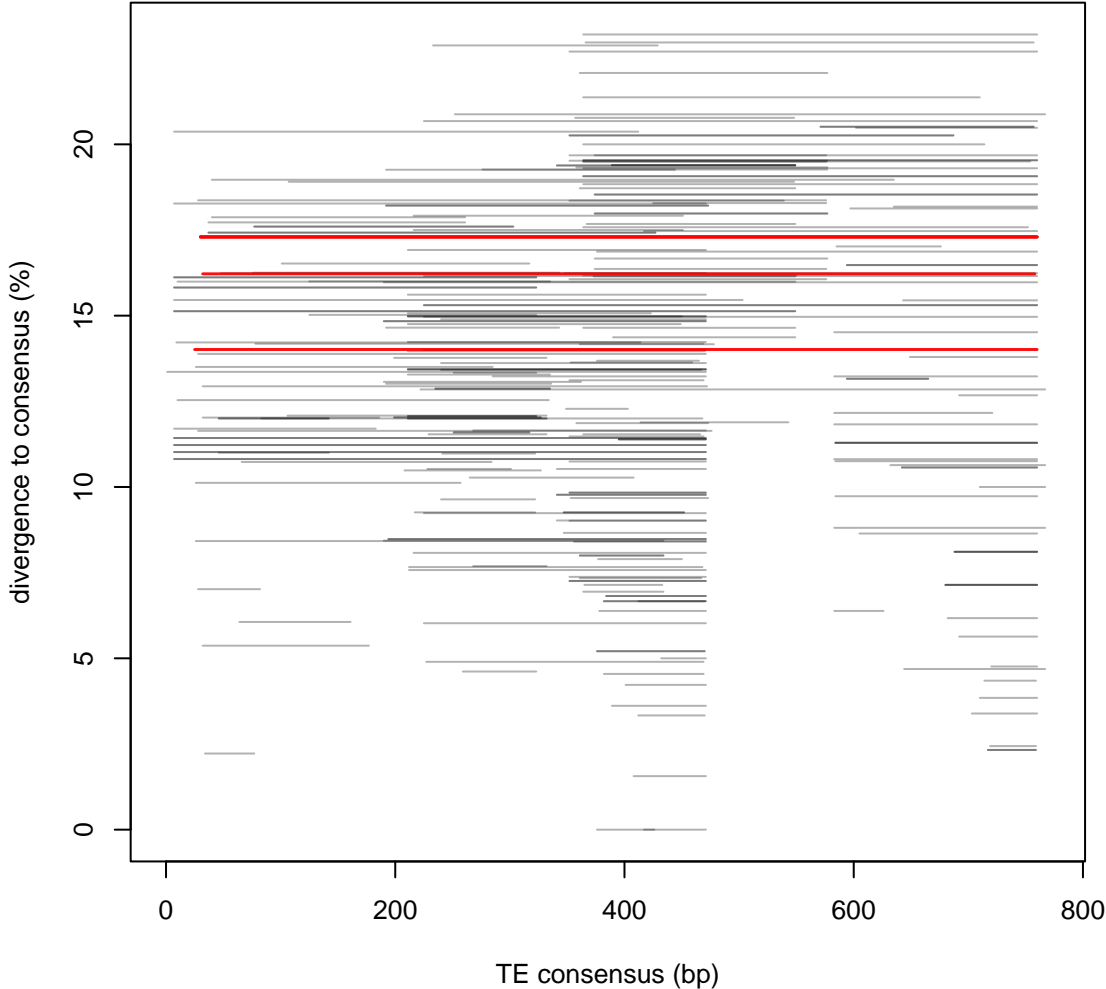
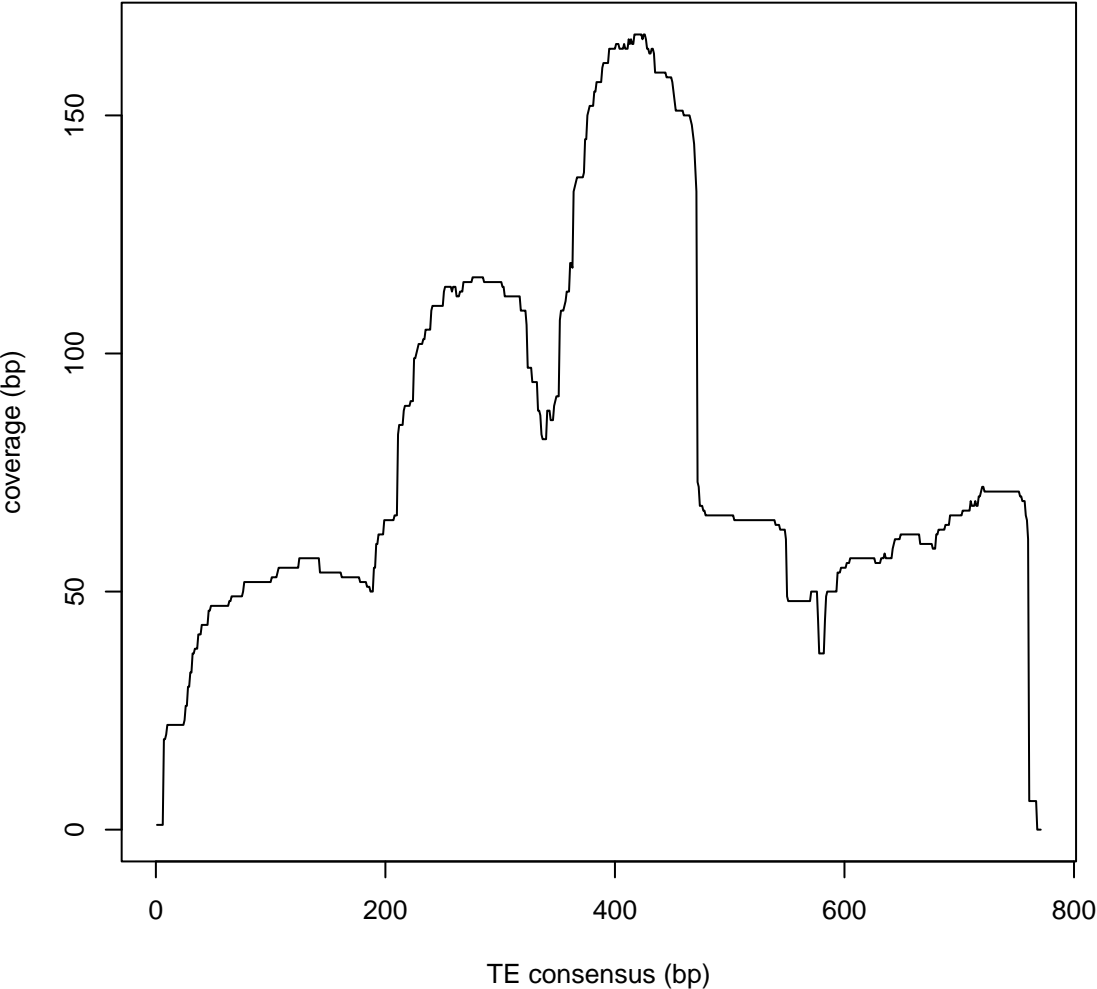


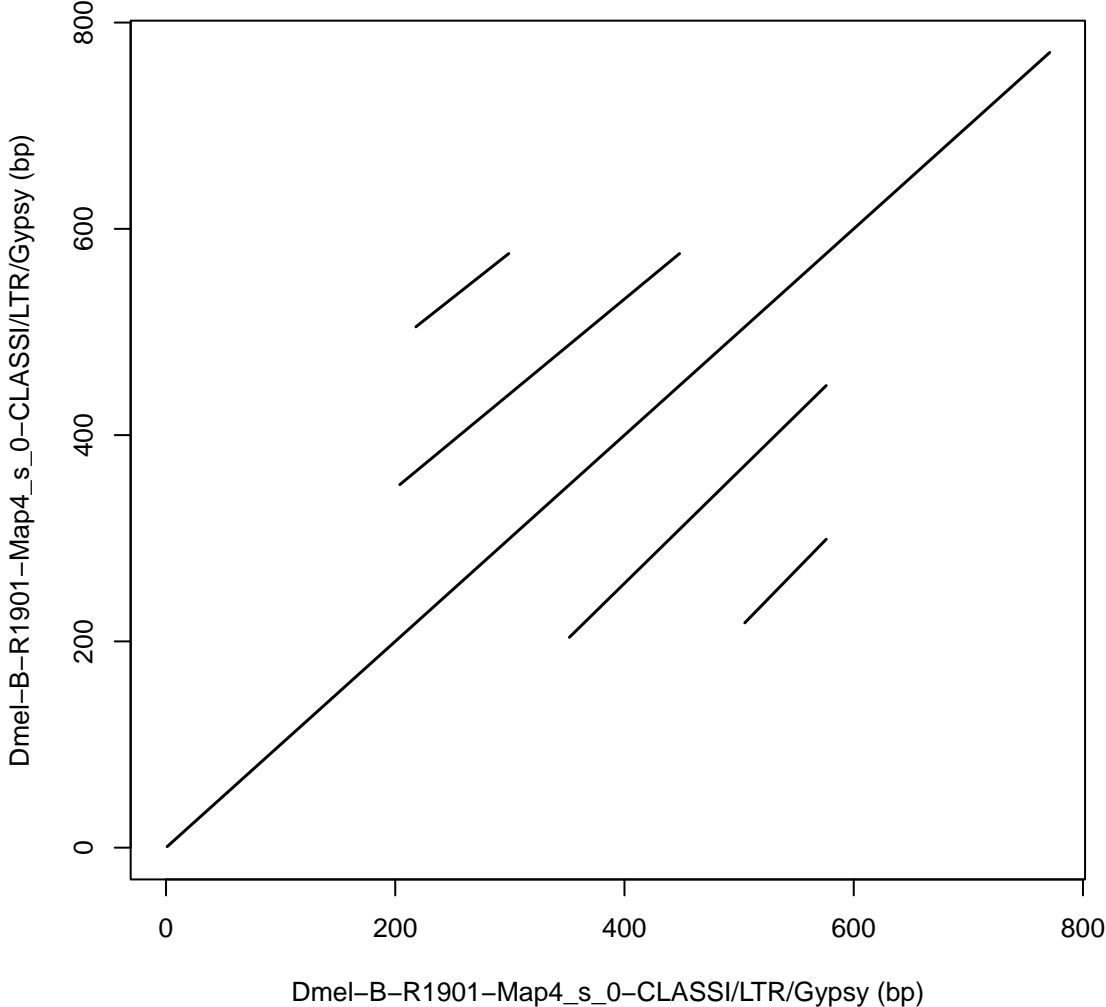
TE: Dmel-B-R1901-Map4_s_0-CLASSI/LTR/Gypsy
consensus size: 771bp; fragments: 269; full length: 5 (>=693.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

