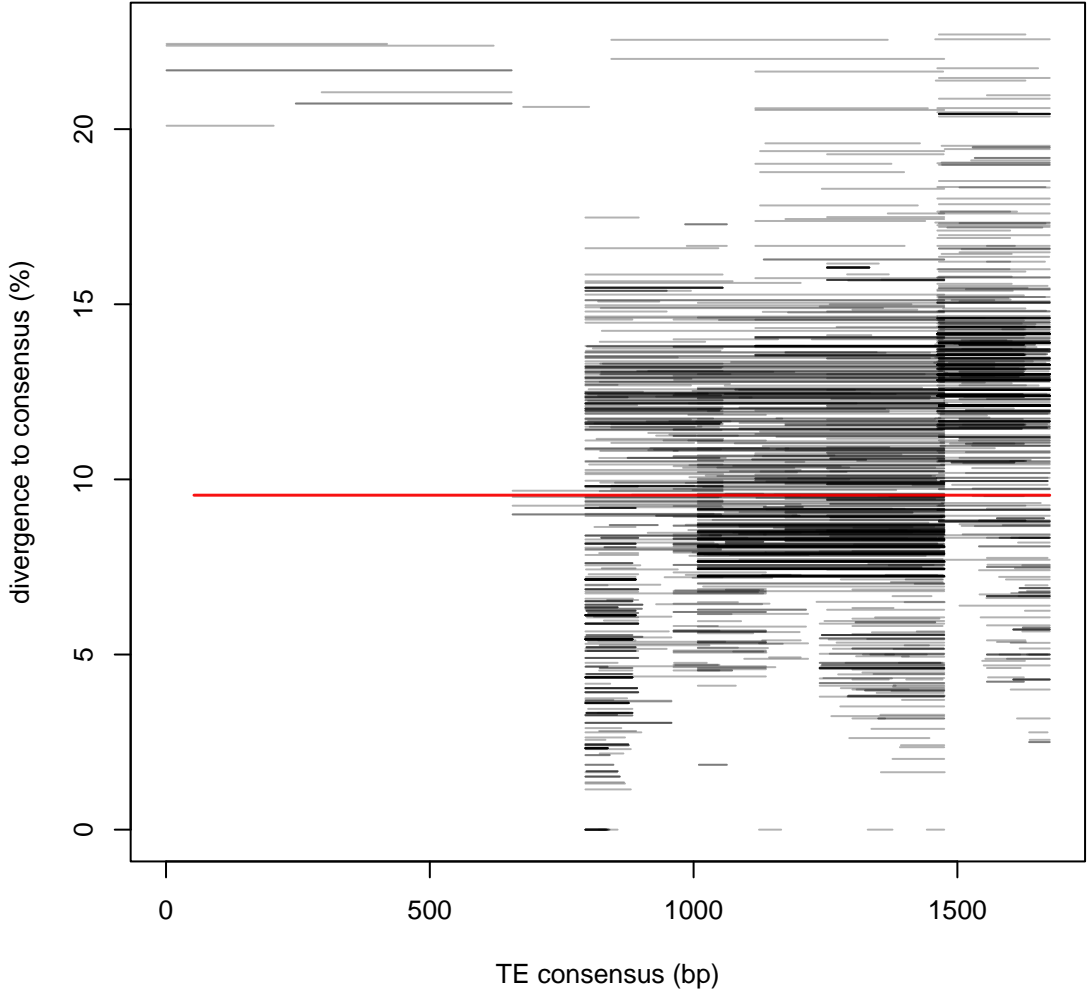
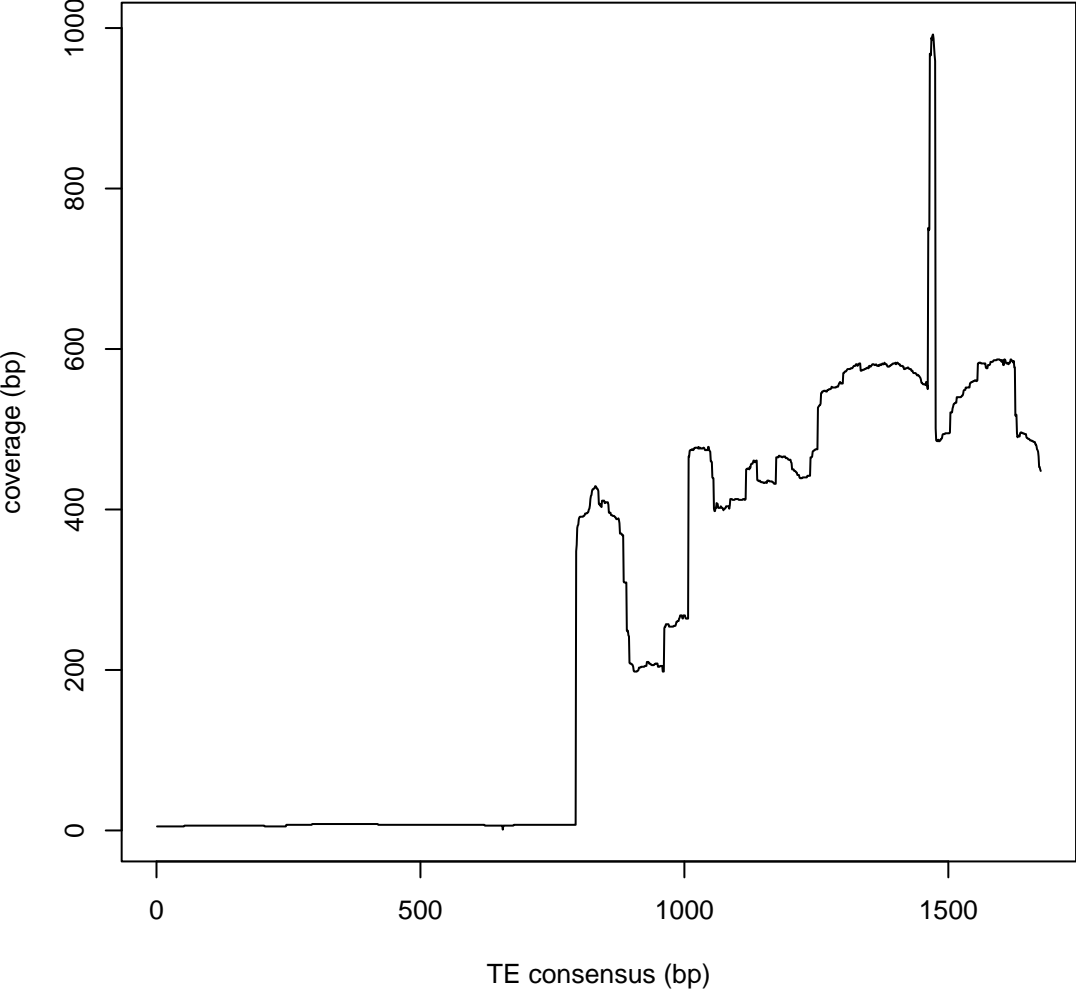


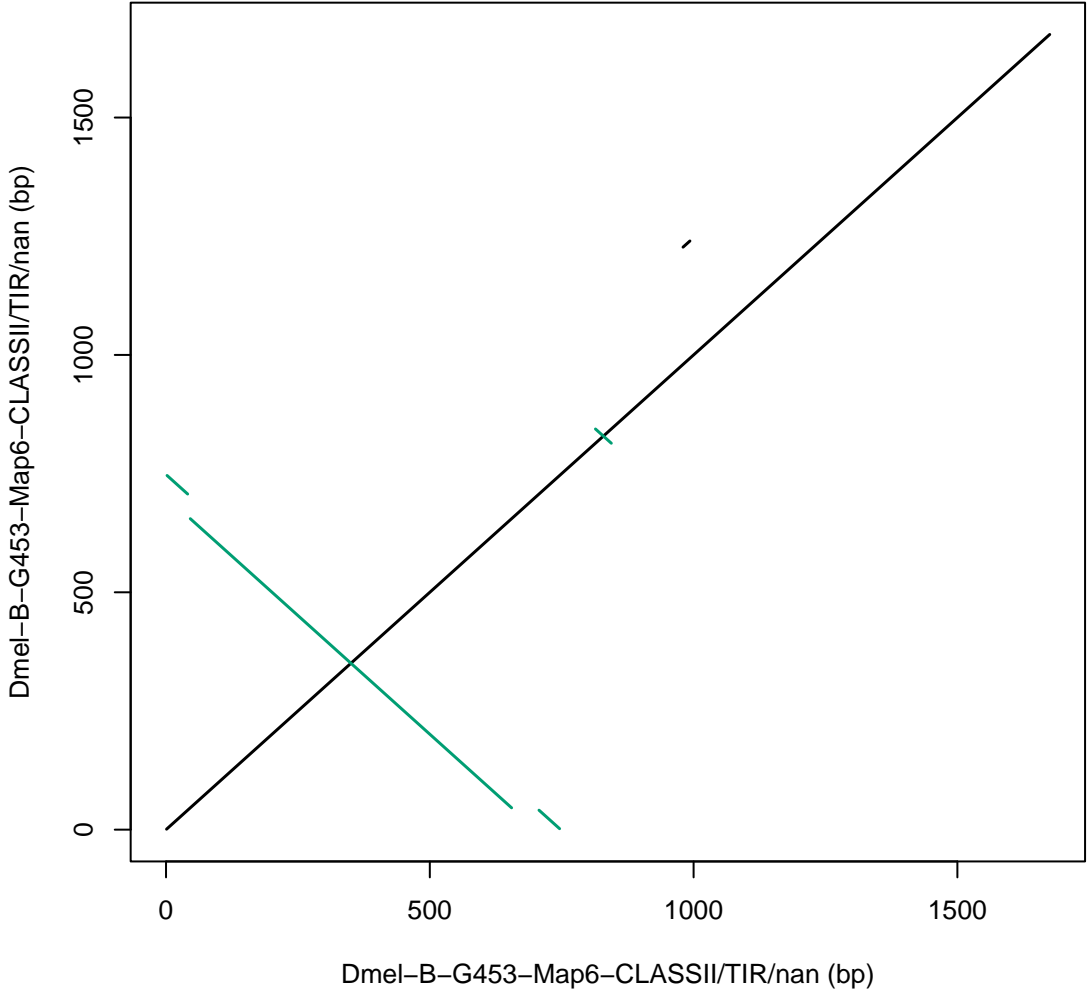
TE: Dmel-B-G453-Map6-CLASSII/TIR/nan
consensus size: 1675bp; fragments: 1734; full length: 1 (>=1507.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

