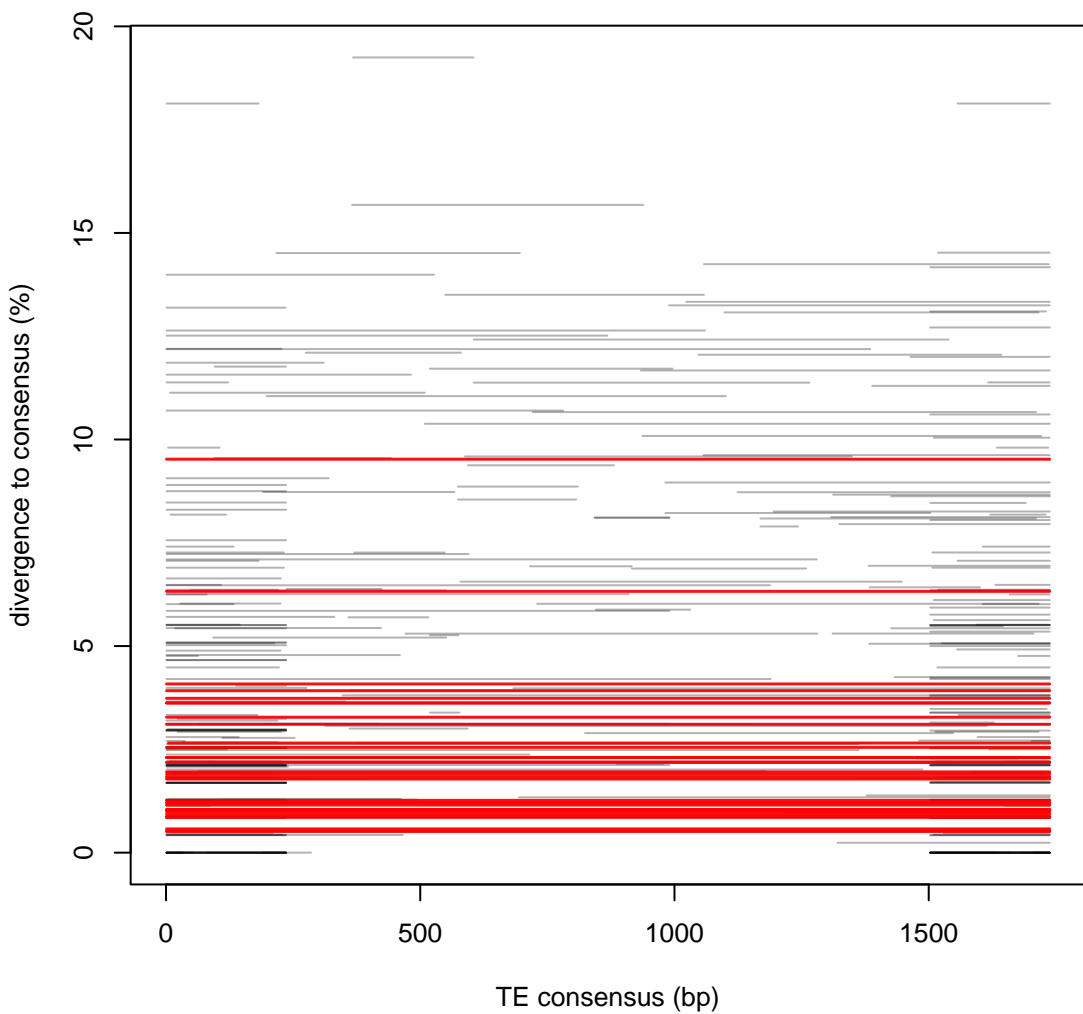
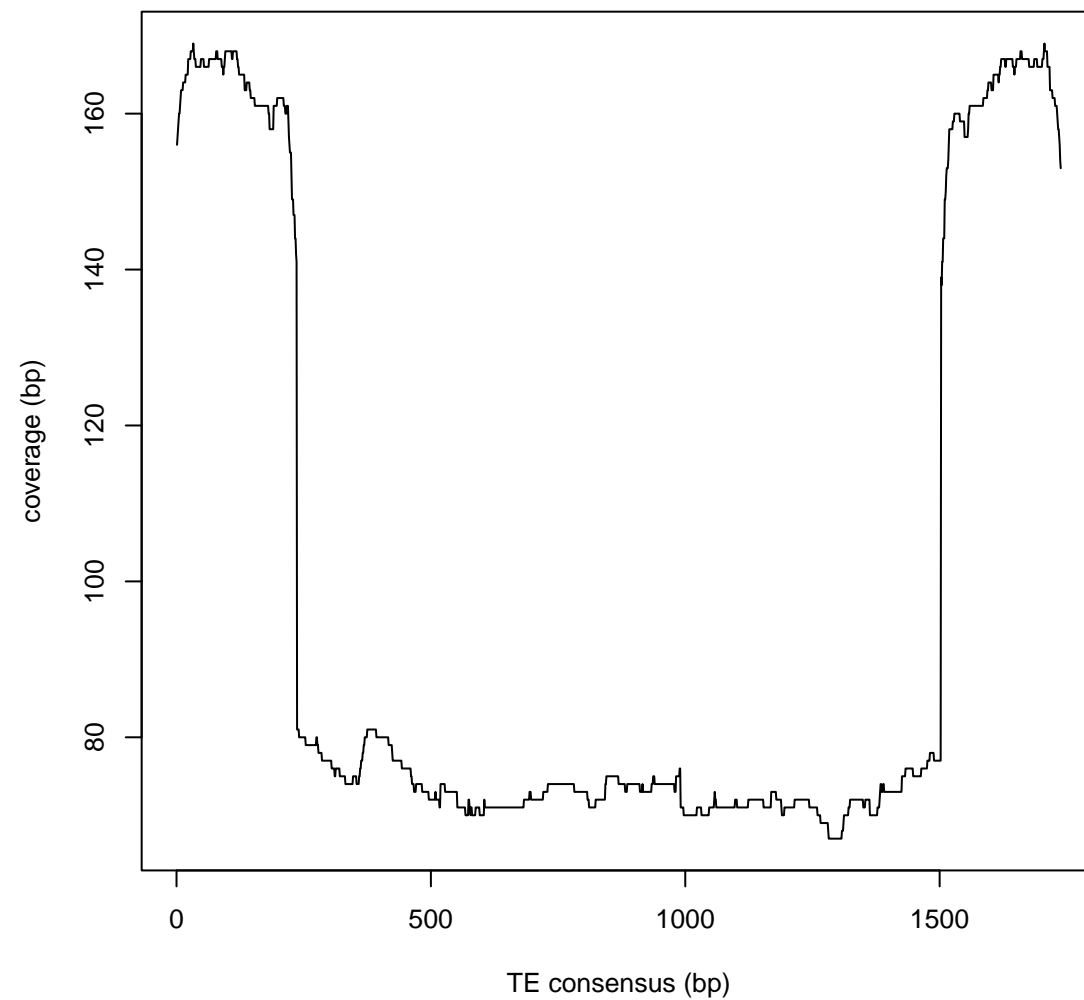


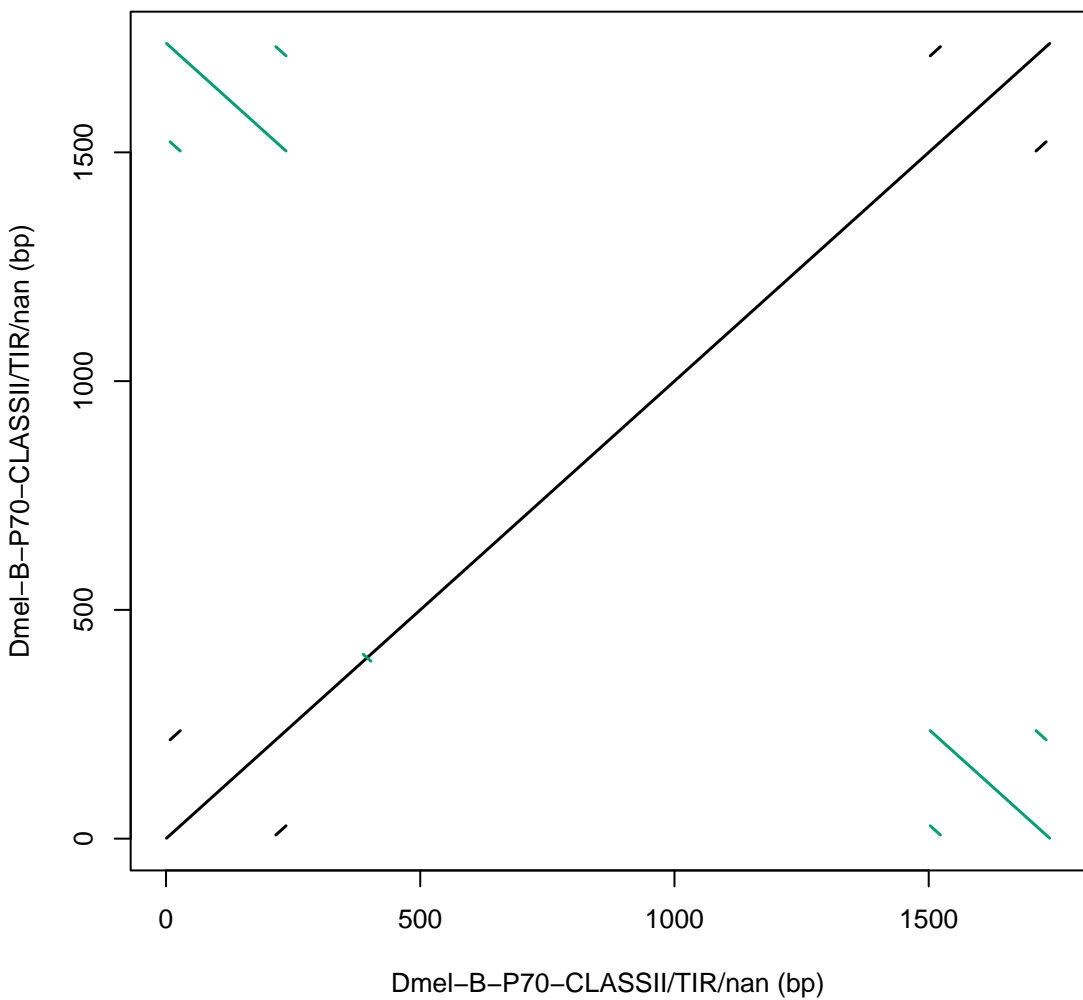
TE: Dmel-B-P70-CLASSII/TIR/nan
consensus size: 1738bp; fragments: 358; full length: 35 (≥ 1564.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

