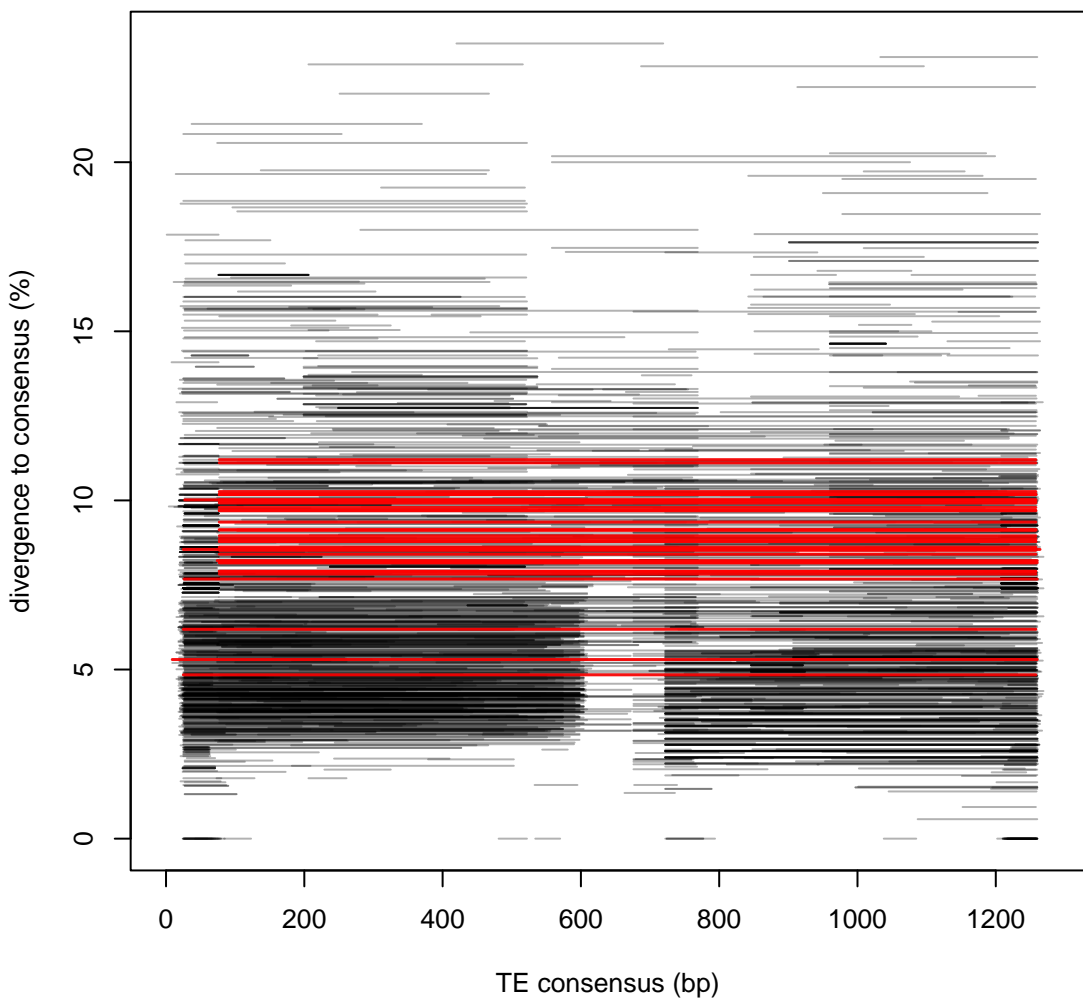
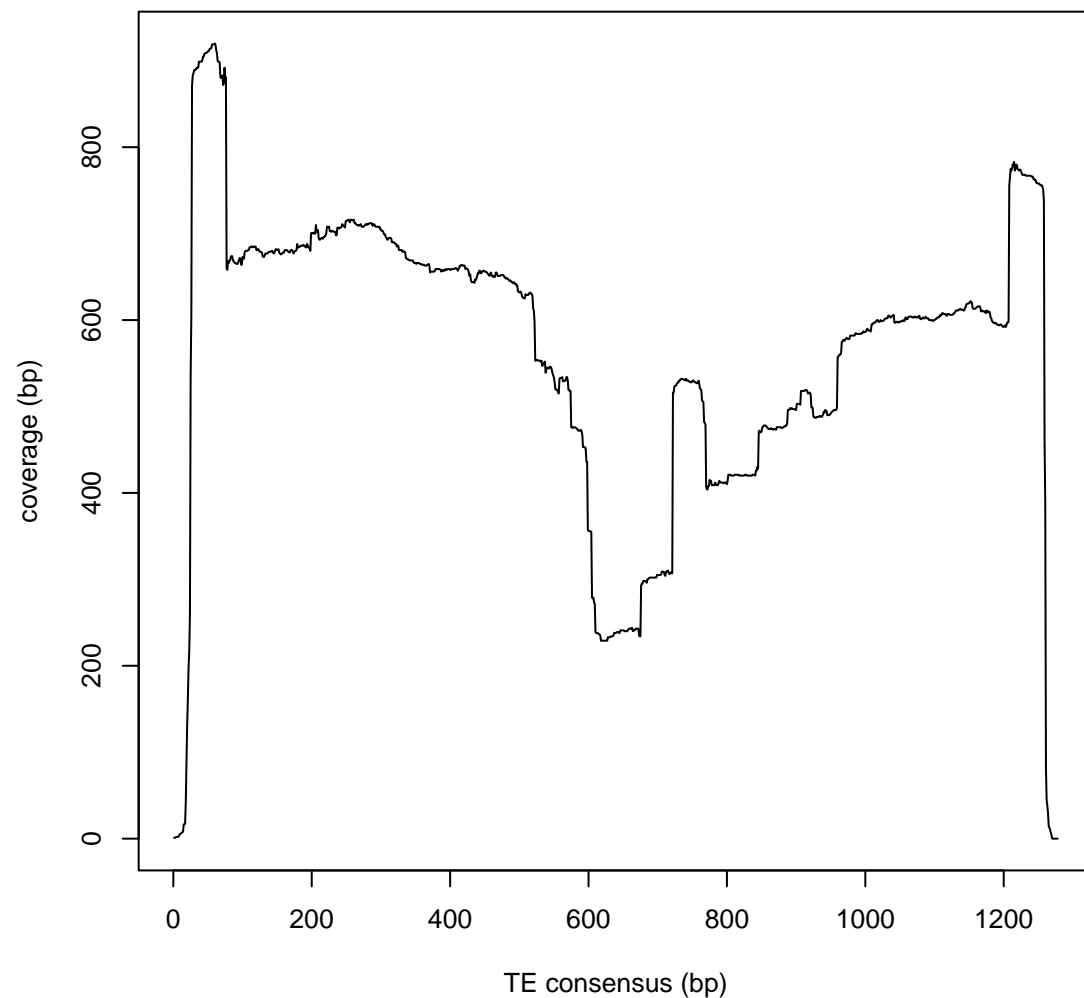


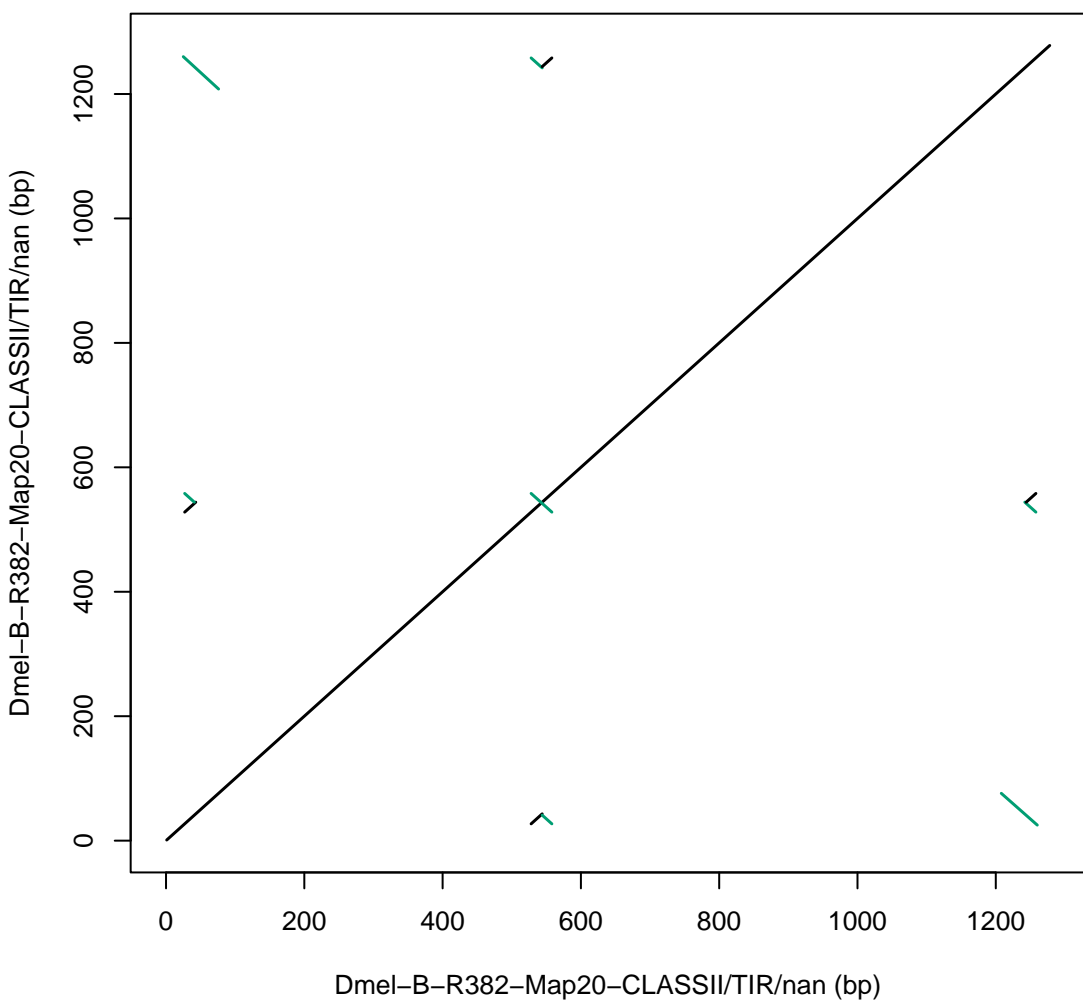
TE: Dmel-B-R382-Map20-CLASSII/TIR/nan
consensus size: 1278bp; fragments: 2379; full length: 39 (>=1150.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

