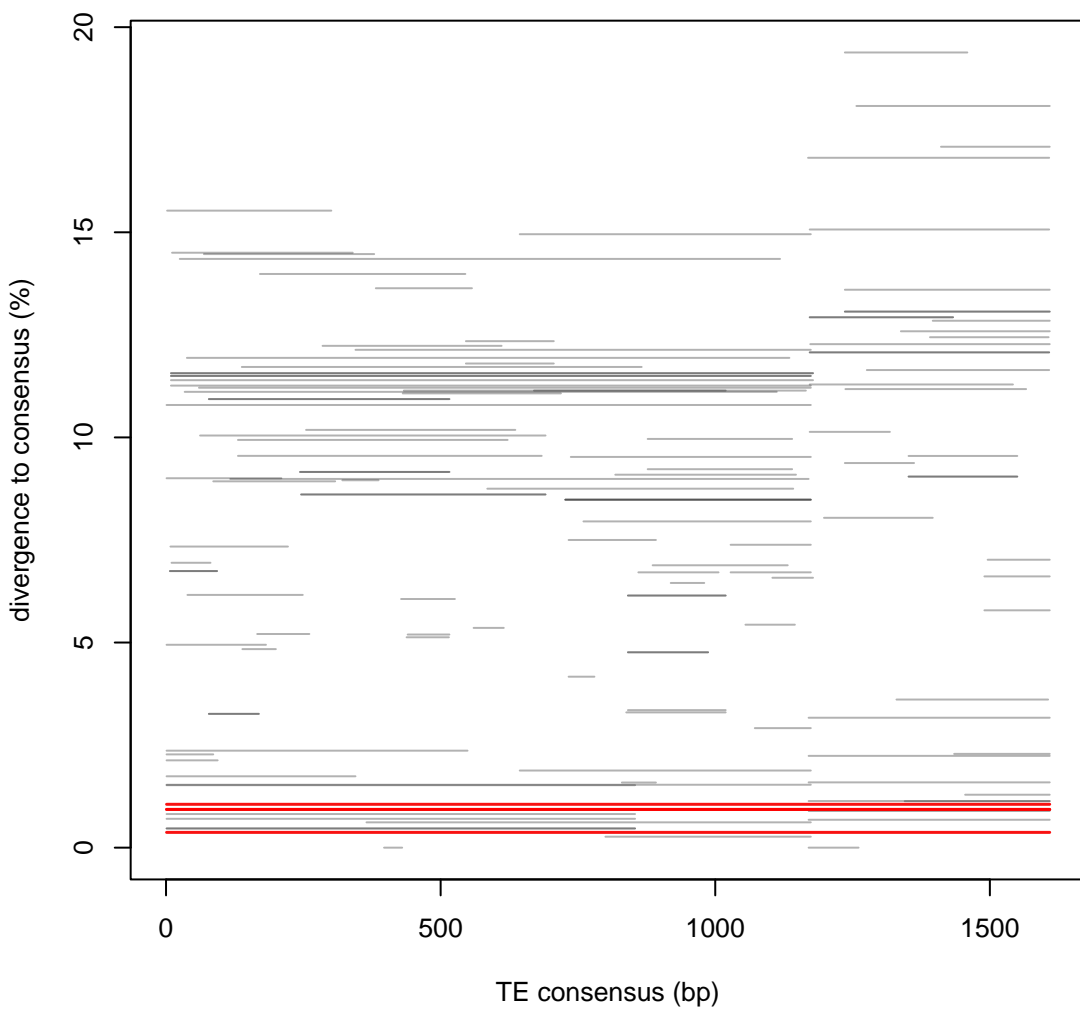
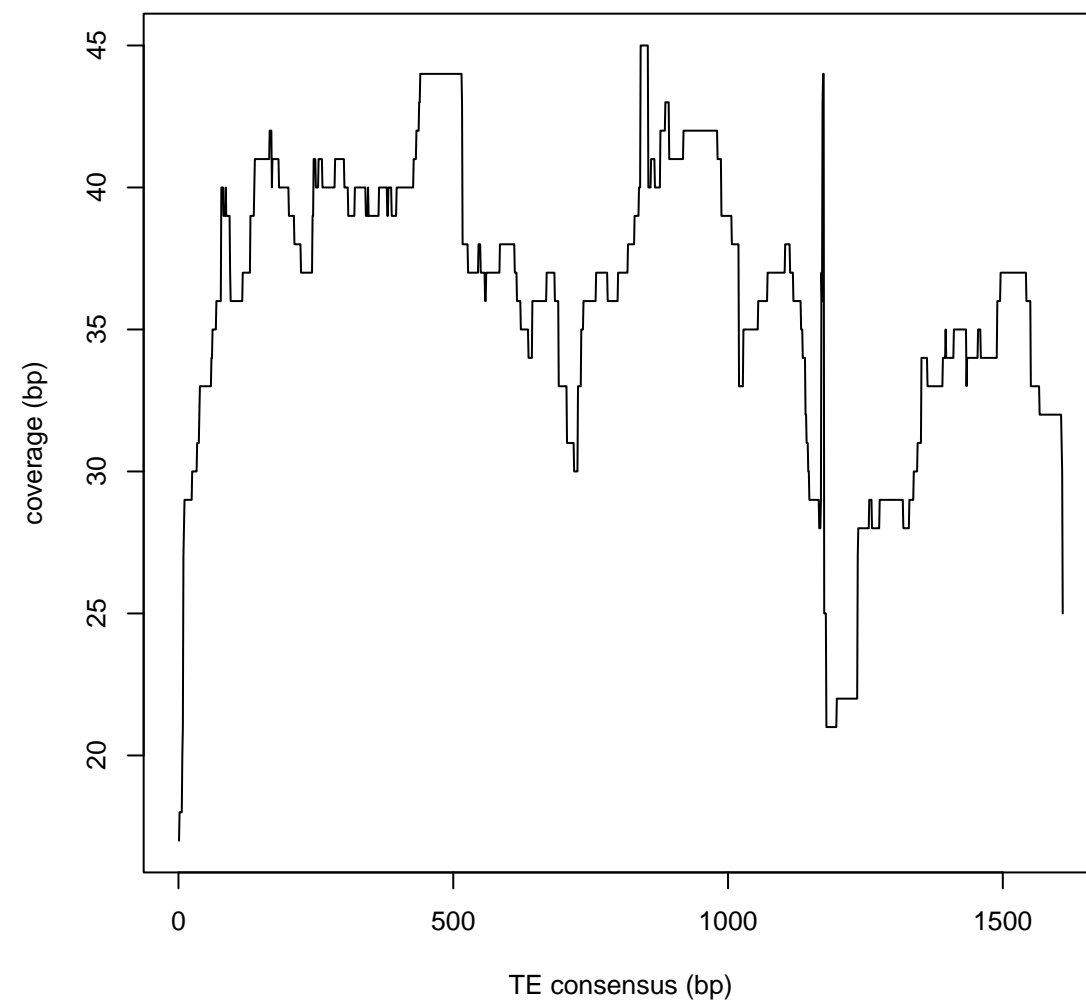


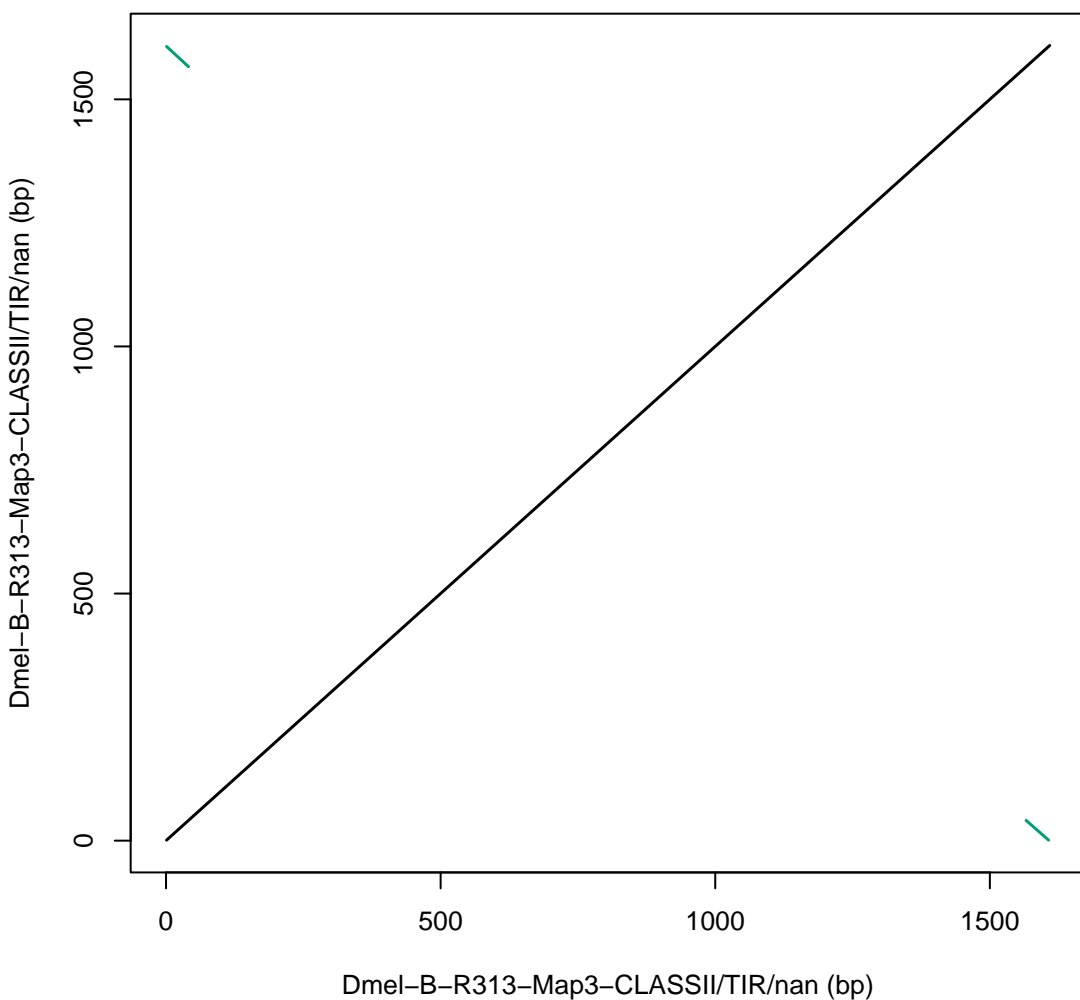
TE: Dmel-B-R313-Map3-CLASSII/TIR/nan
consensus size: 1609bp; fragments: 137; full length: 4 (>=1448.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

