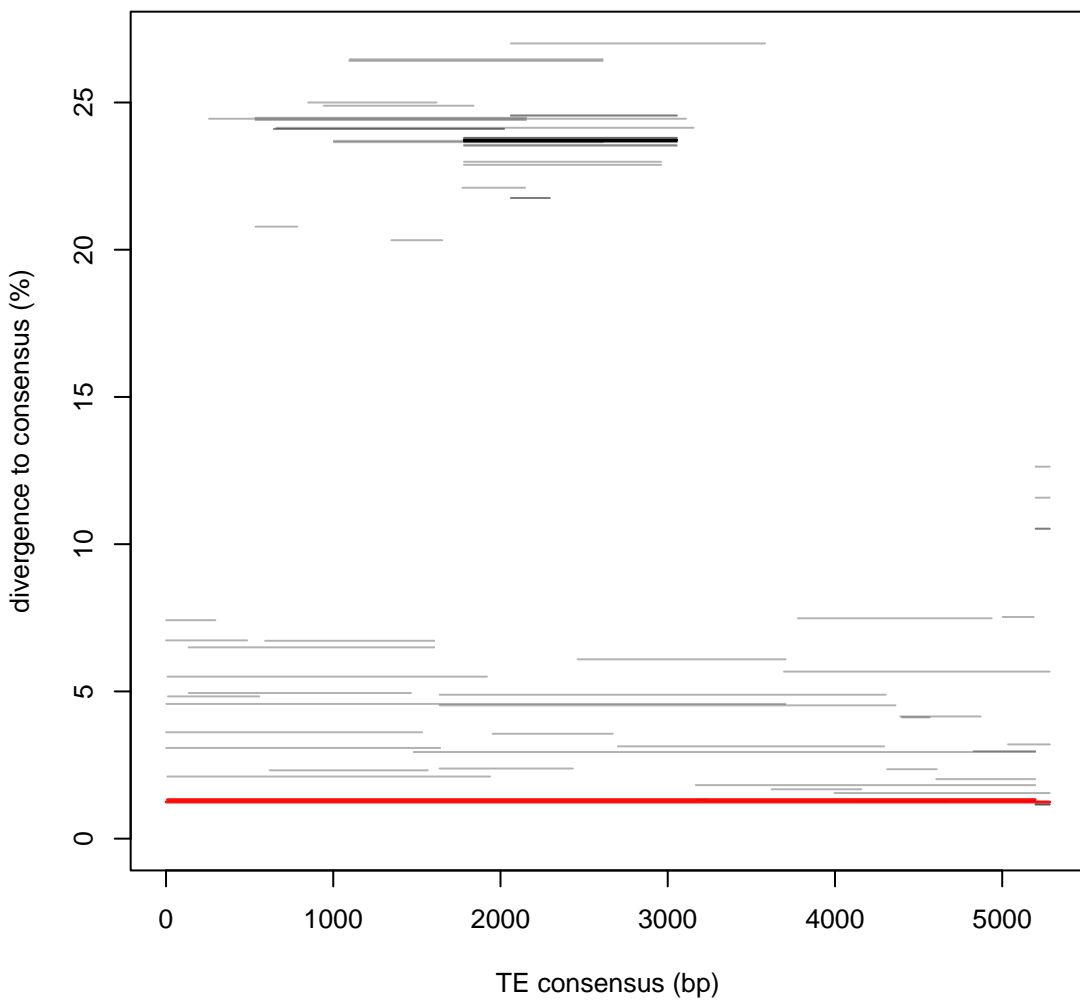
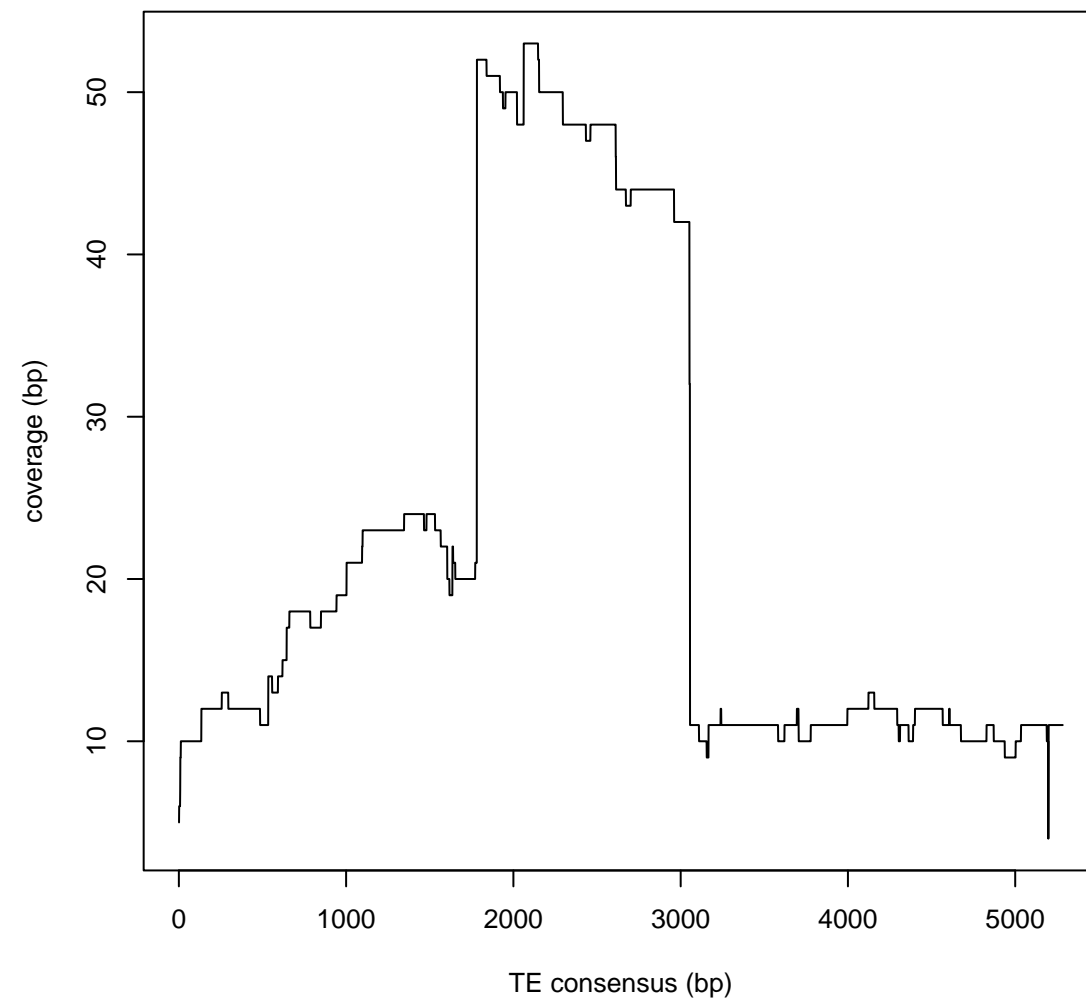


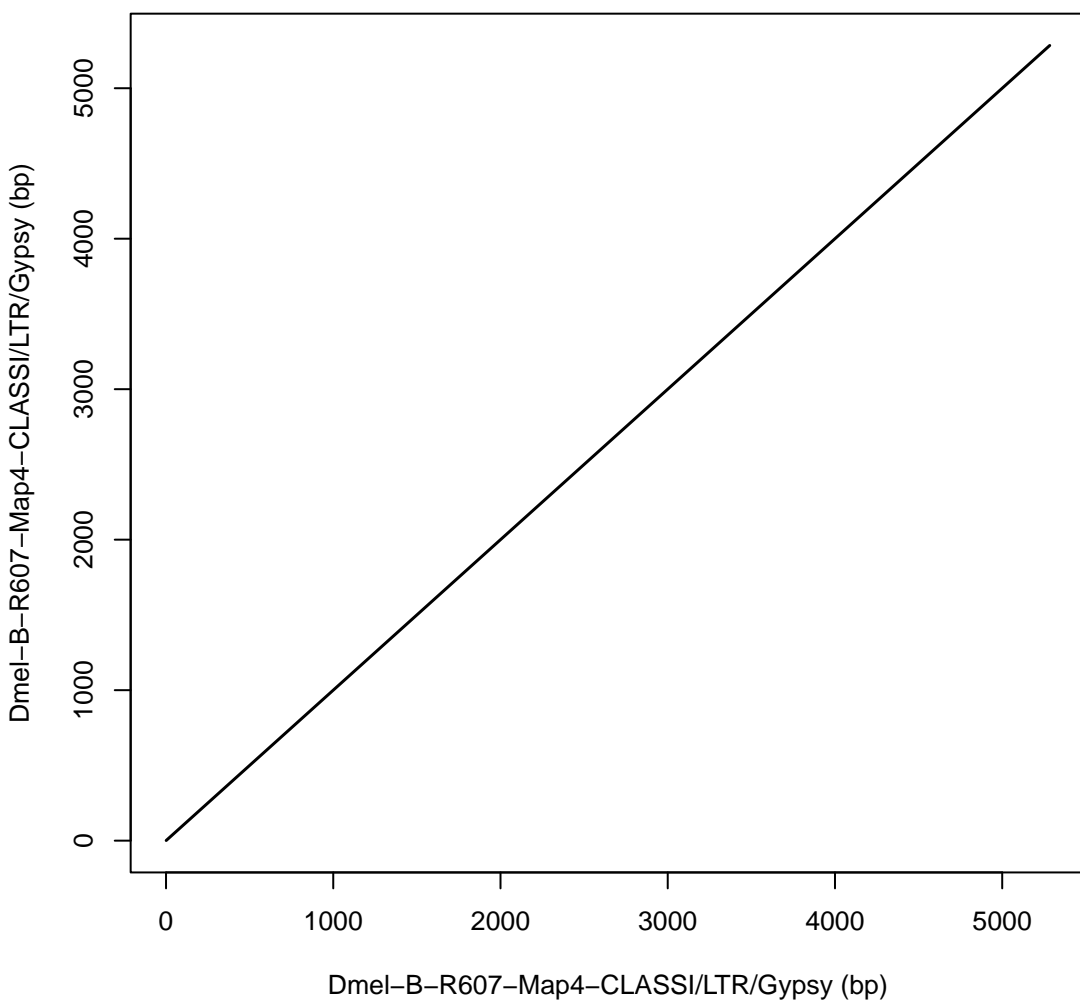
TE: Dmel-B-R607-Map4-CLASSI/LTR/Gypsy
consensus size: 5284bp; fragments: 94; full length: 2 (≥ 4755.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

