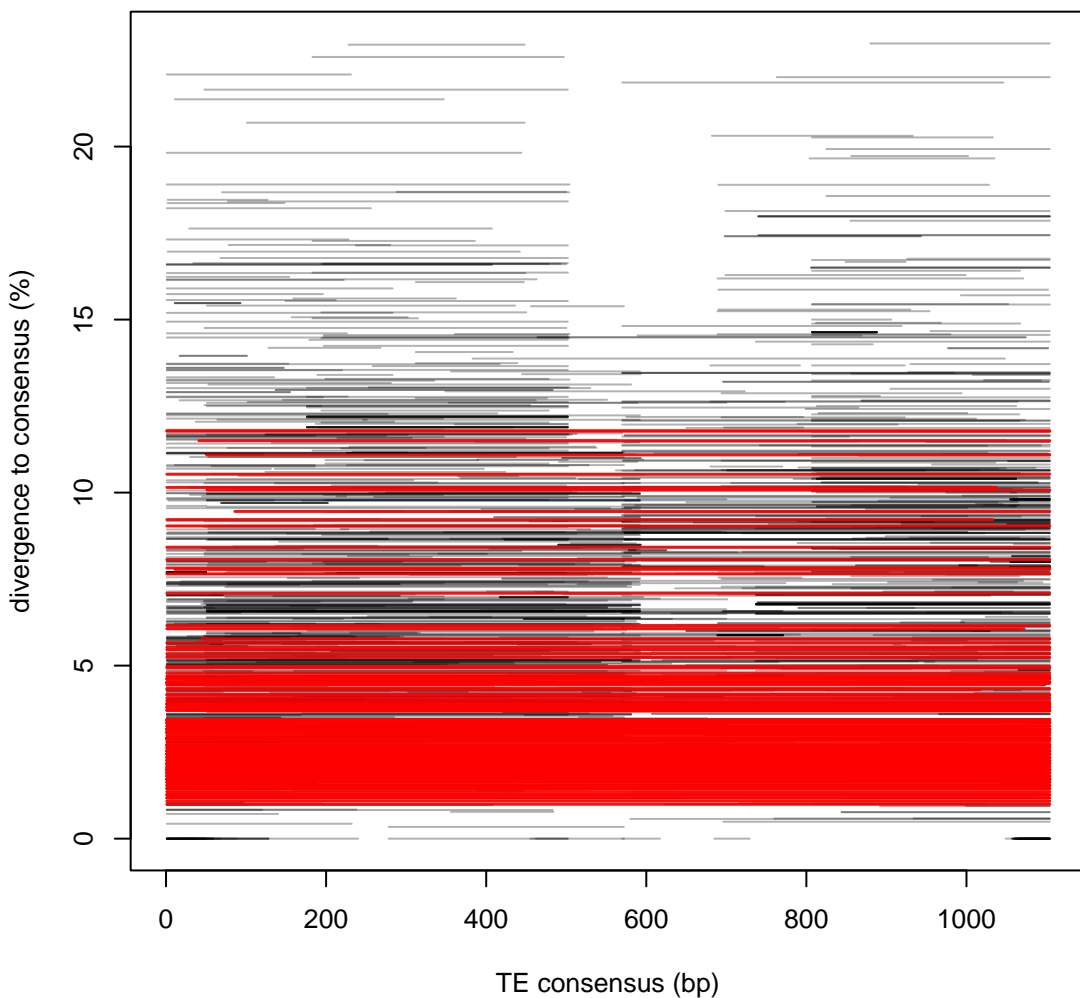
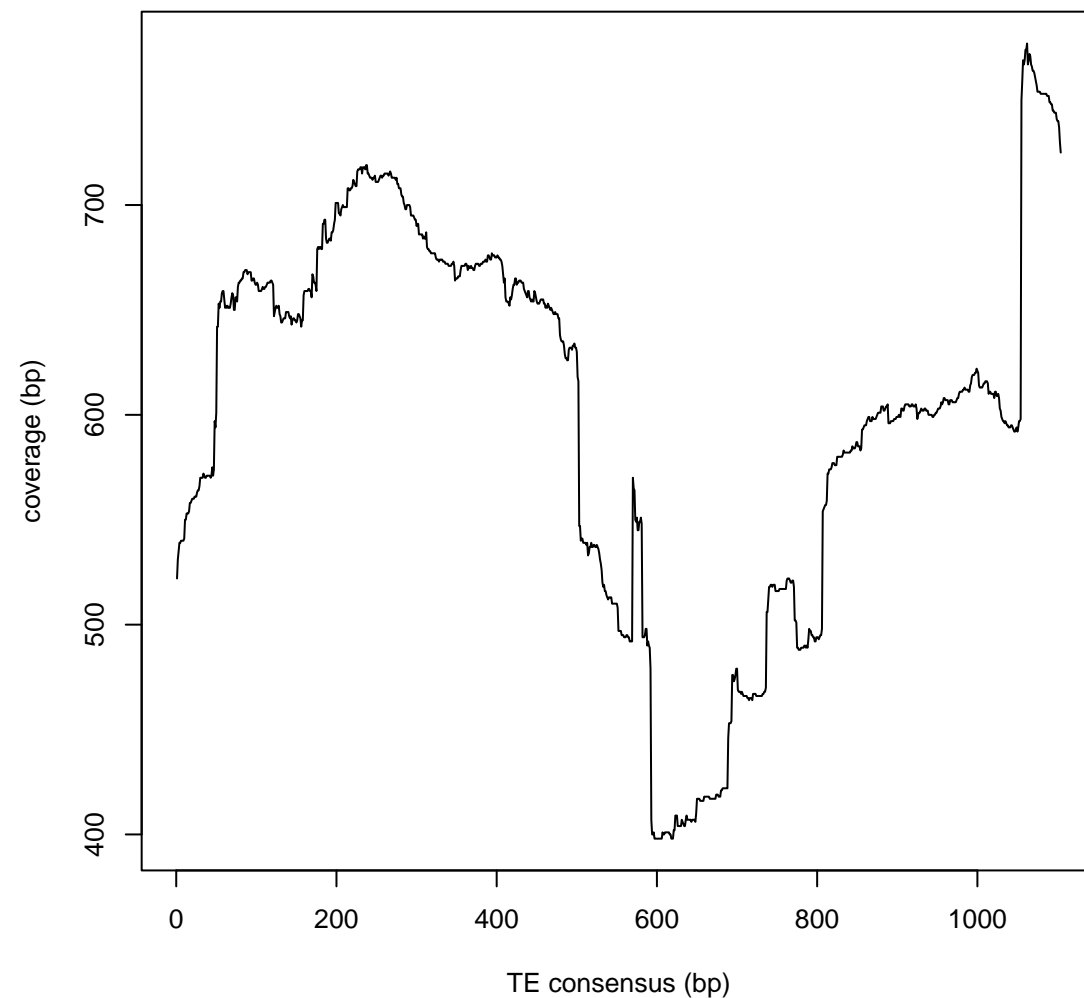


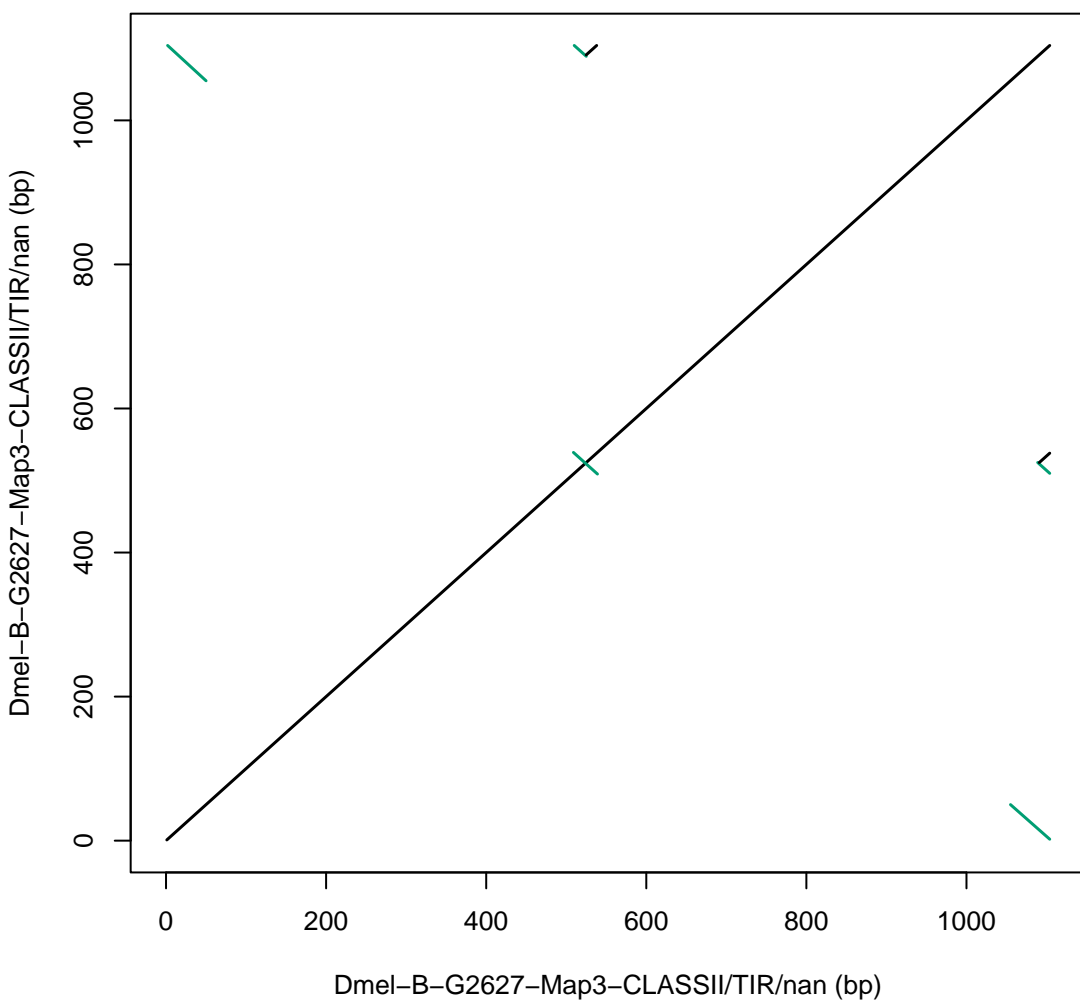
TE: Dmel-B-G2627-Map3-CLASSII/TIR/nan  
consensus size: 1104bp; fragments: 1827; full length: 165 (>=993.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

