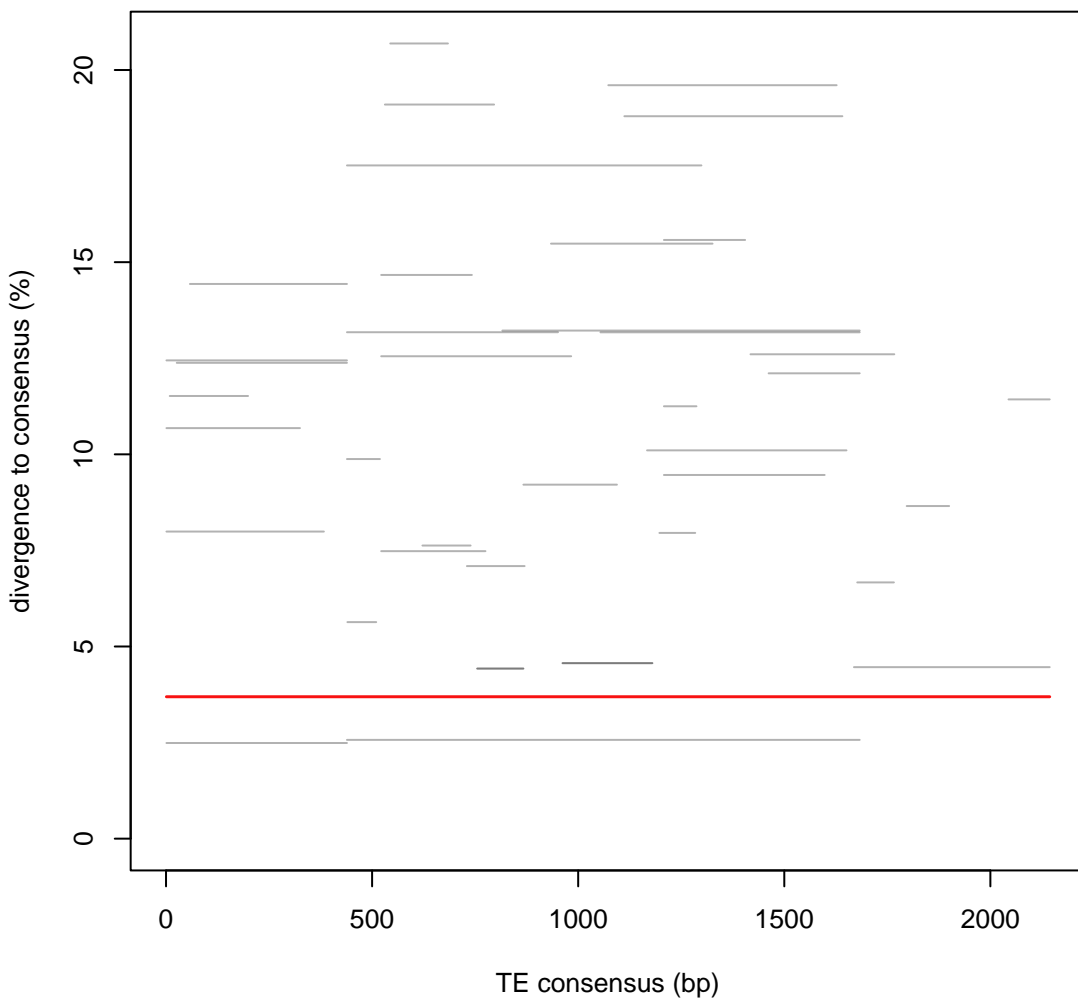
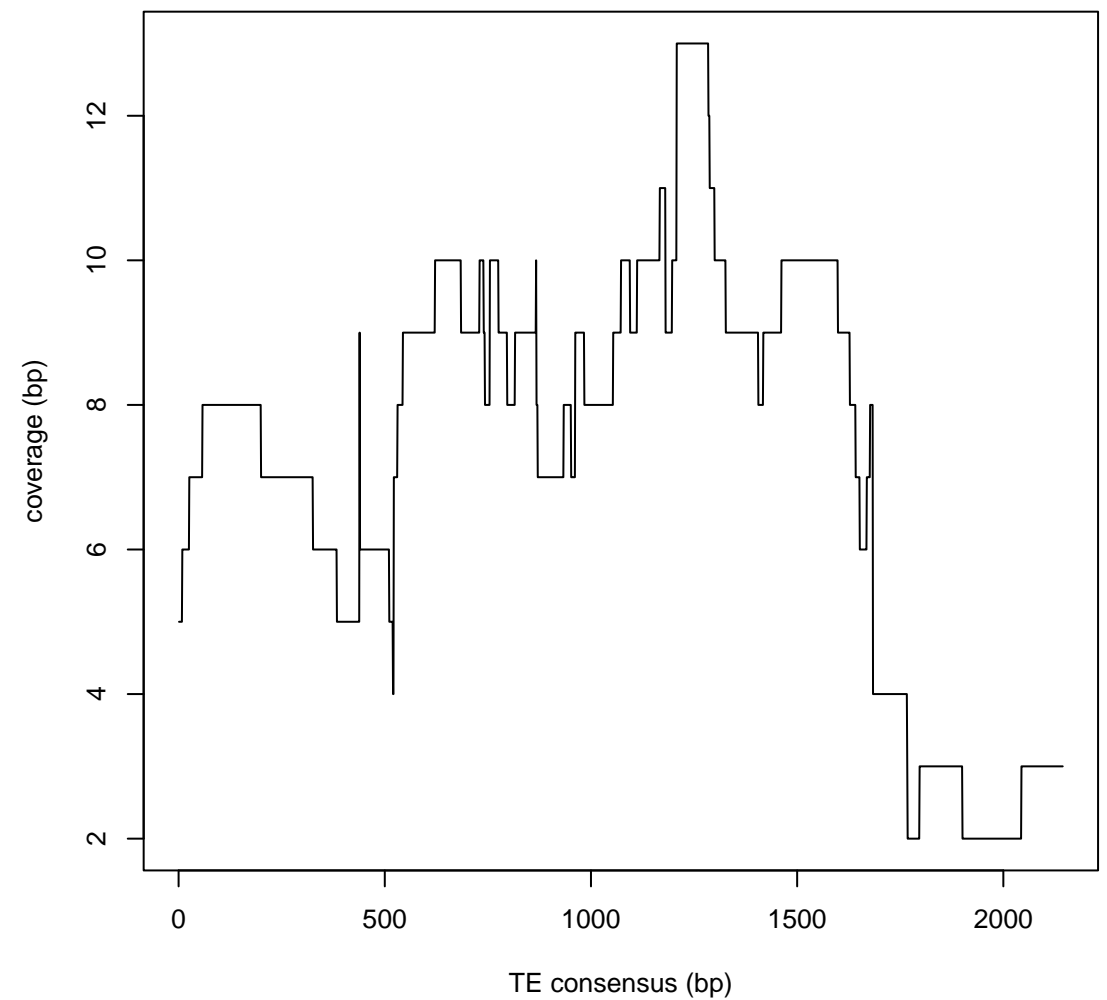


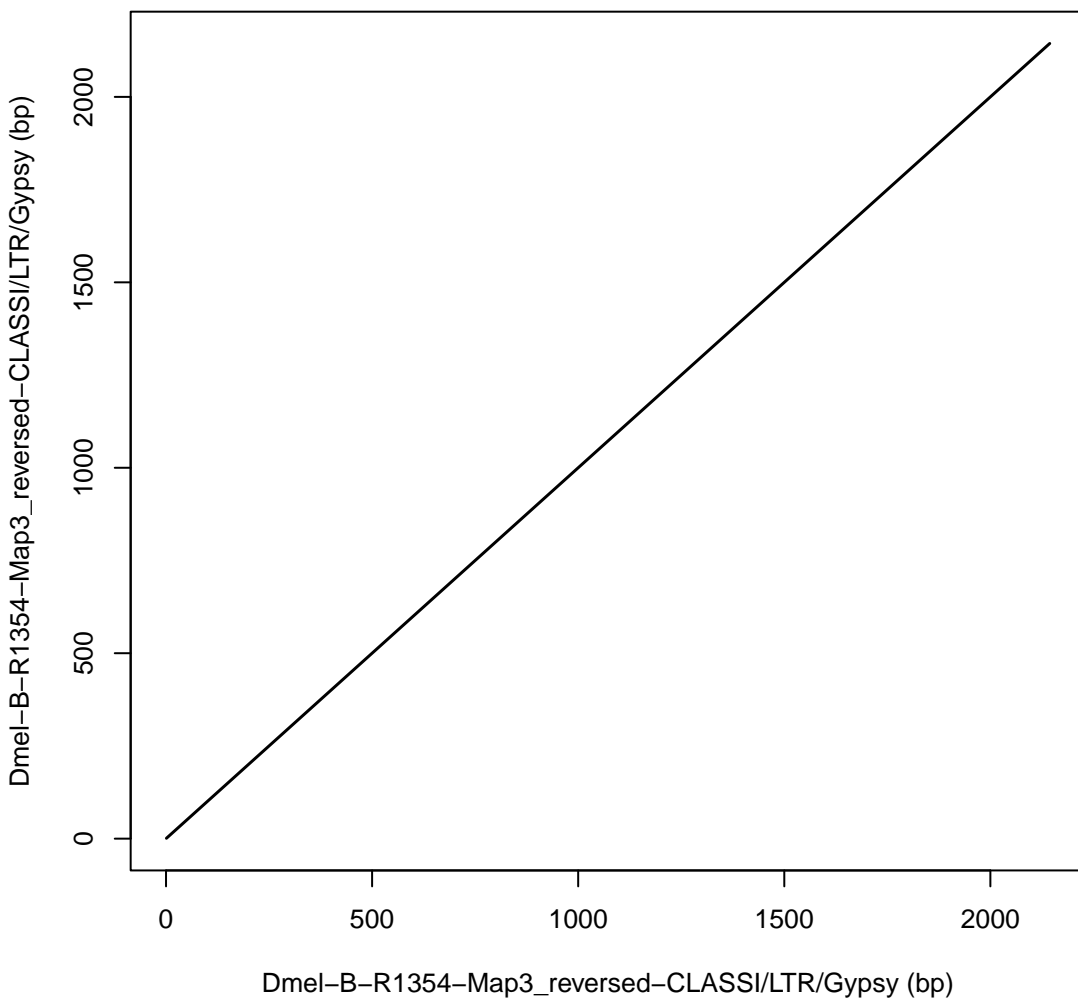
TE: Dmel-B-R1354-Map3_reversed-CLASSI/LTR/Gypsy
consensus size: 2144bp; fragments: 41; full length: 1 (≥ 1929.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

