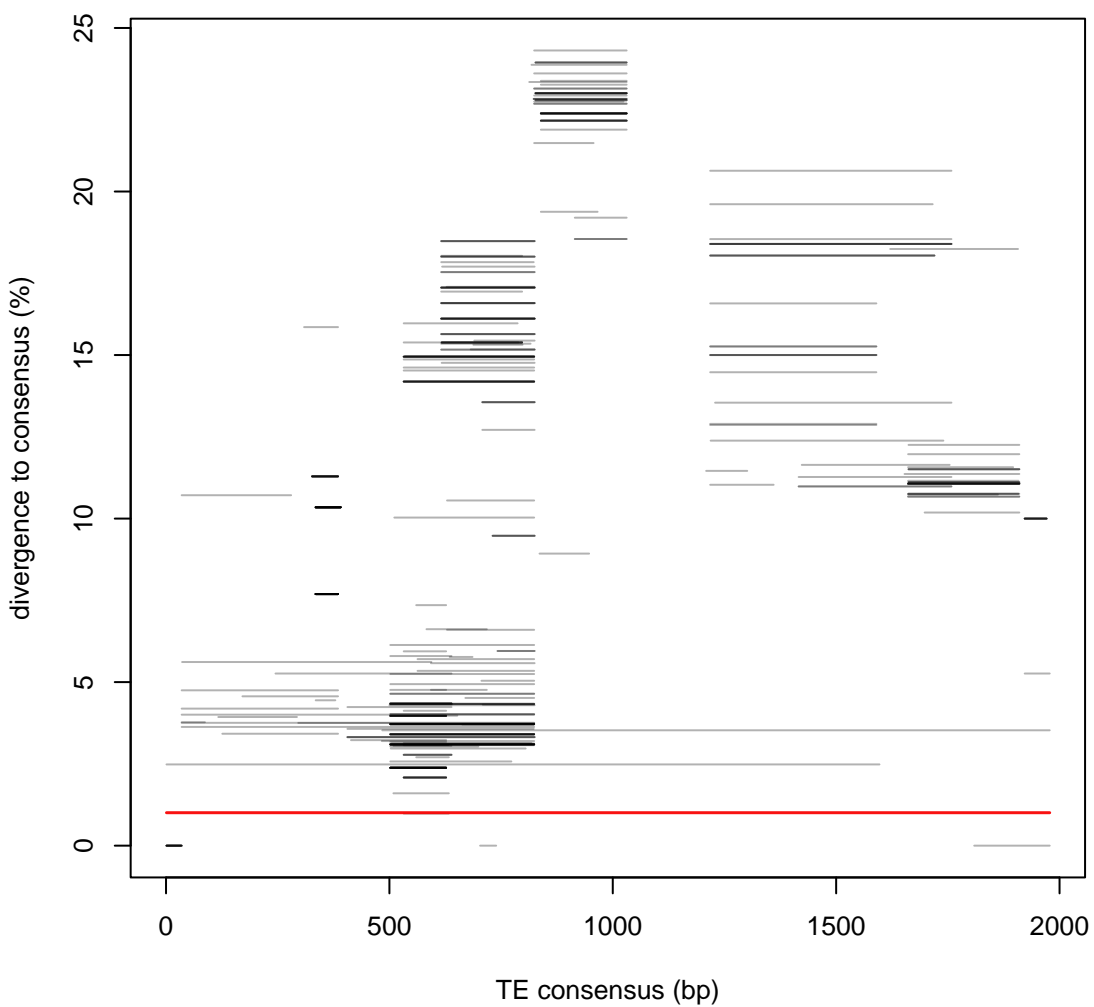
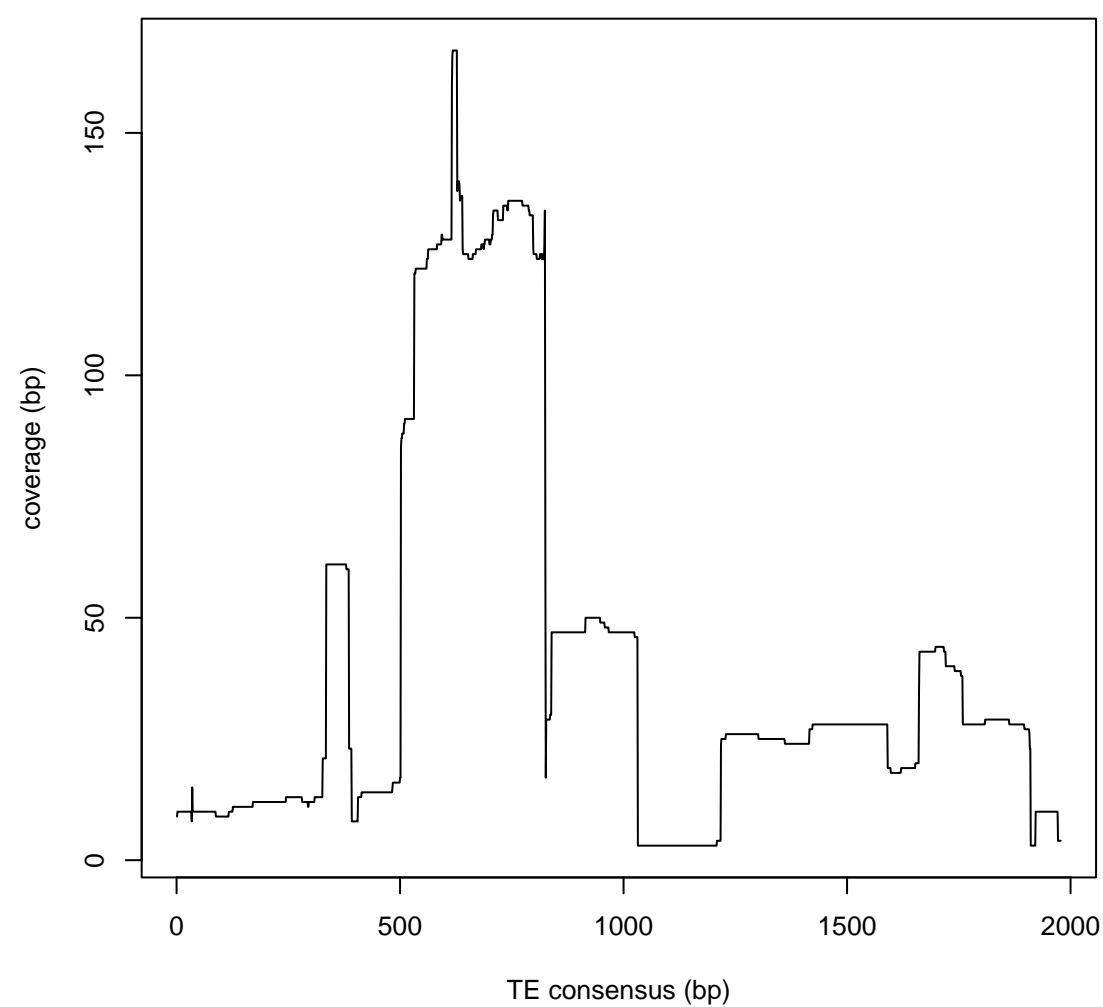


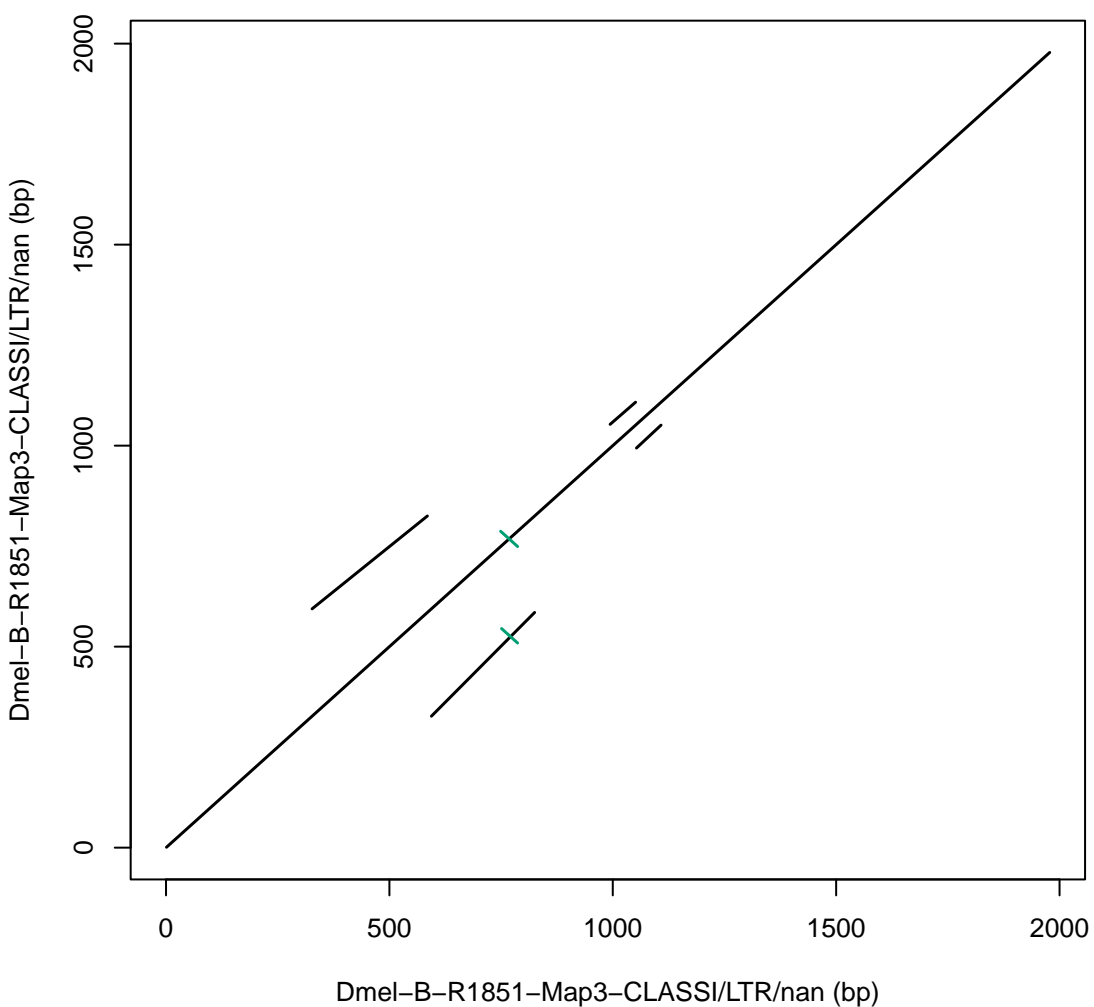
TE: Dmel-B-R1851-Map3-CLASSI/LTR/nan
consensus size: 1978bp; fragments: 361; full length: 1 (≥ 1780.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

