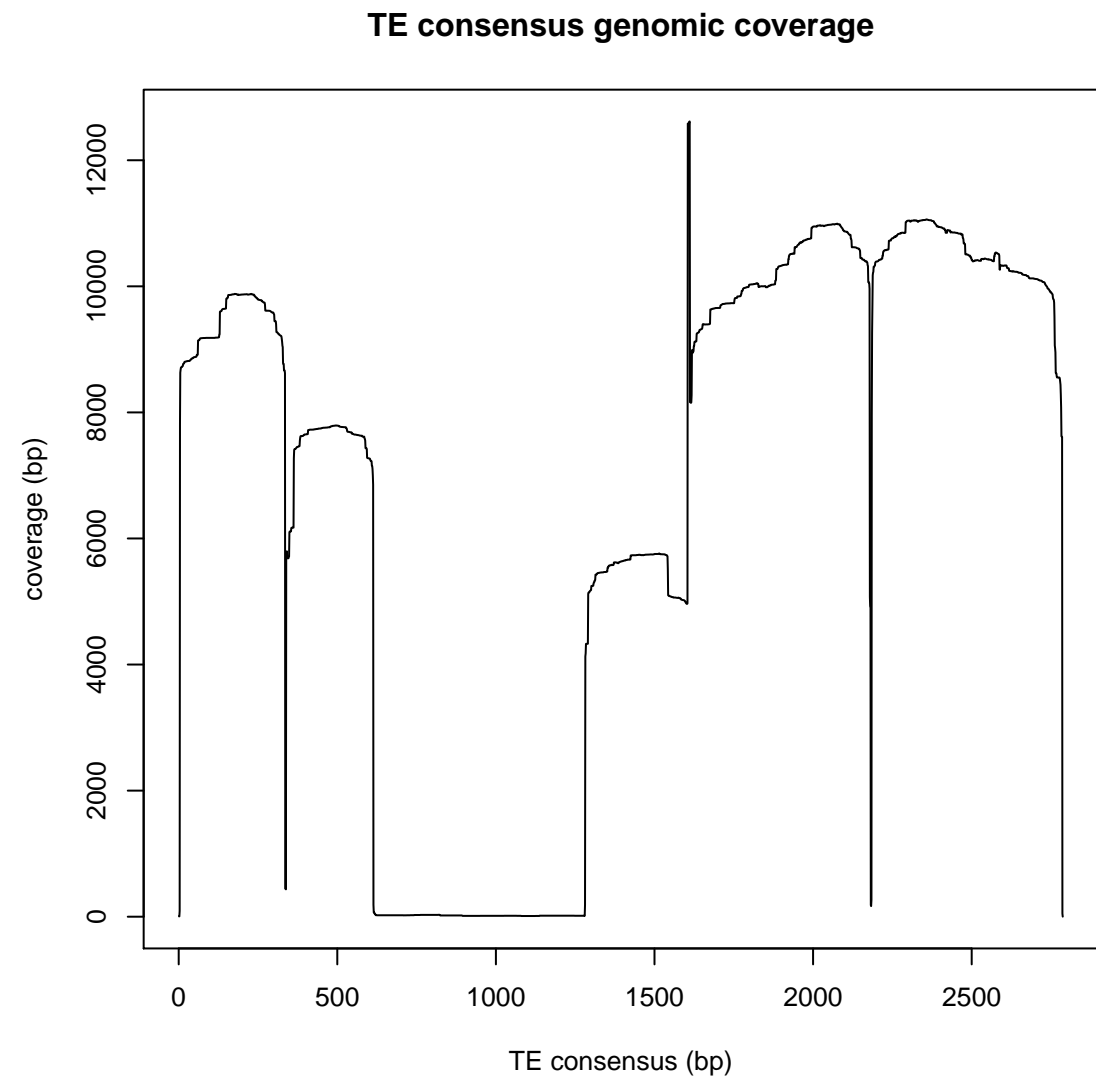
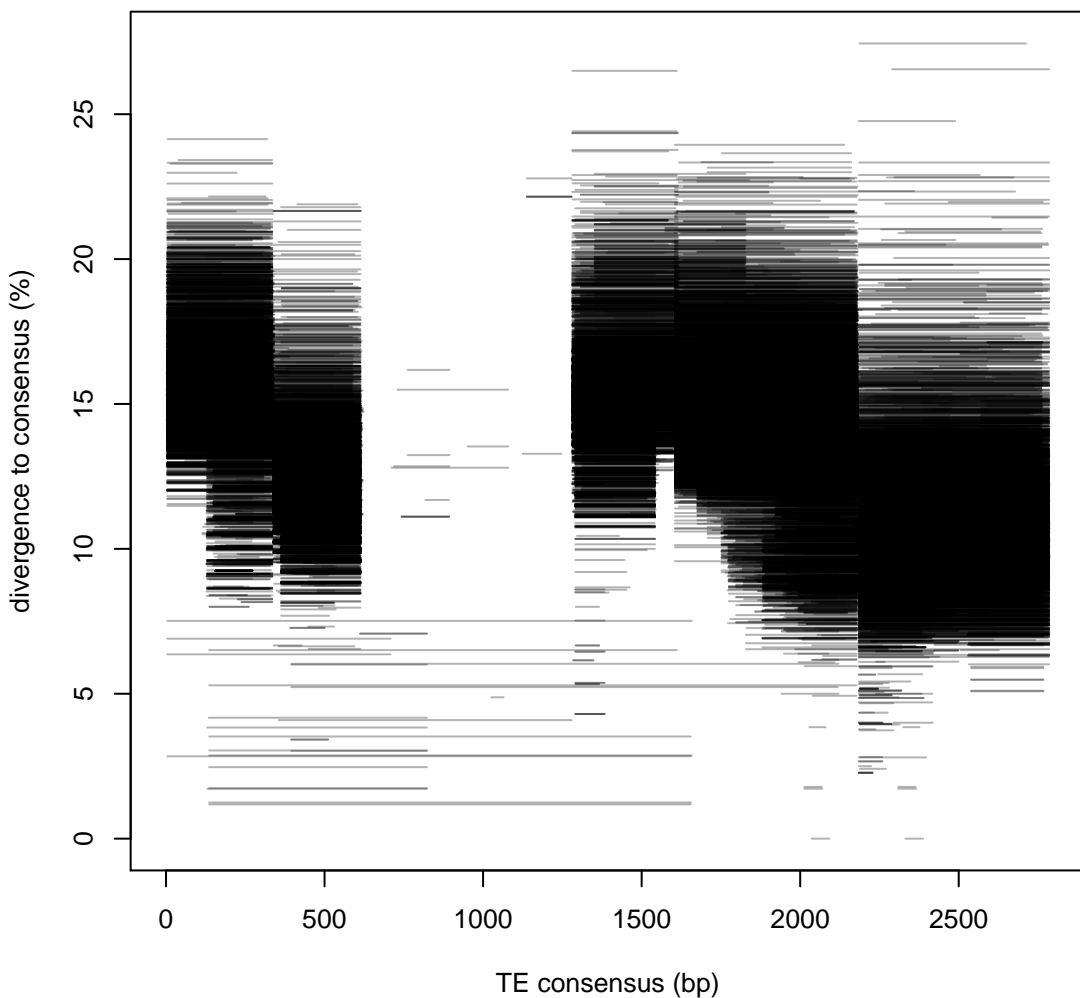
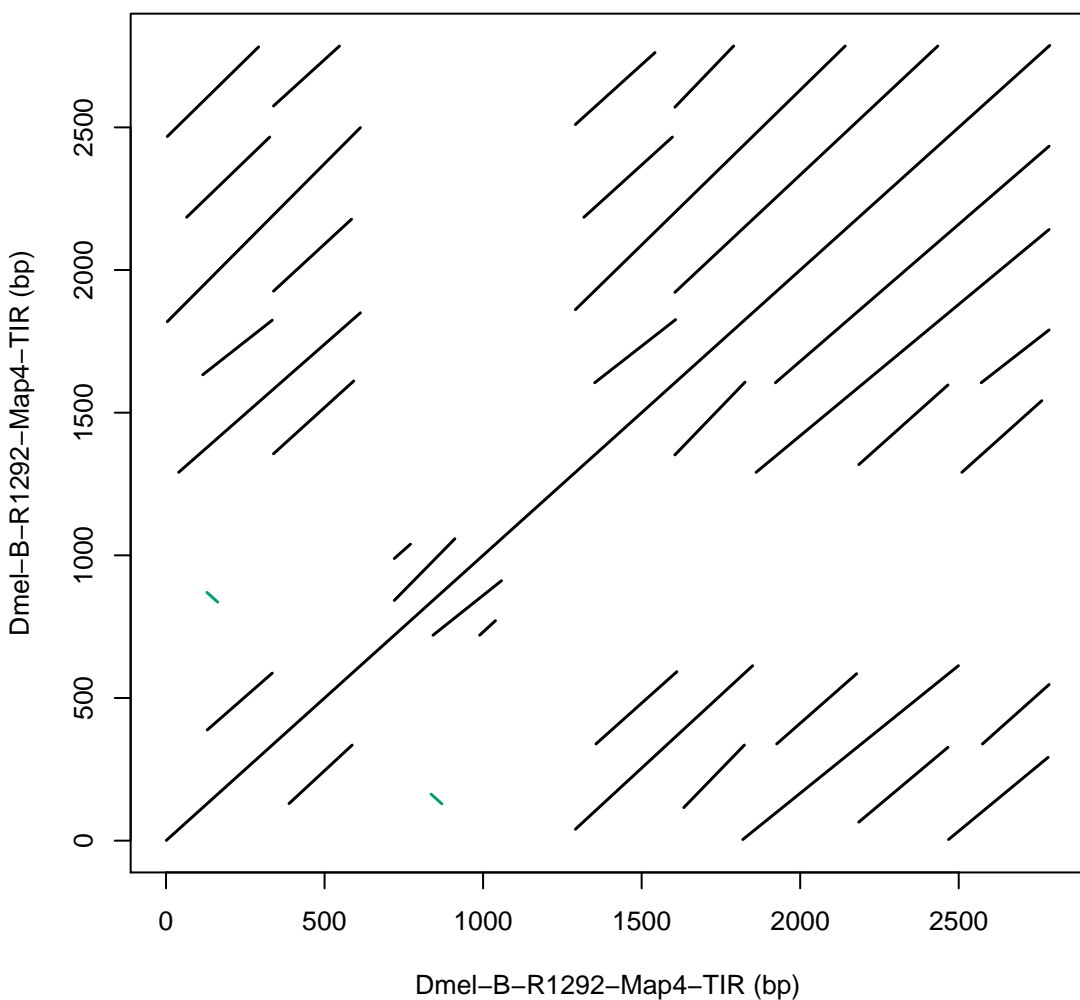


TE: Dmel-B-R1292-Map4-TIR  
consensus size: 2787bp; fragments: 45982; full length: 0 (>=2508.3bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

