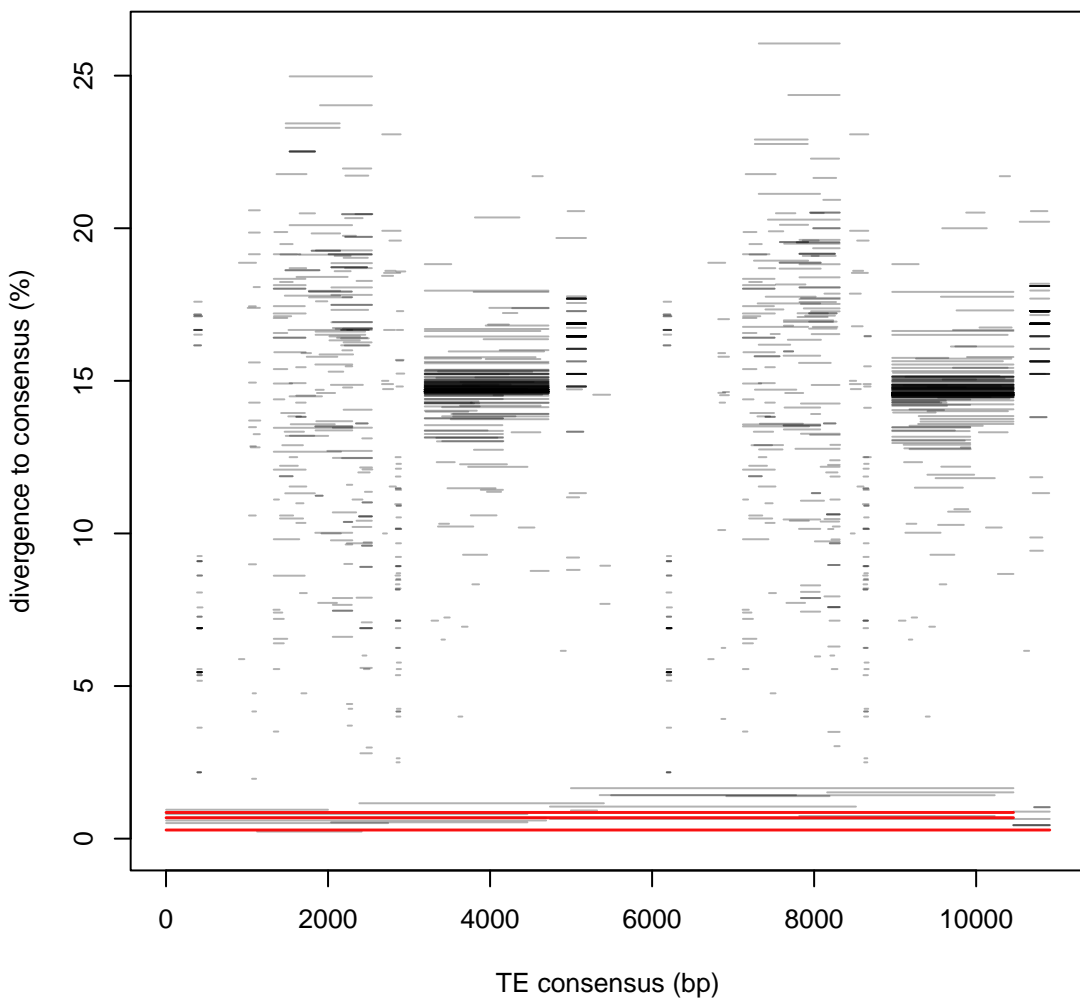
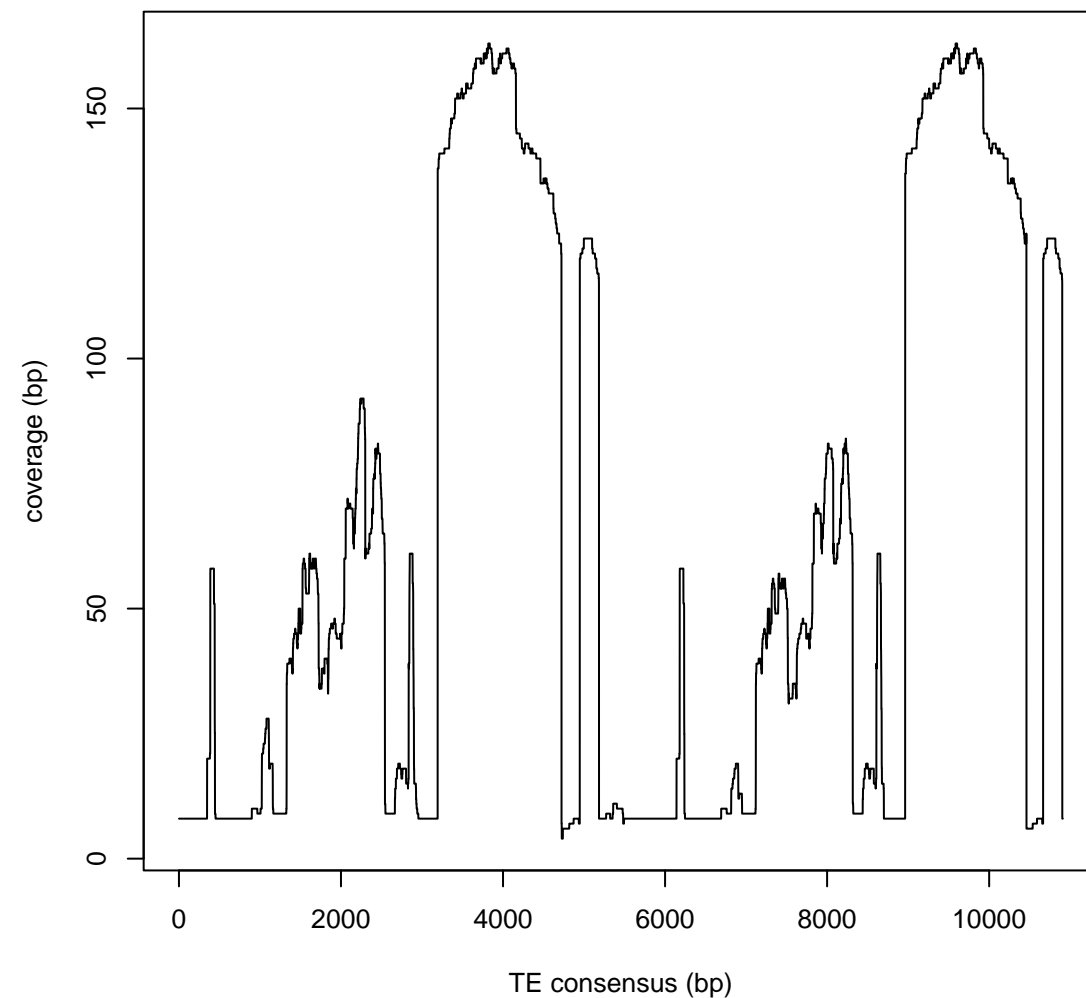


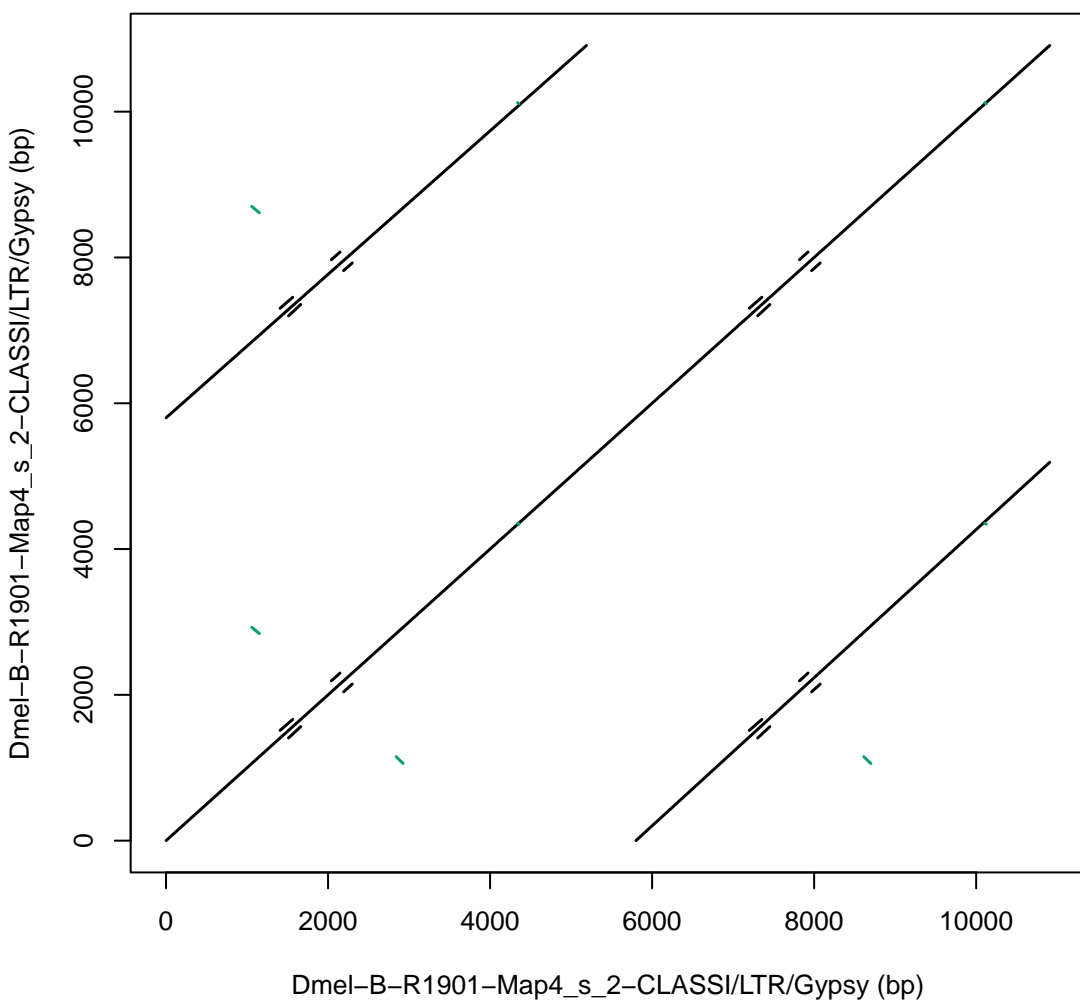
TE: Dmel-B-R1901-Map4_s_2-CLASSI/LTR/Gypsy
consensus size: 10908bp; fragments: 1267; full length: 3 (>=9817.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

