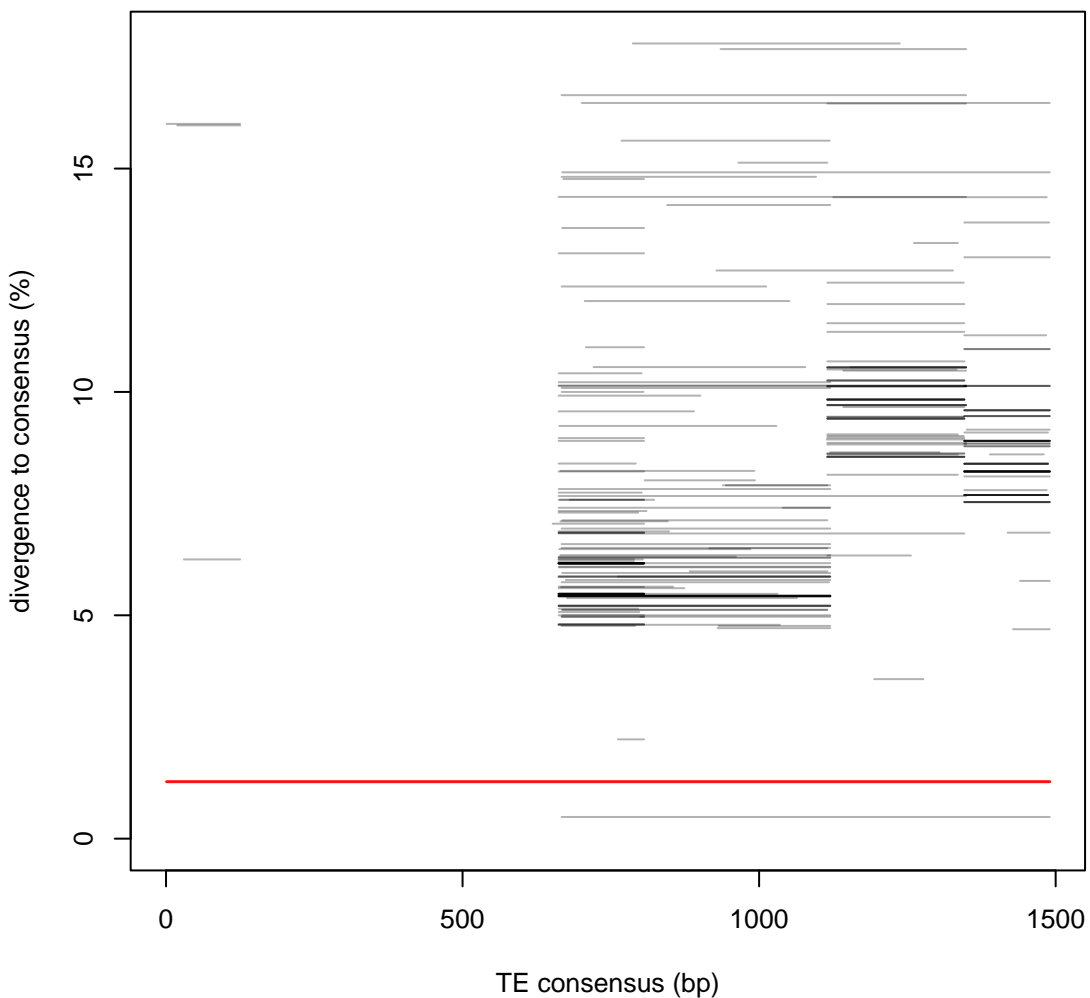
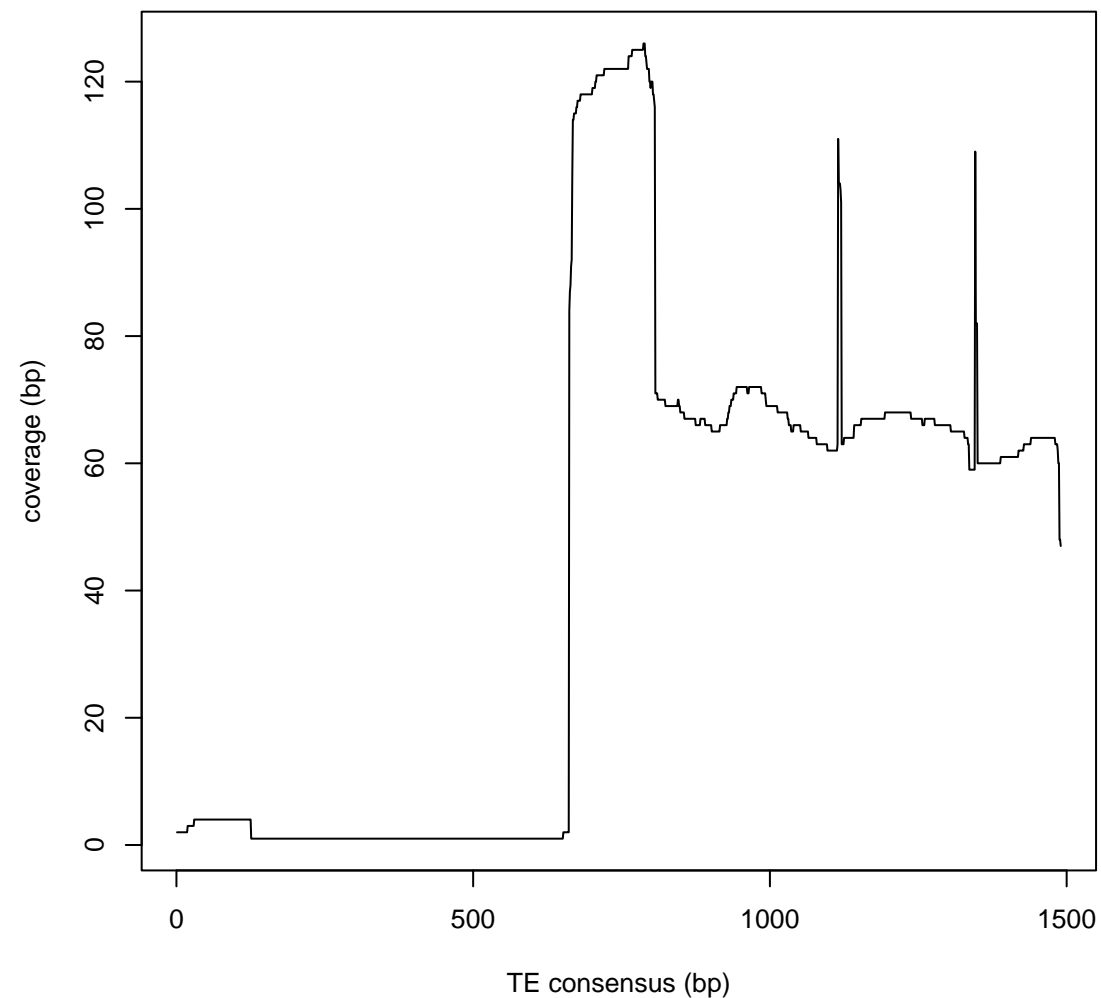


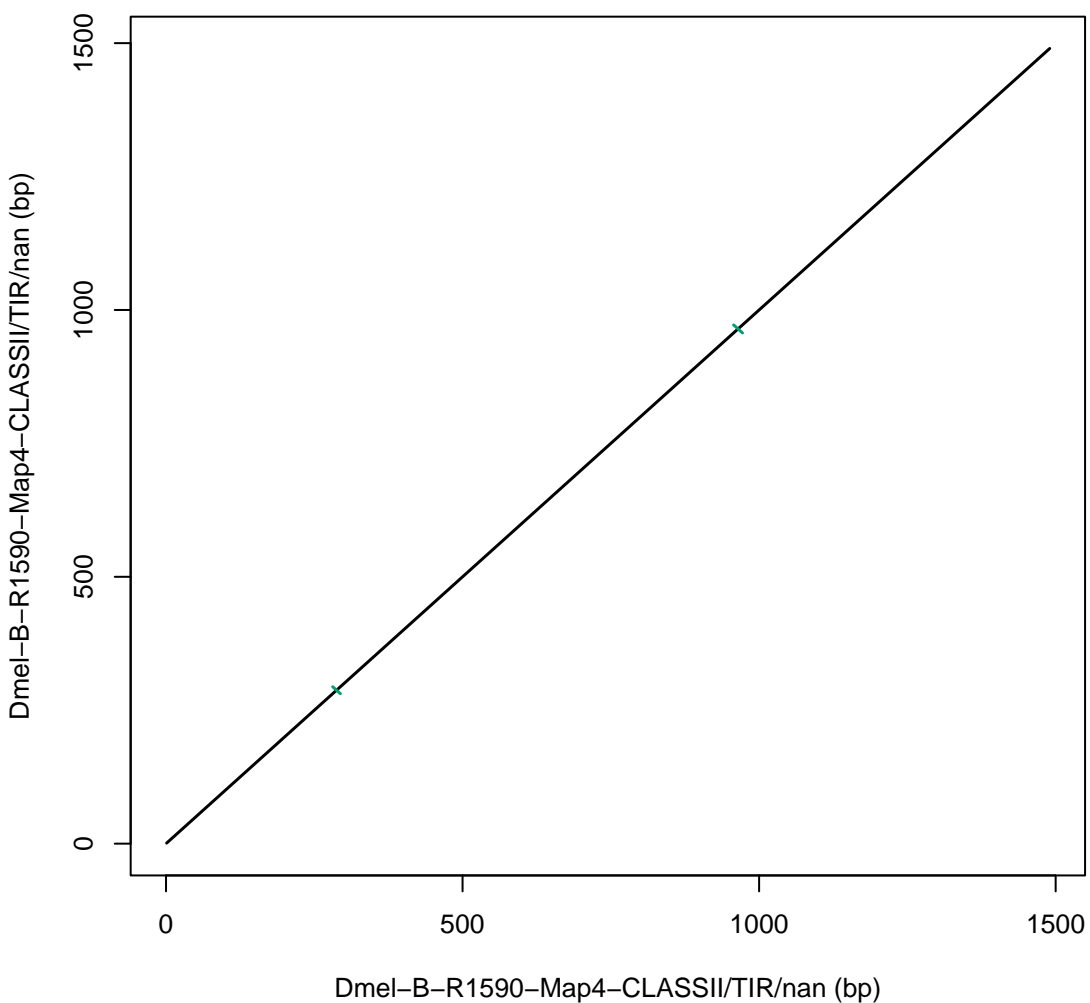
TE: Dmel-B-R1590-Map4-CLASSII/TIR/nan
consensus size: 1490bp; fragments: 257; full length: 1 (>=1341bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

