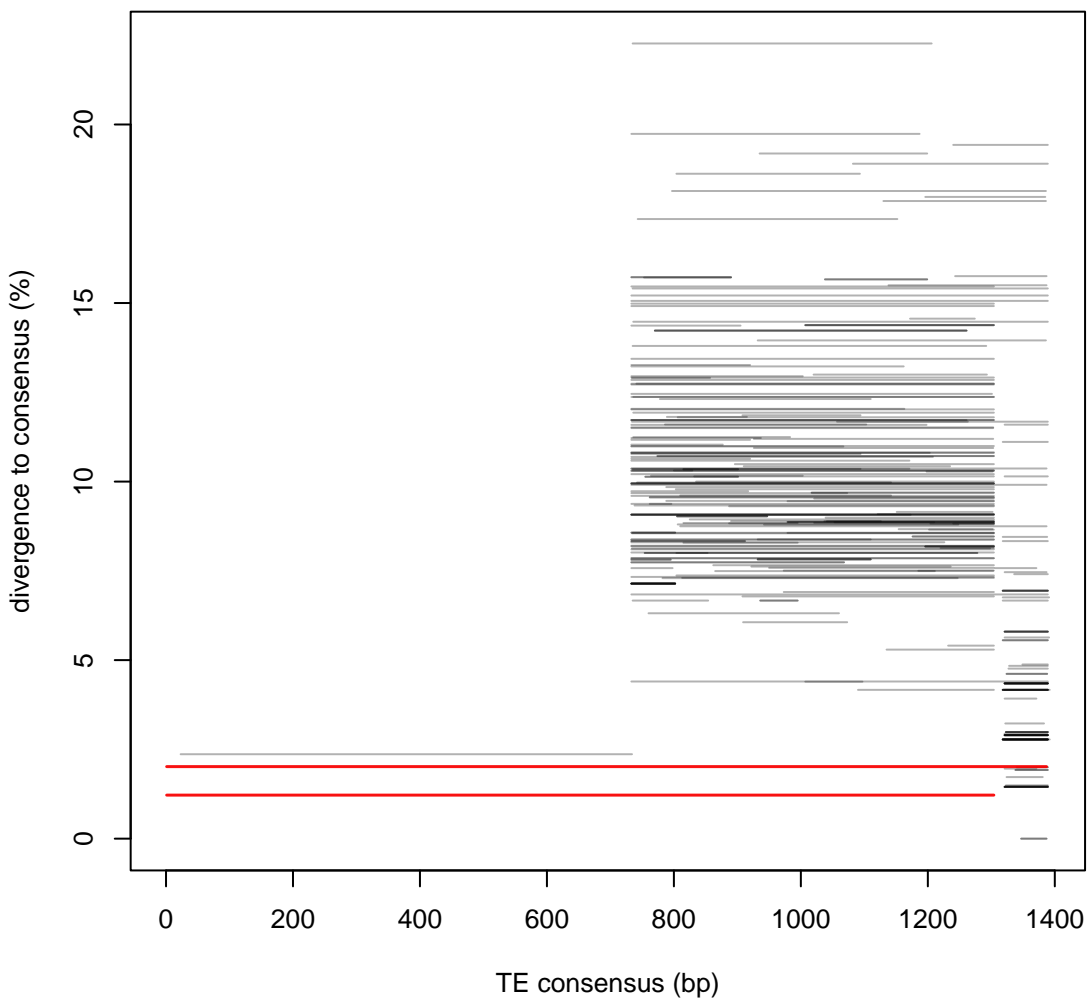
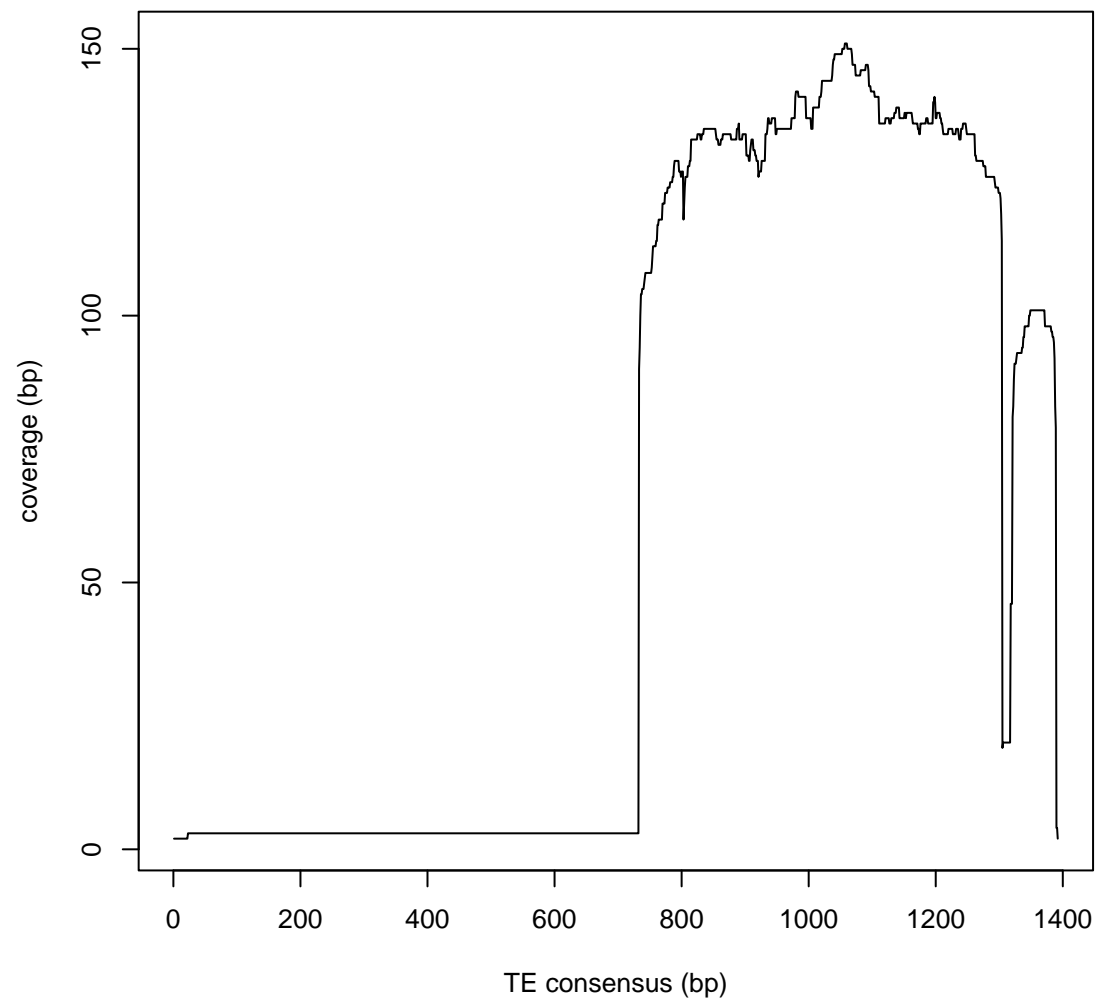


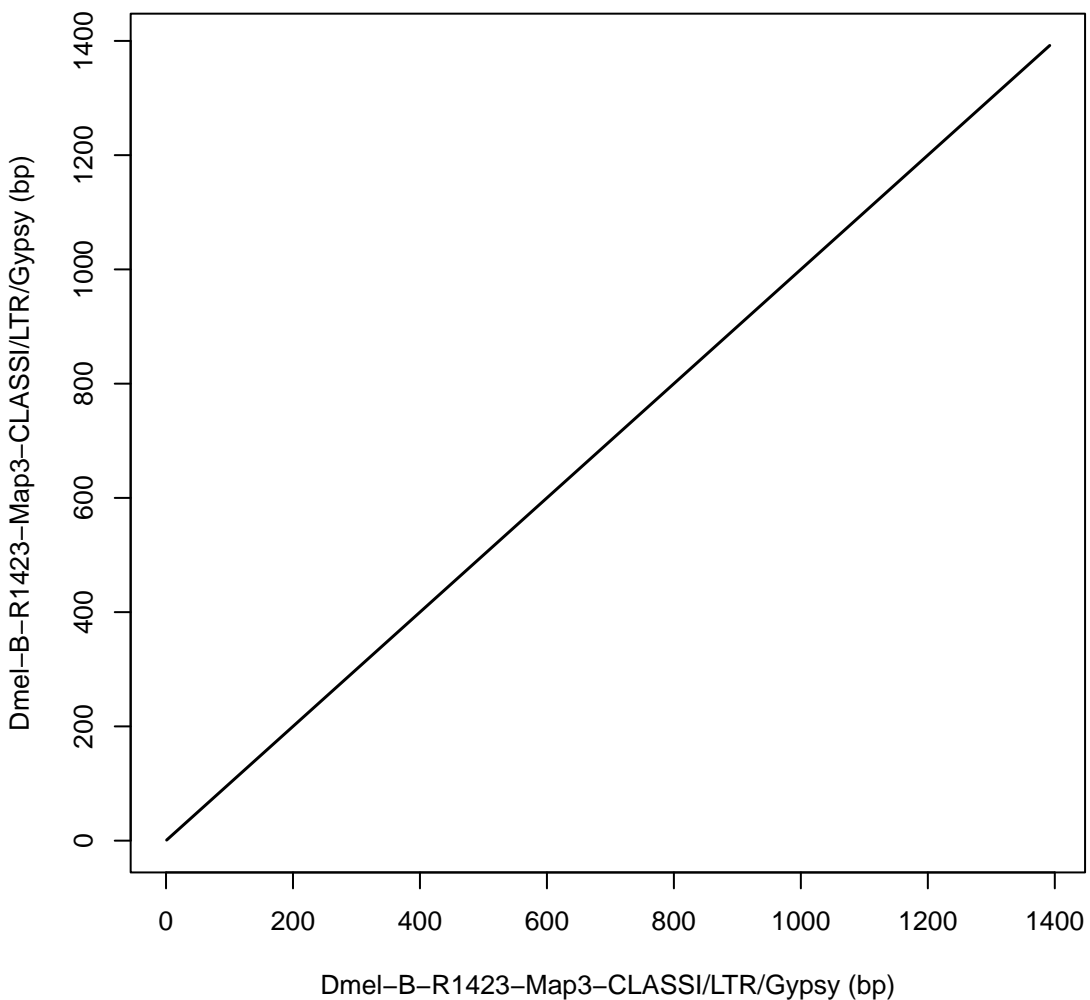
TE: Dmel-B-R1423-Map3-CLASSI/LTR/Gypsy
consensus size: 1392bp; fragments: 303; full length: 2 (≥ 1252.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

