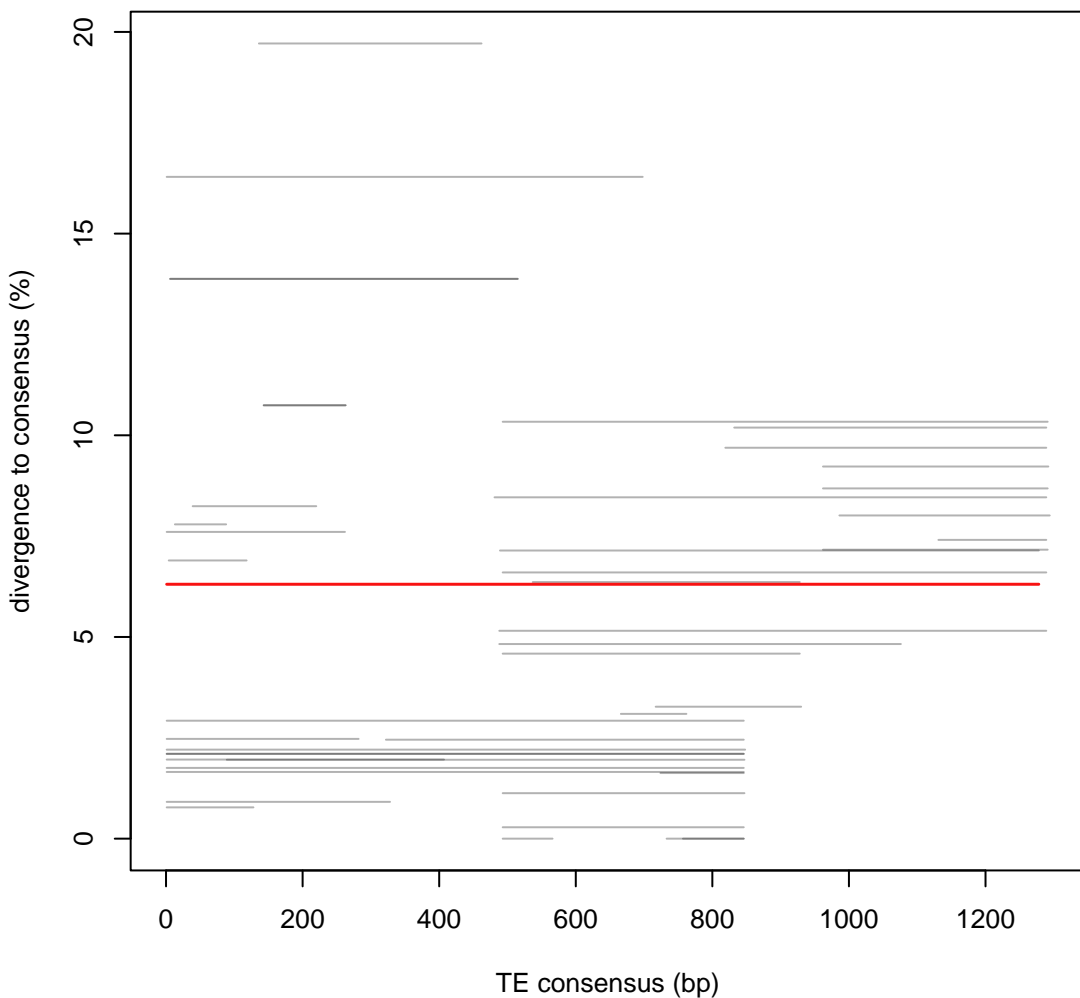
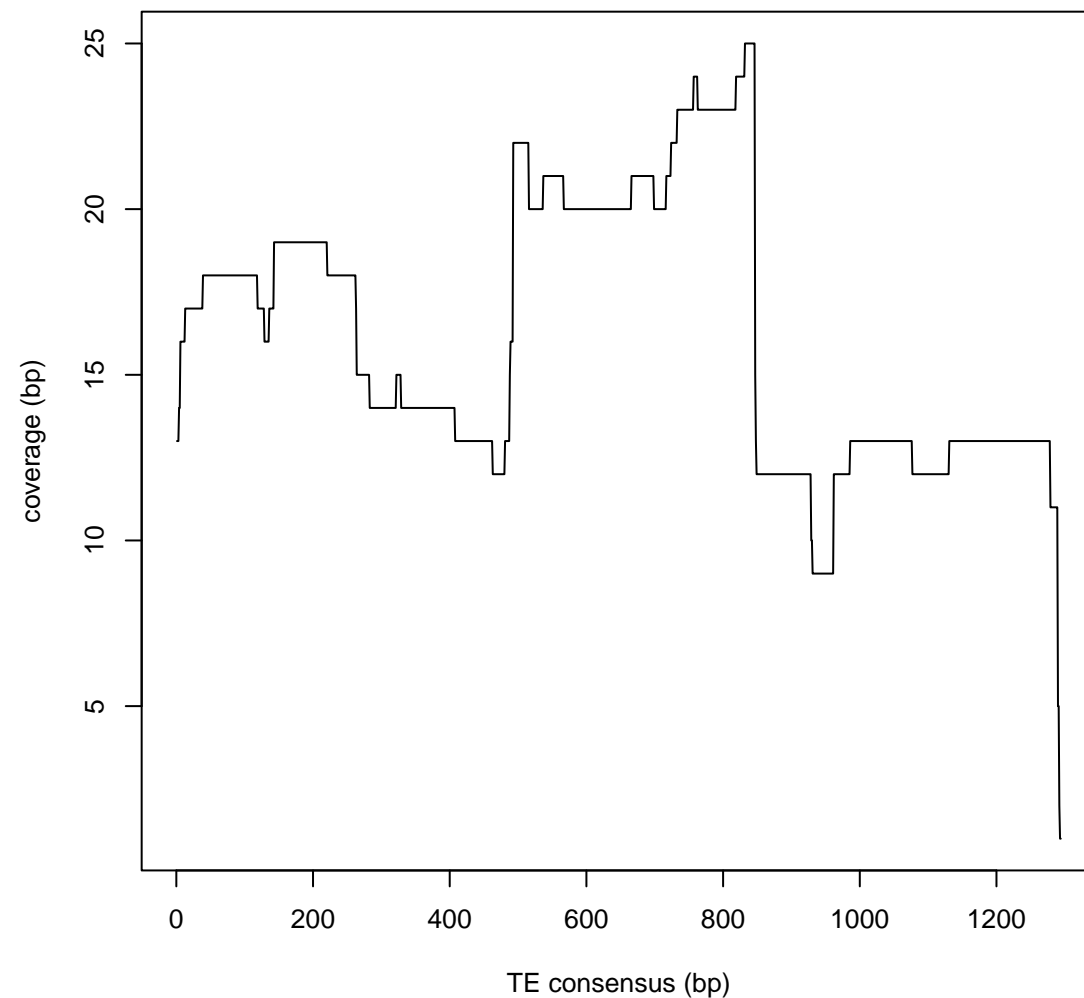


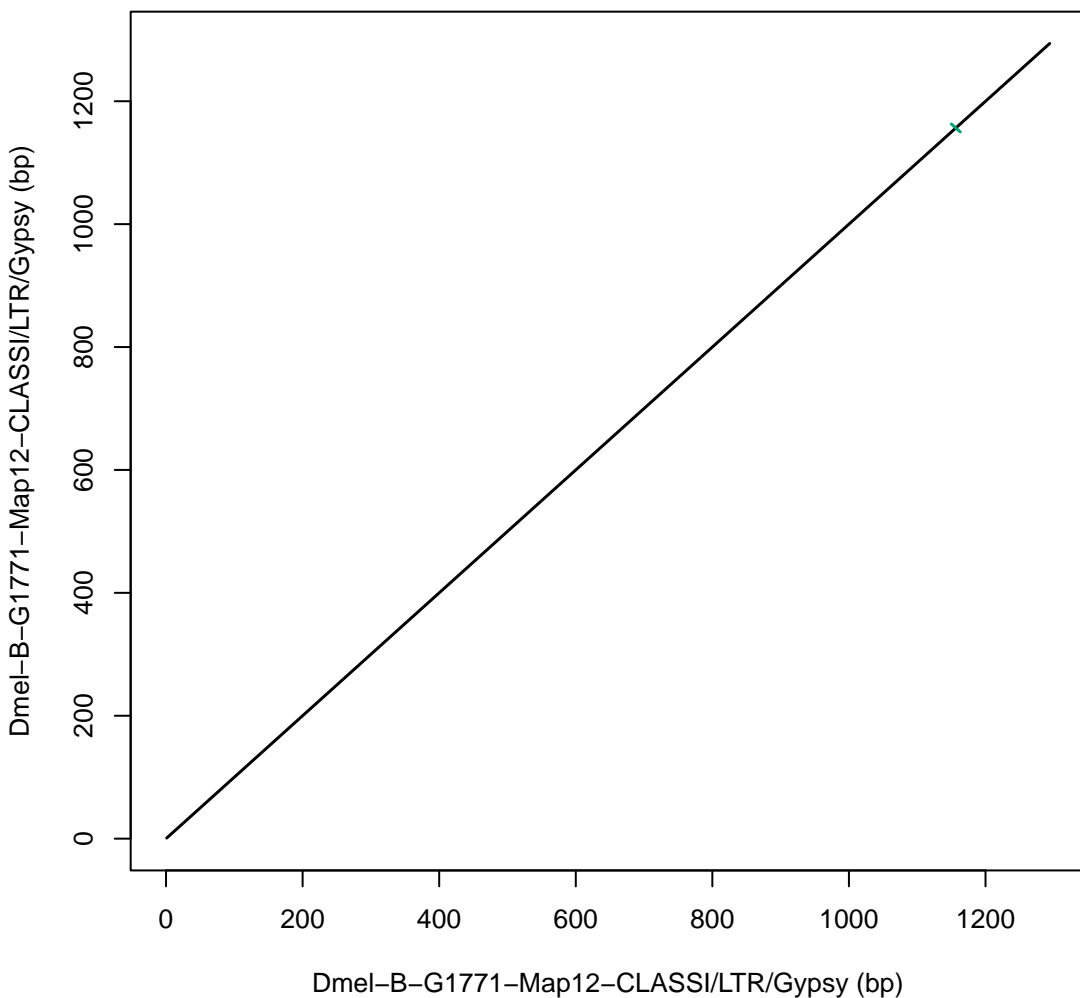
TE: Dmel-B-G1771-Map12-CLASSI/LTR/Gypsy
consensus size: 1294bp; fragments: 46; full length: 1 (>=1164.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

