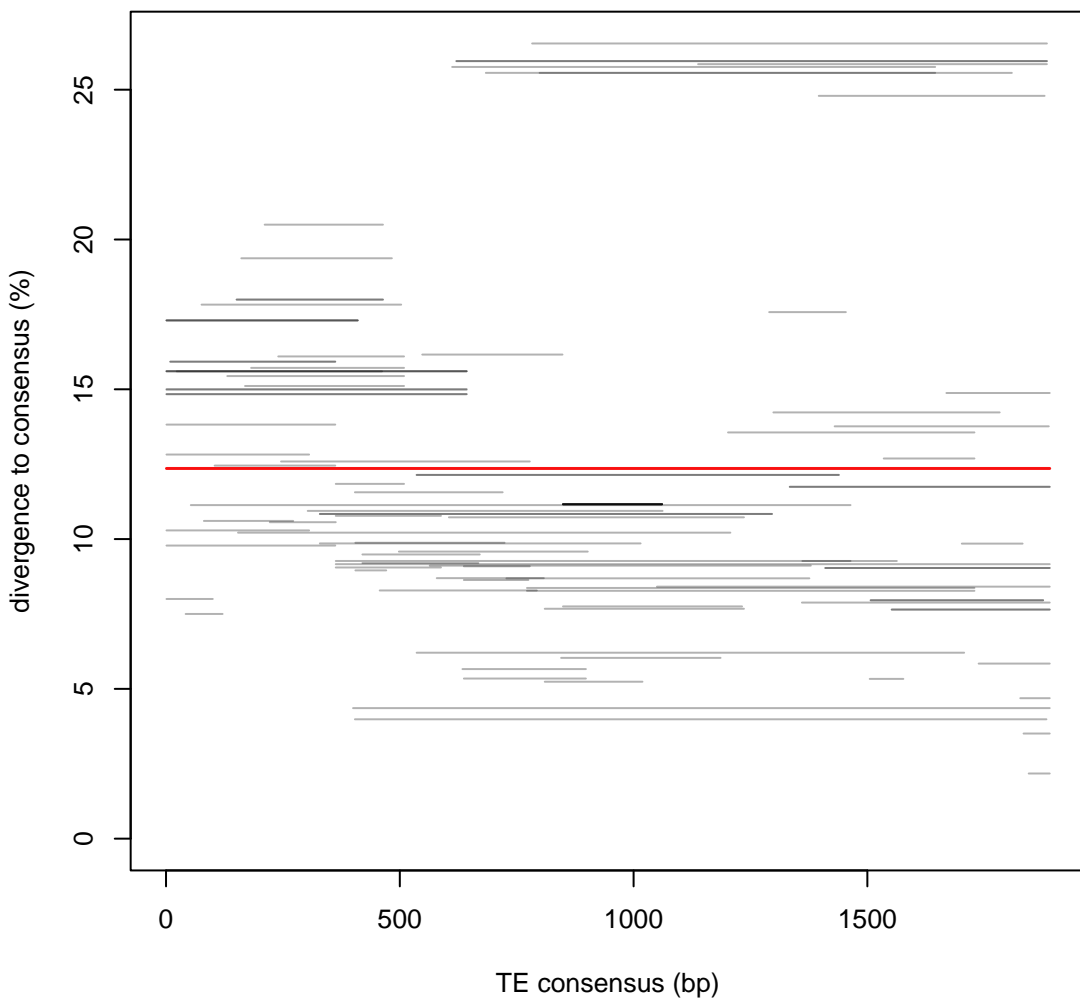
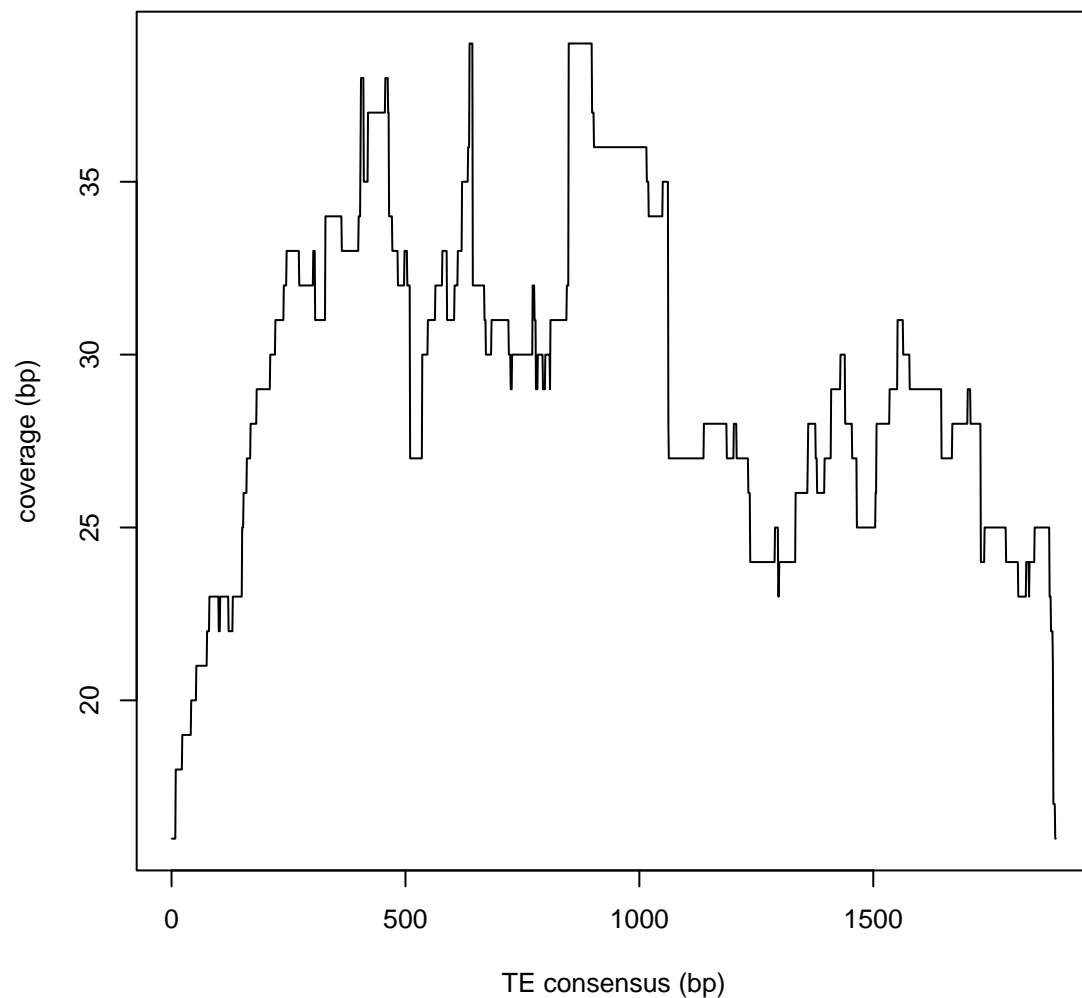


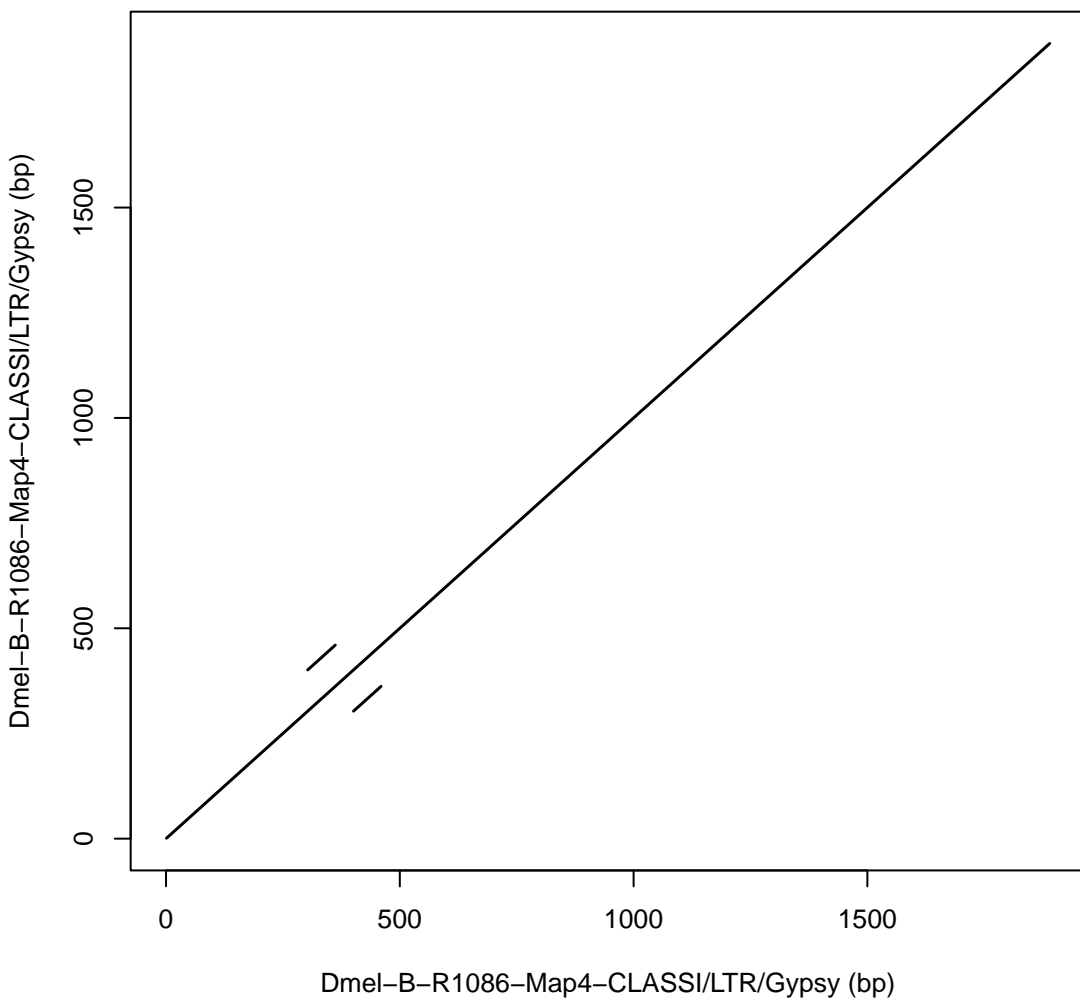
TE: Dmel-B-R1086-Map4-CLASSI/LTR/Gypsy
consensus size: 1890bp; fragments: 109; full length: 1 (>=1701bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

