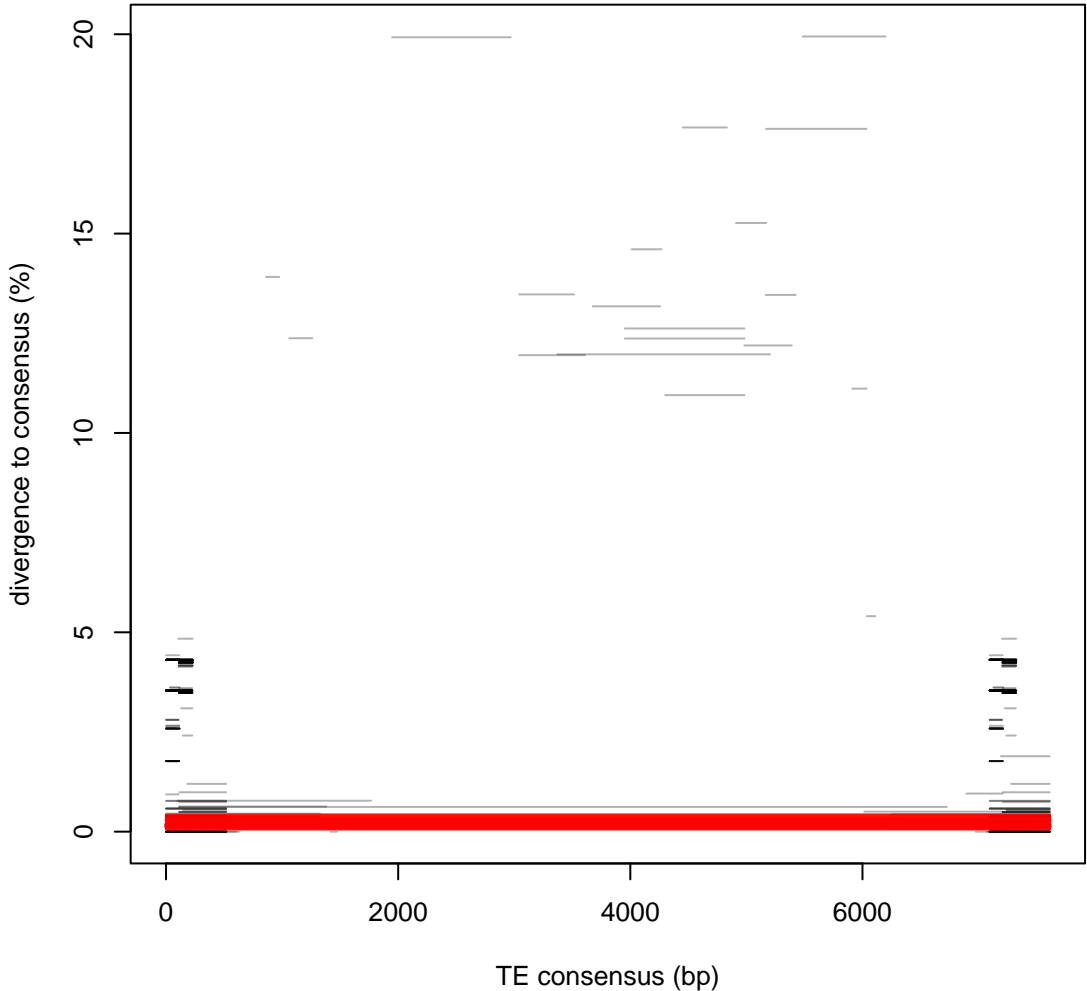
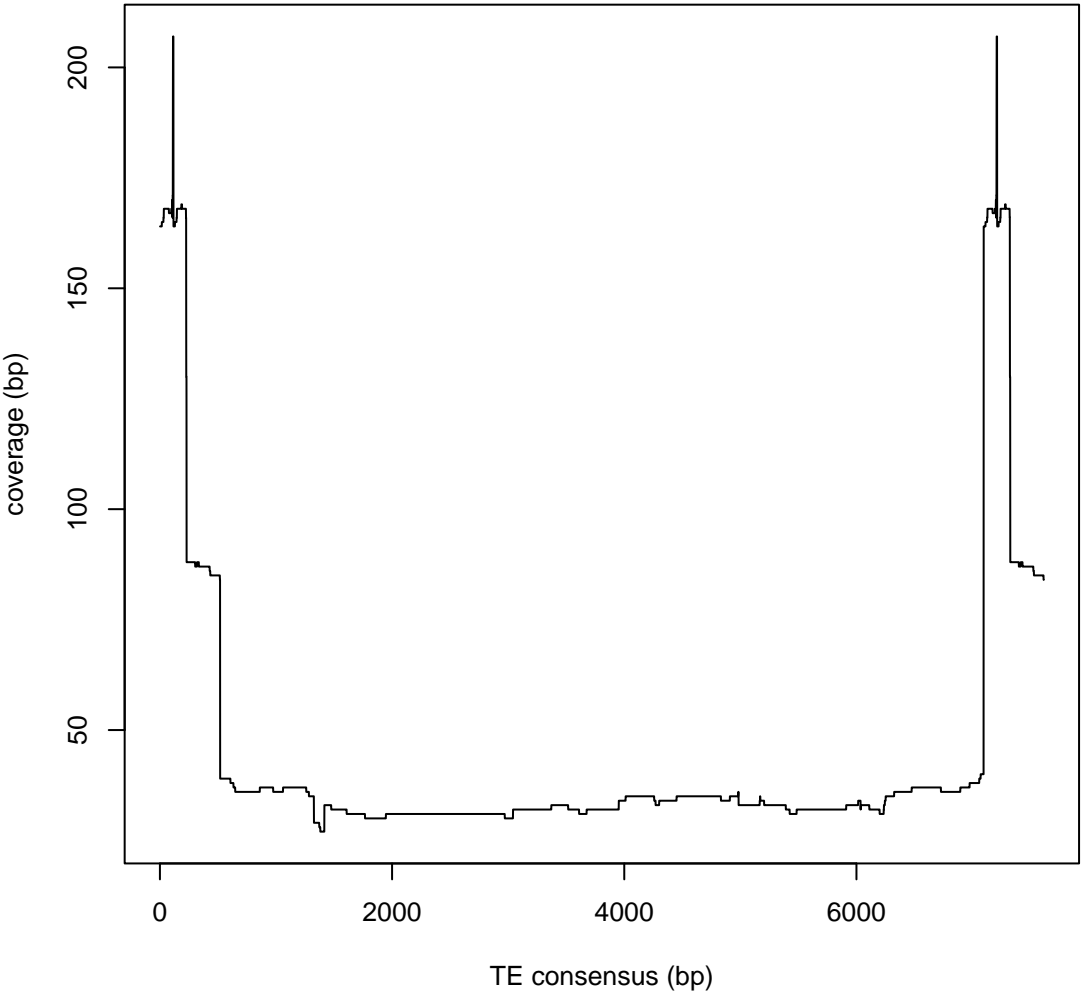


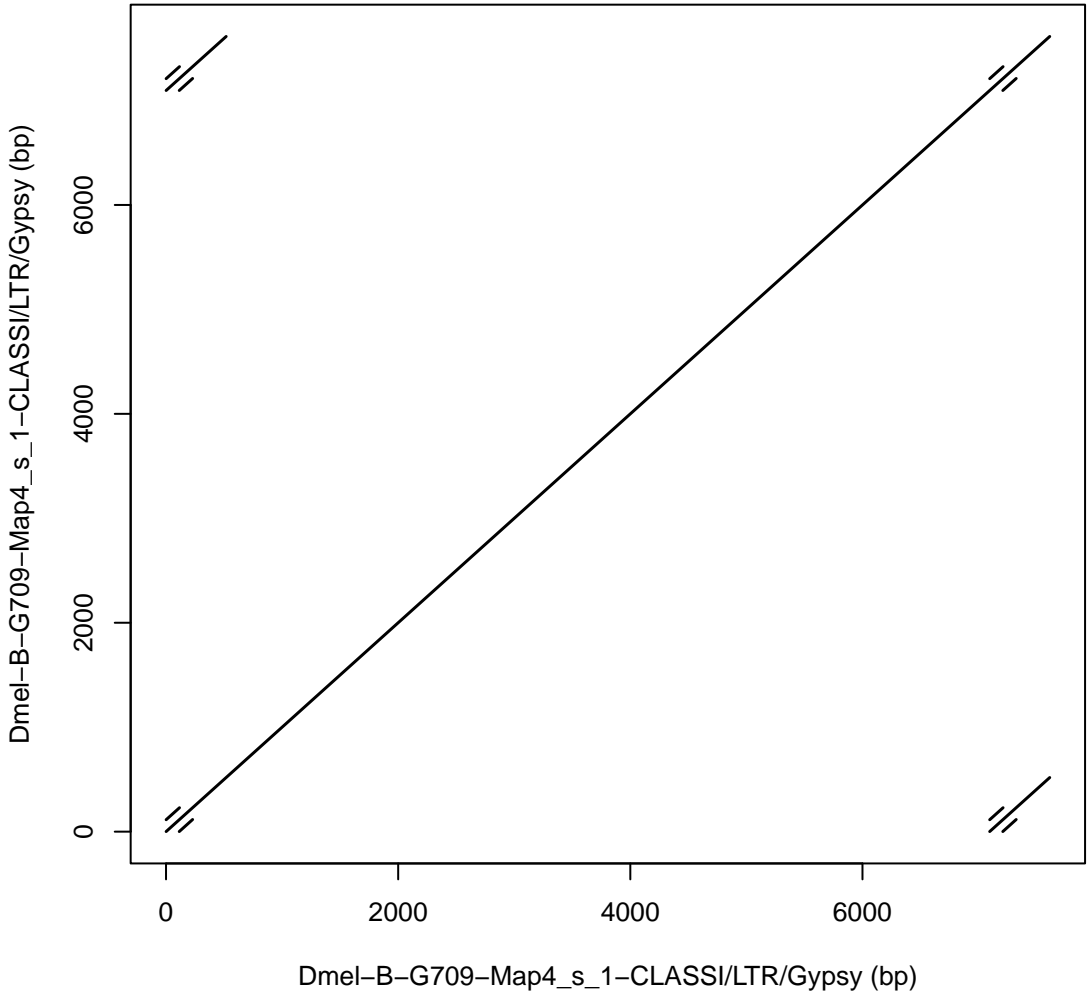
TE: Dmel-B-G709-Map4_s_1-CLASSI/LTR/Gypsy
consensus size: 7613bp; fragments: 516; full length: 24 (>=6851.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

