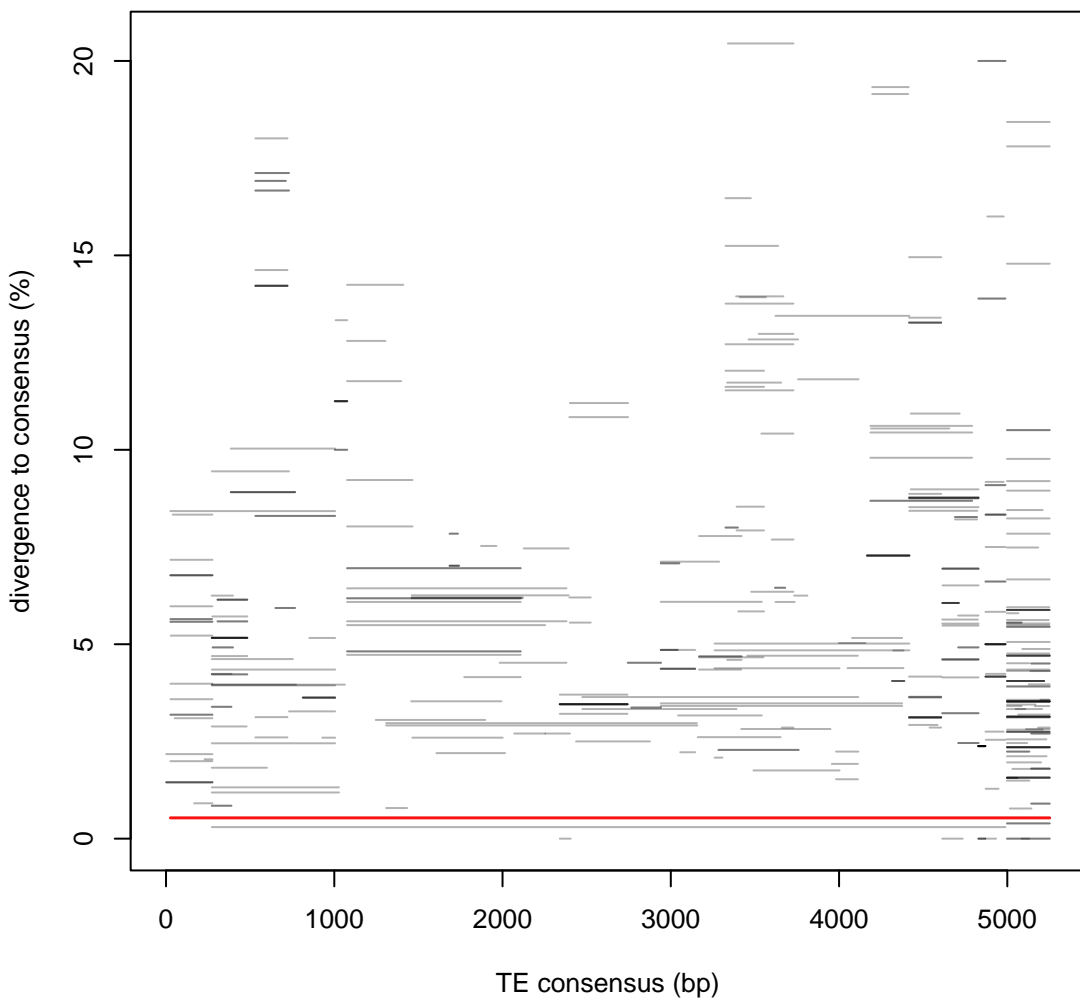
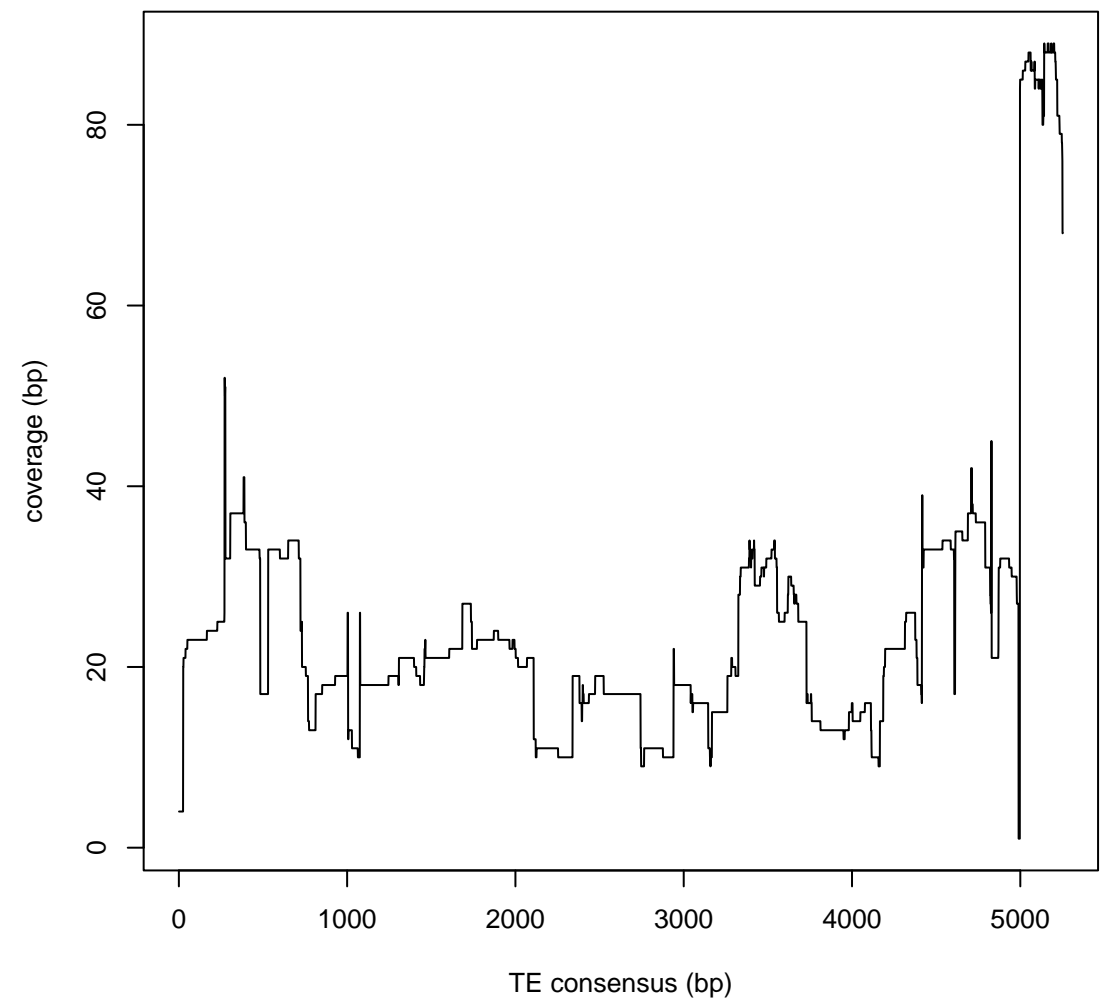


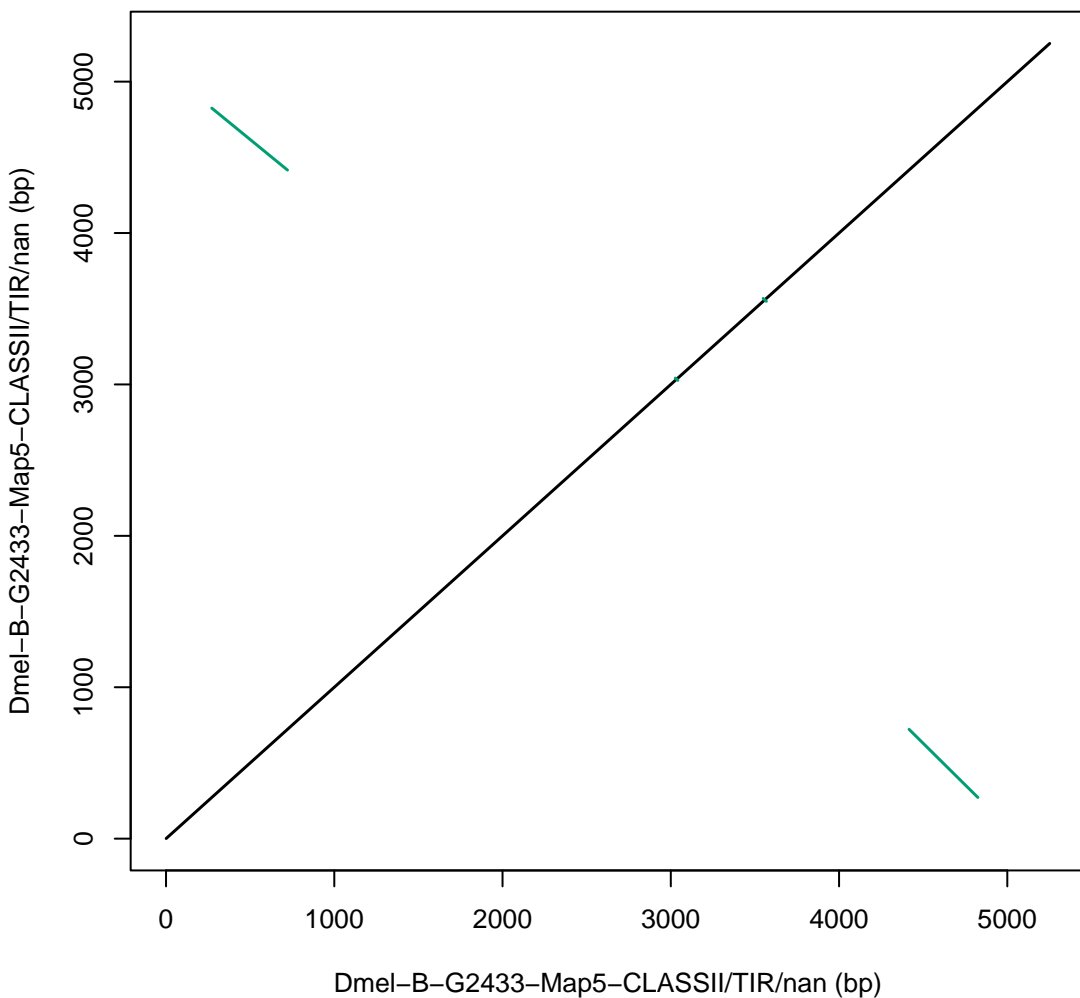
TE: Dmel-B-G2433-Map5-CLASSII/TIR/nan
consensus size: 5252bp; fragments: 436; full length: 1 (≥ 4726.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

