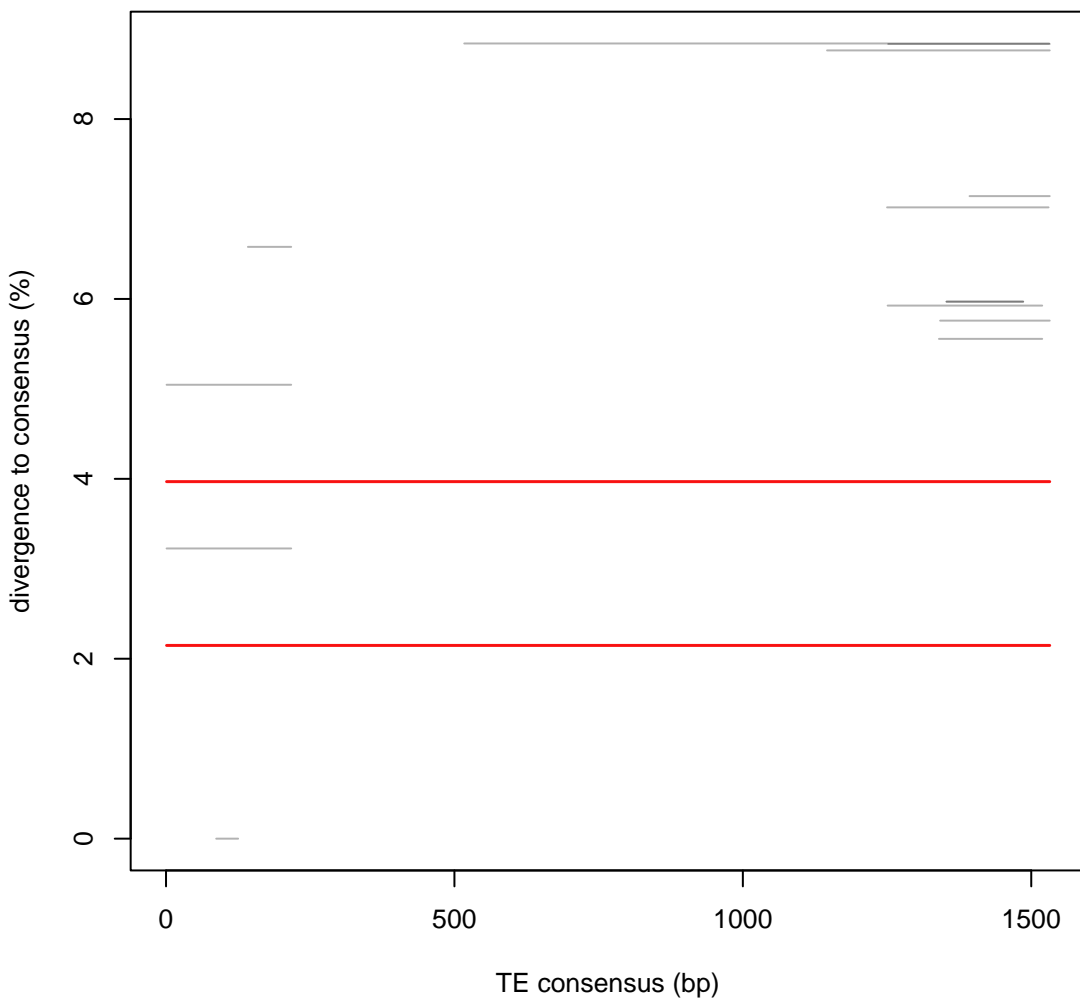
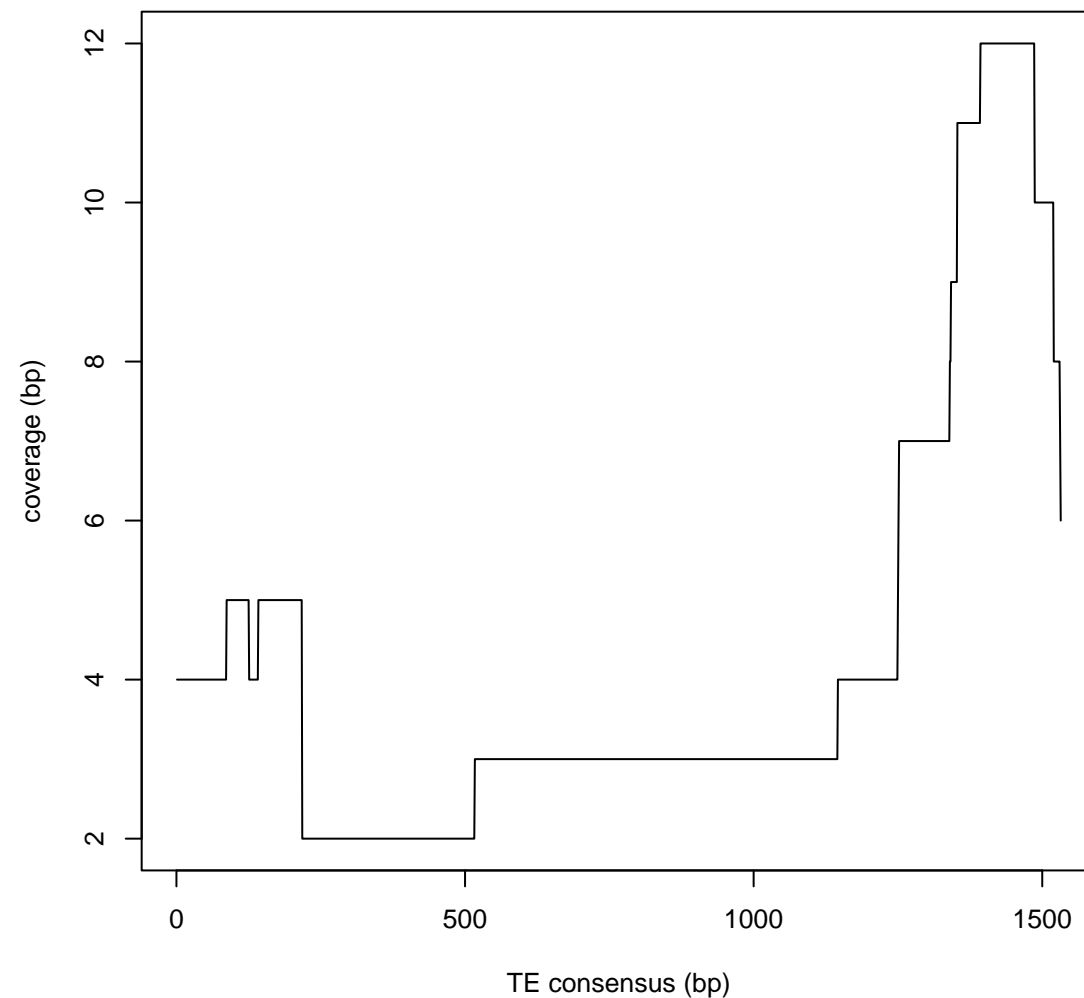


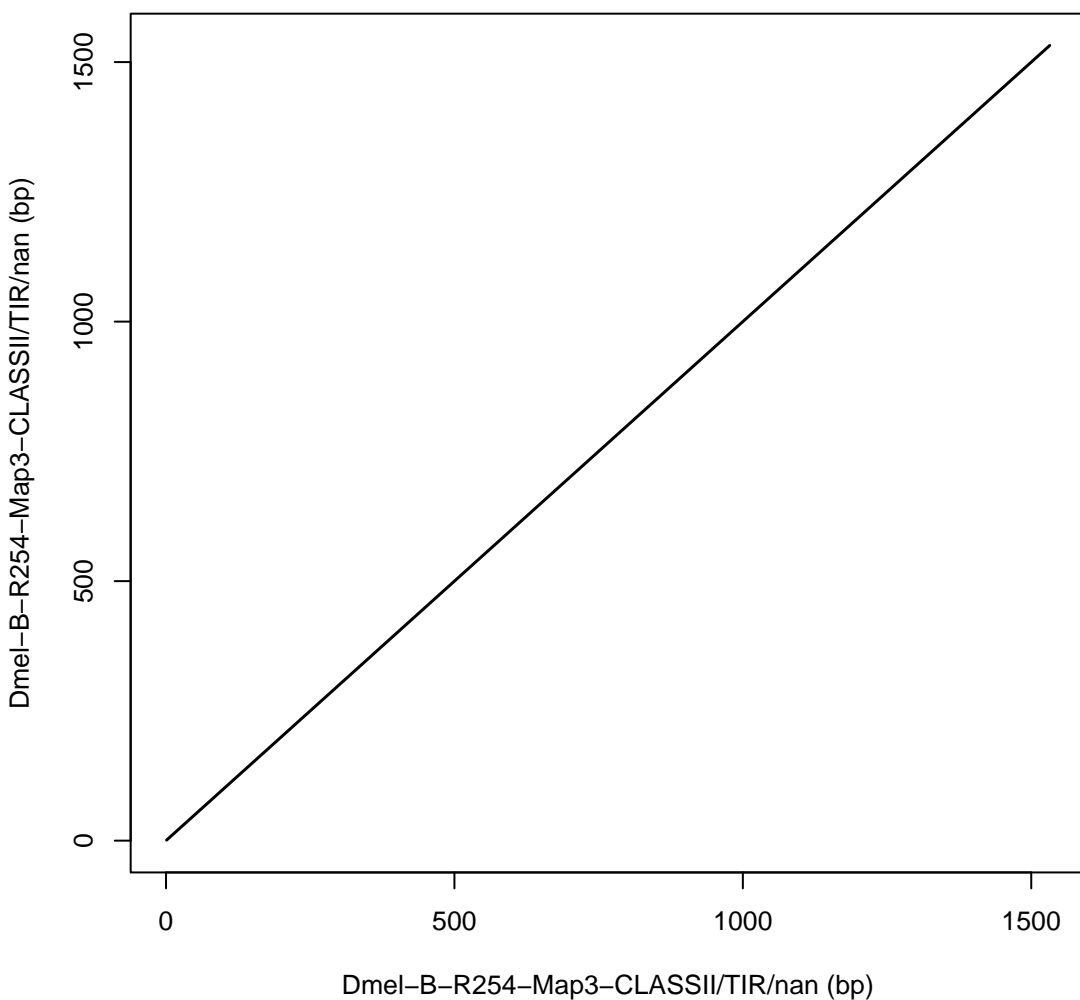
TE: Dmel-B-R254-Map3-CLASSII/TIR/nan
consensus size: 1532bp; fragments: 16; full length: 2 (≥ 1378.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

