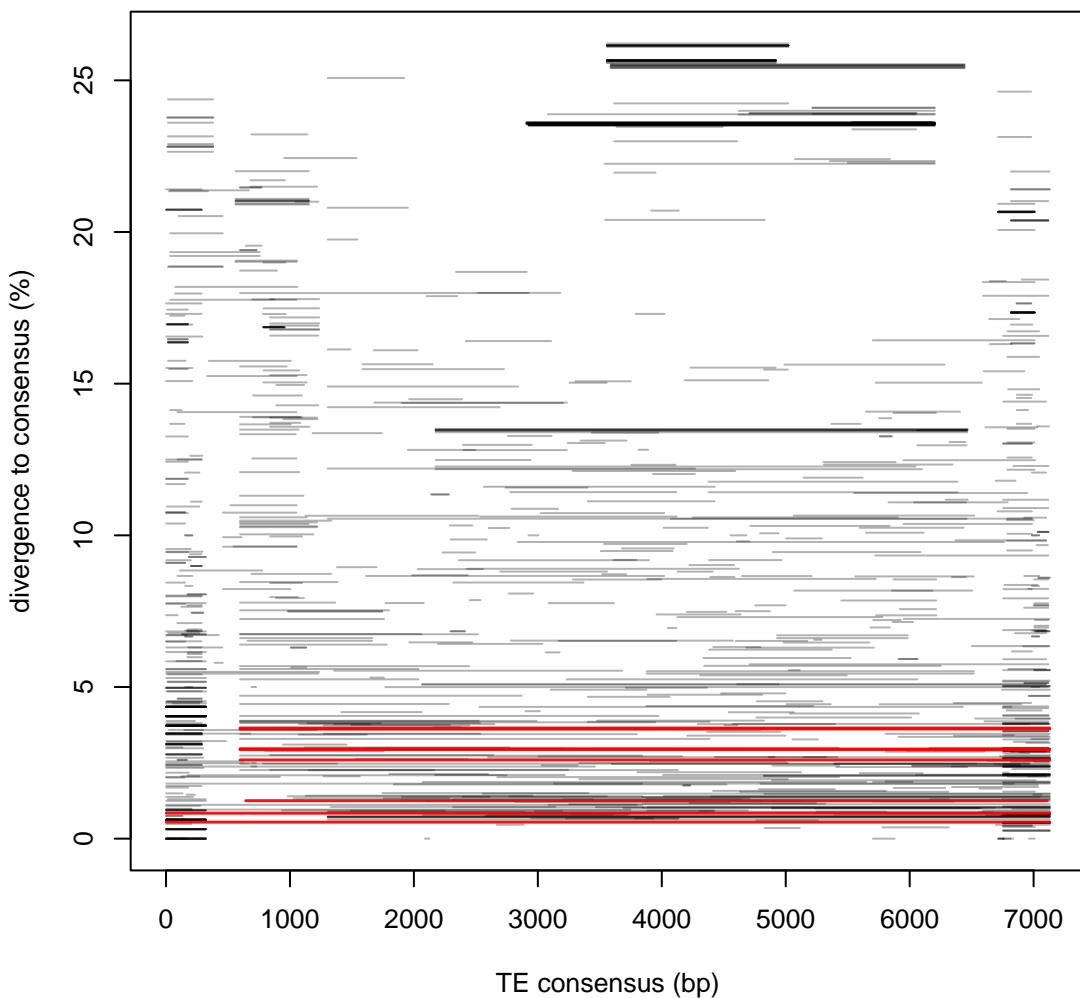
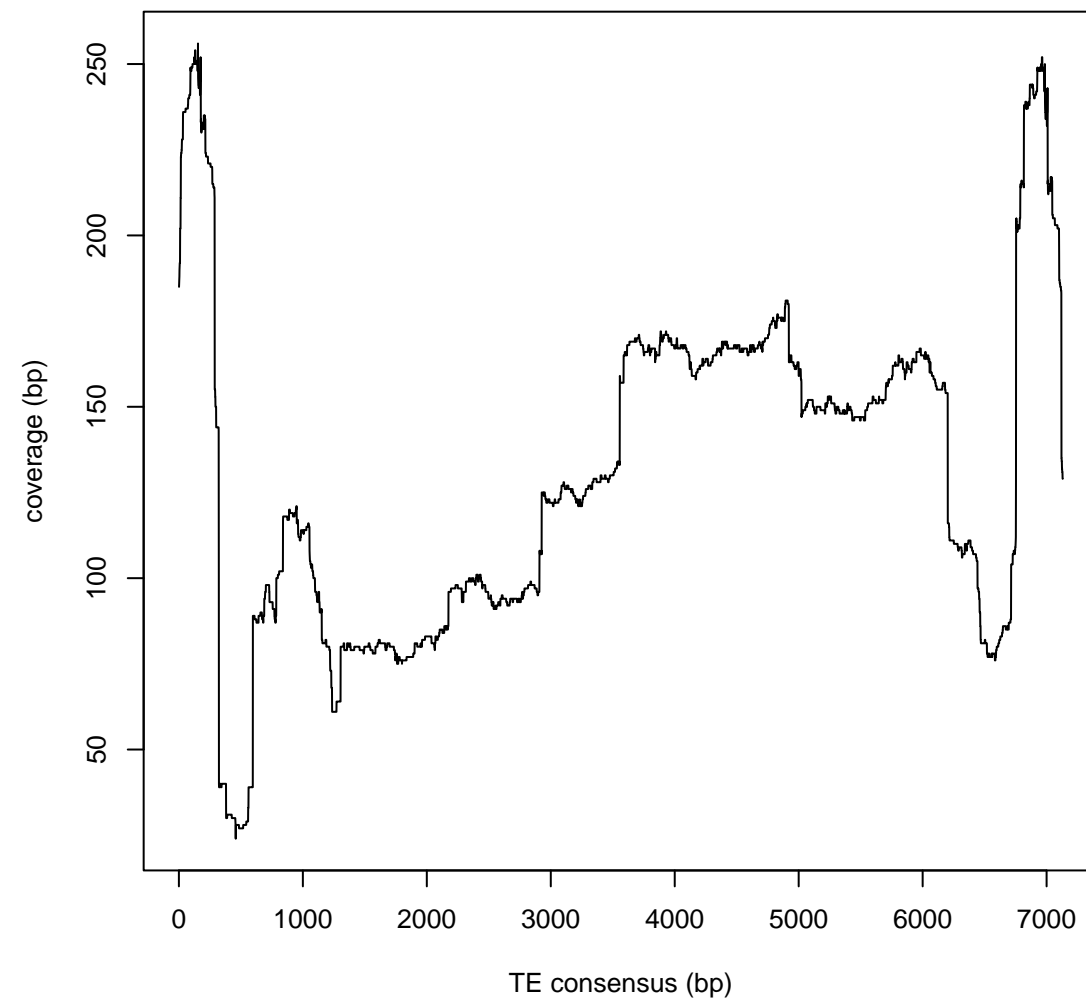


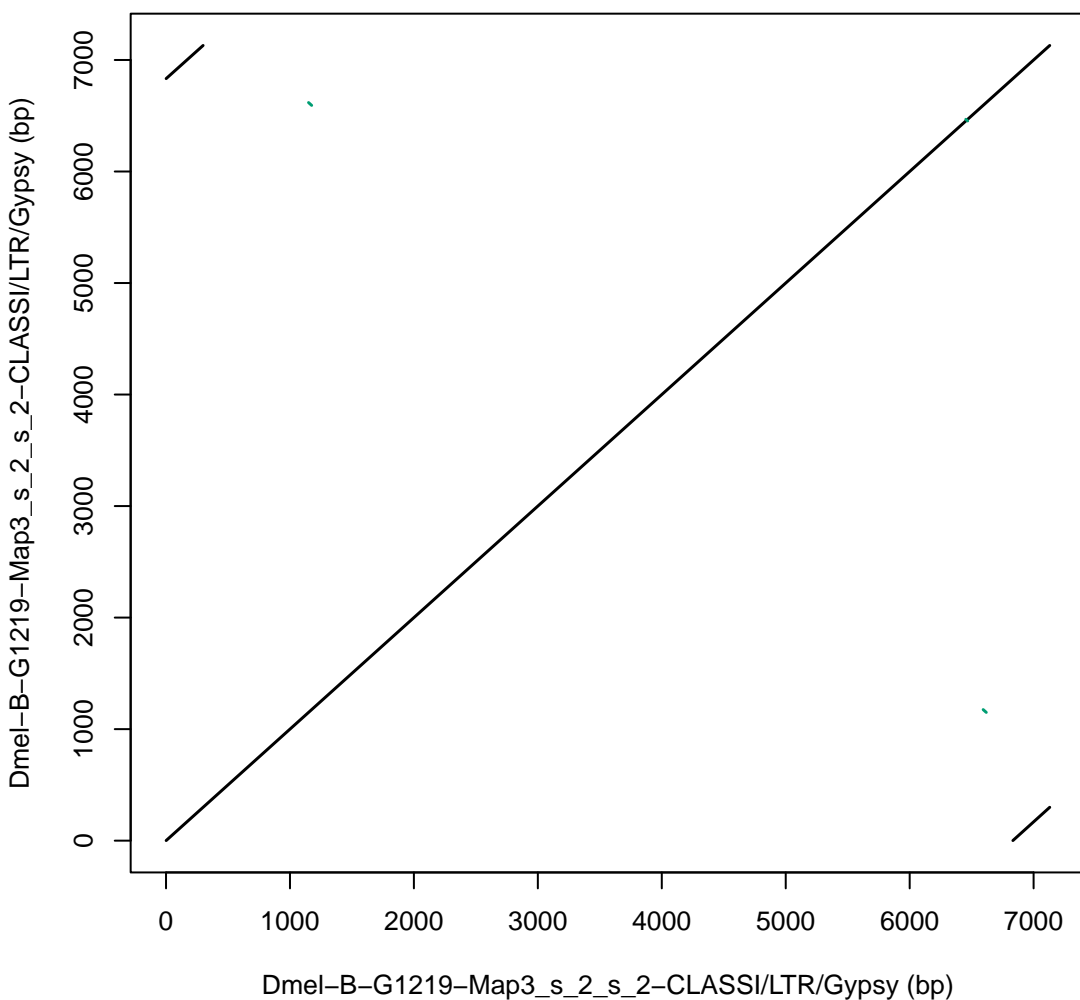
TE: Dmel-B-G1219-Map3_s_2_s_2-CLASSI/LTR/Gypsy
consensus size: 7130bp; fragments: 1063; full length: 8 (>=6417bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

