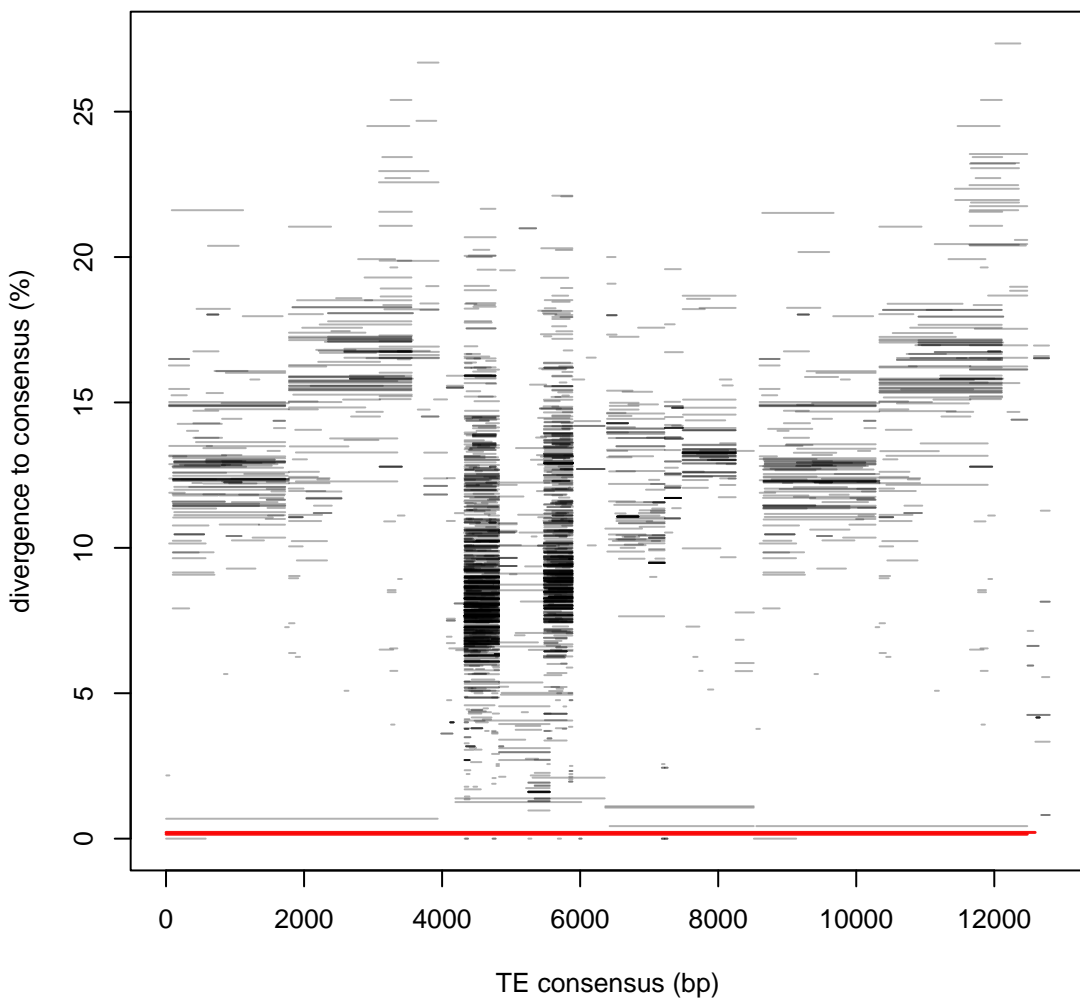
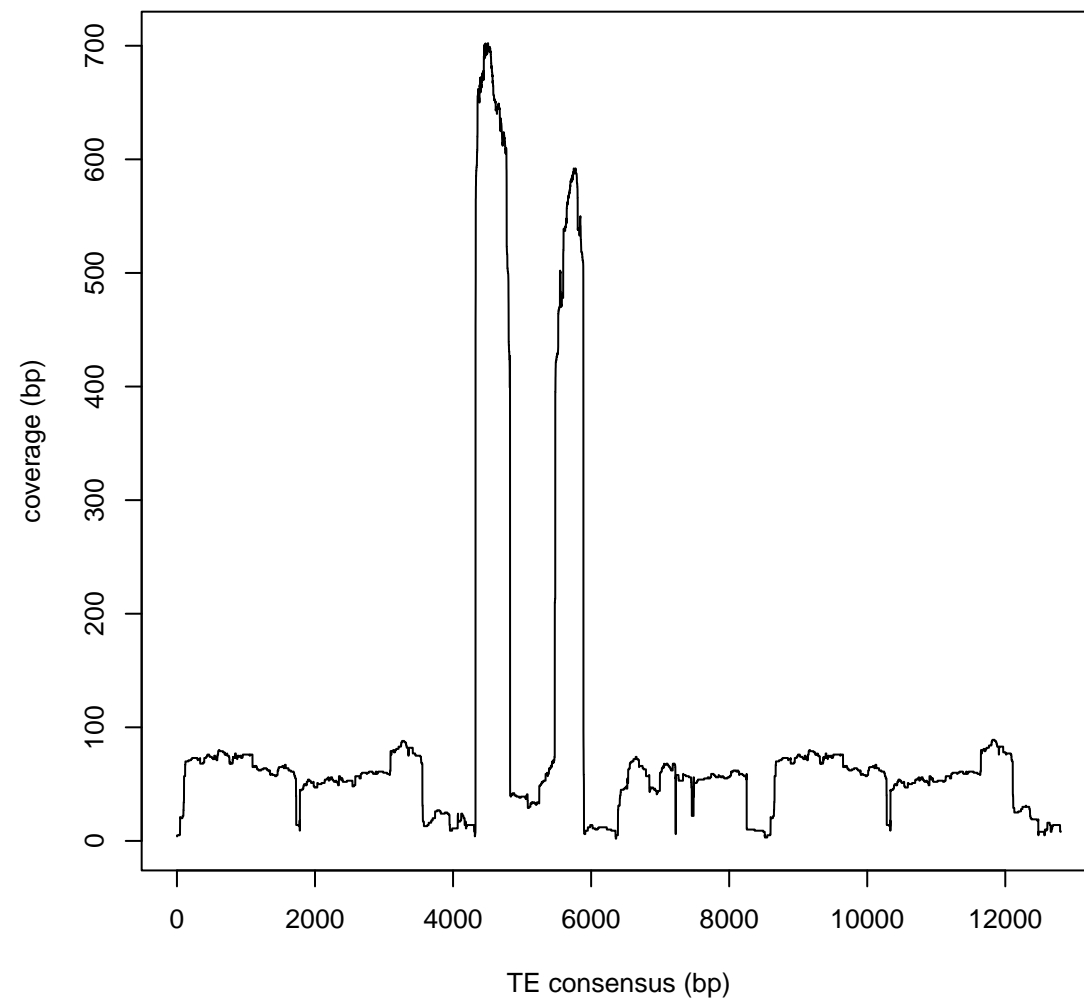


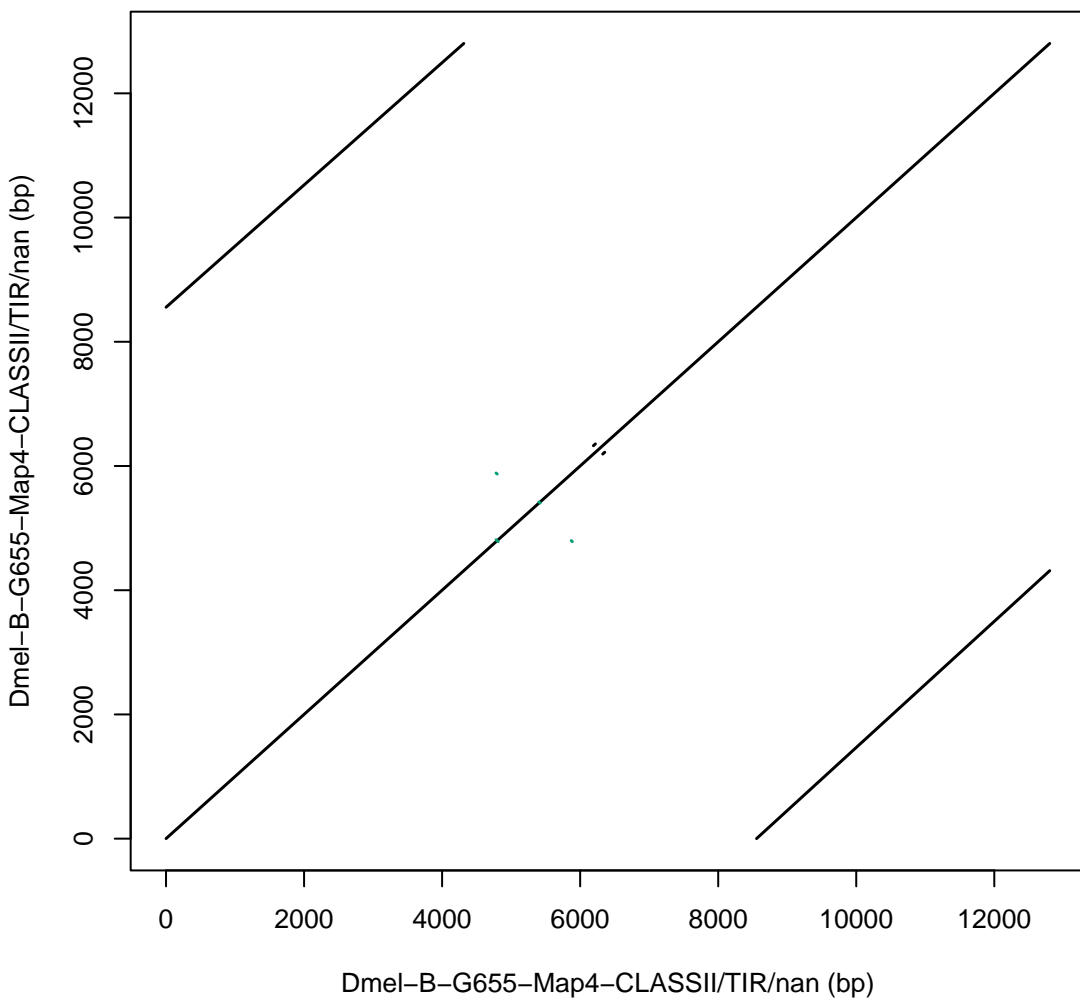
TE: Dmel-B-G655-Map4-CLASSII/TIR/nan  
consensus size: 12803bp; fragments: 2549; full length: 2 (>=11522.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

