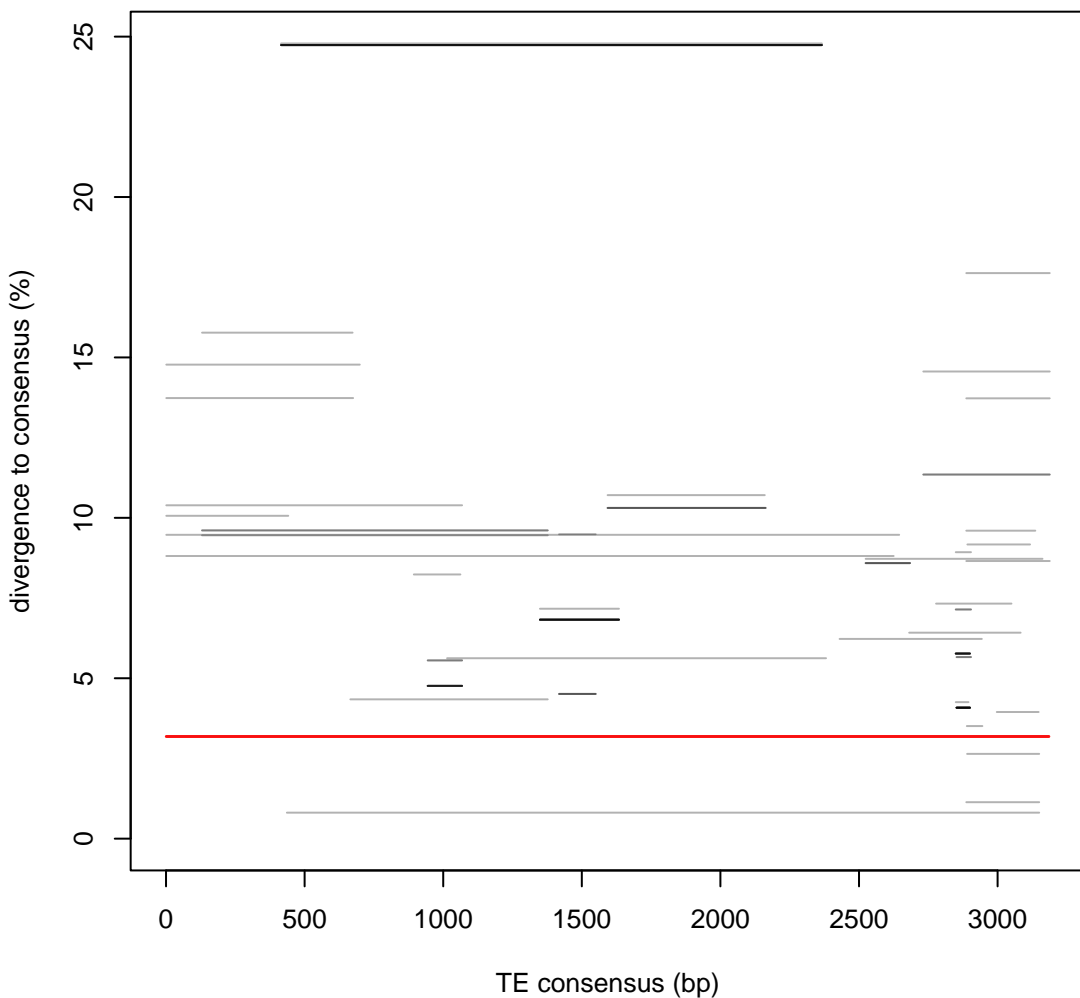
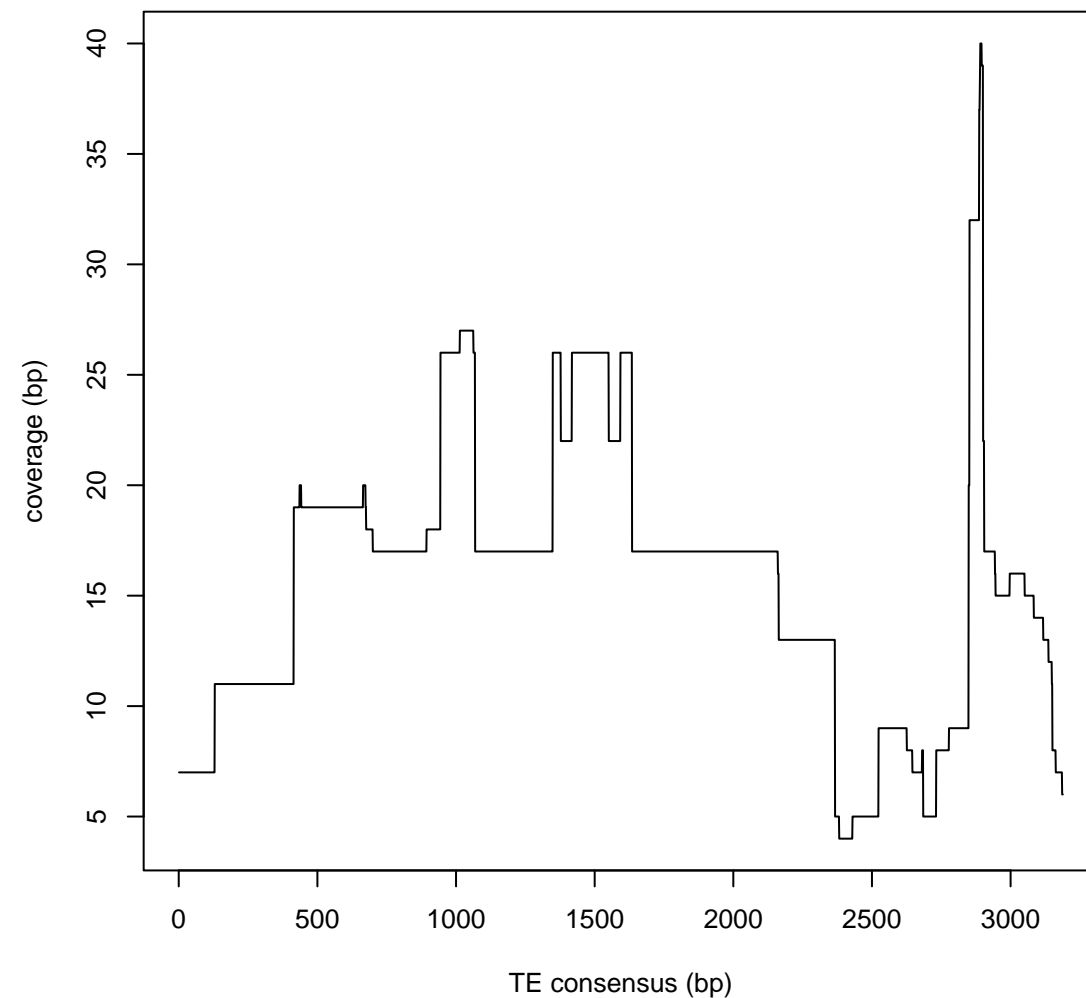


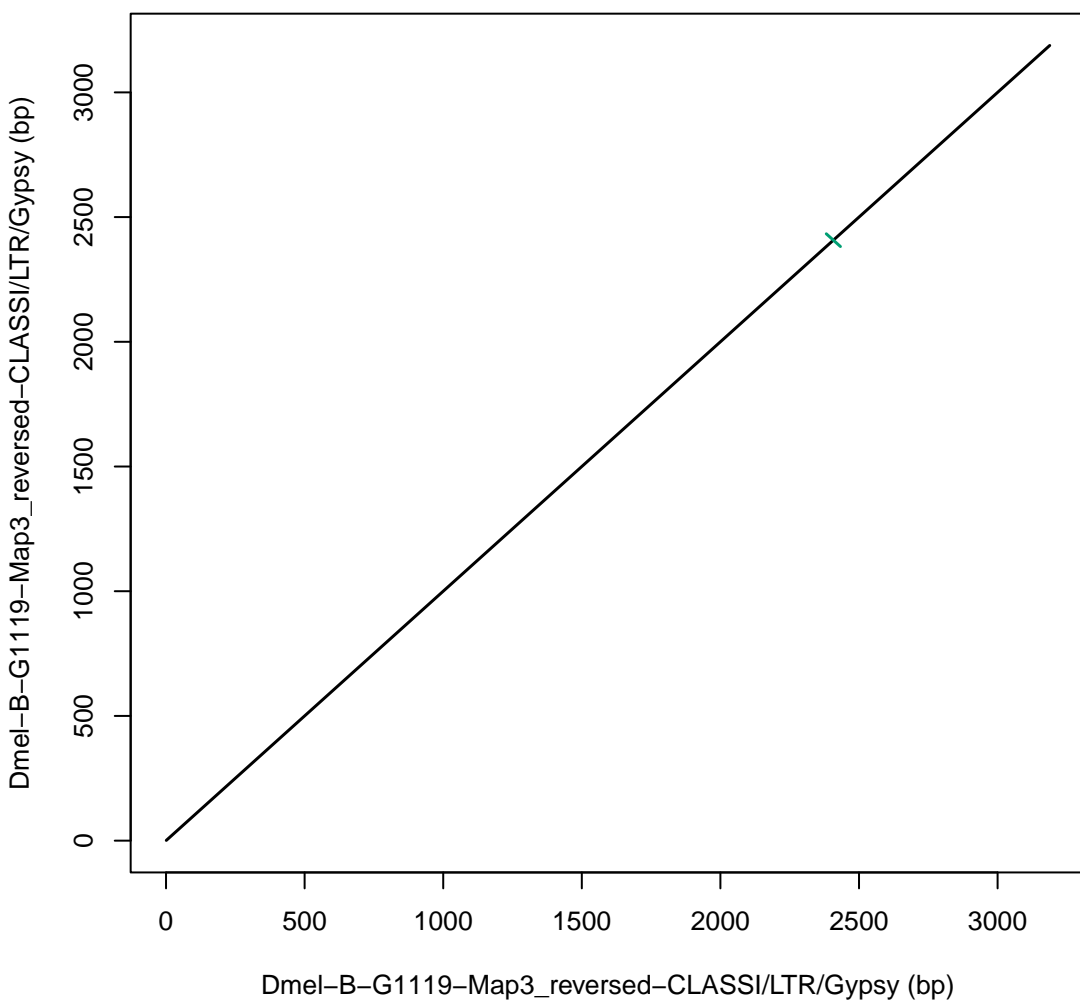
TE: Dmel-B-G1119-Map3_reversed-CLASSI/LTR/Gypsy
consensus size: 3188bp; fragments: 90; full length: 1 (≥ 2869.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

