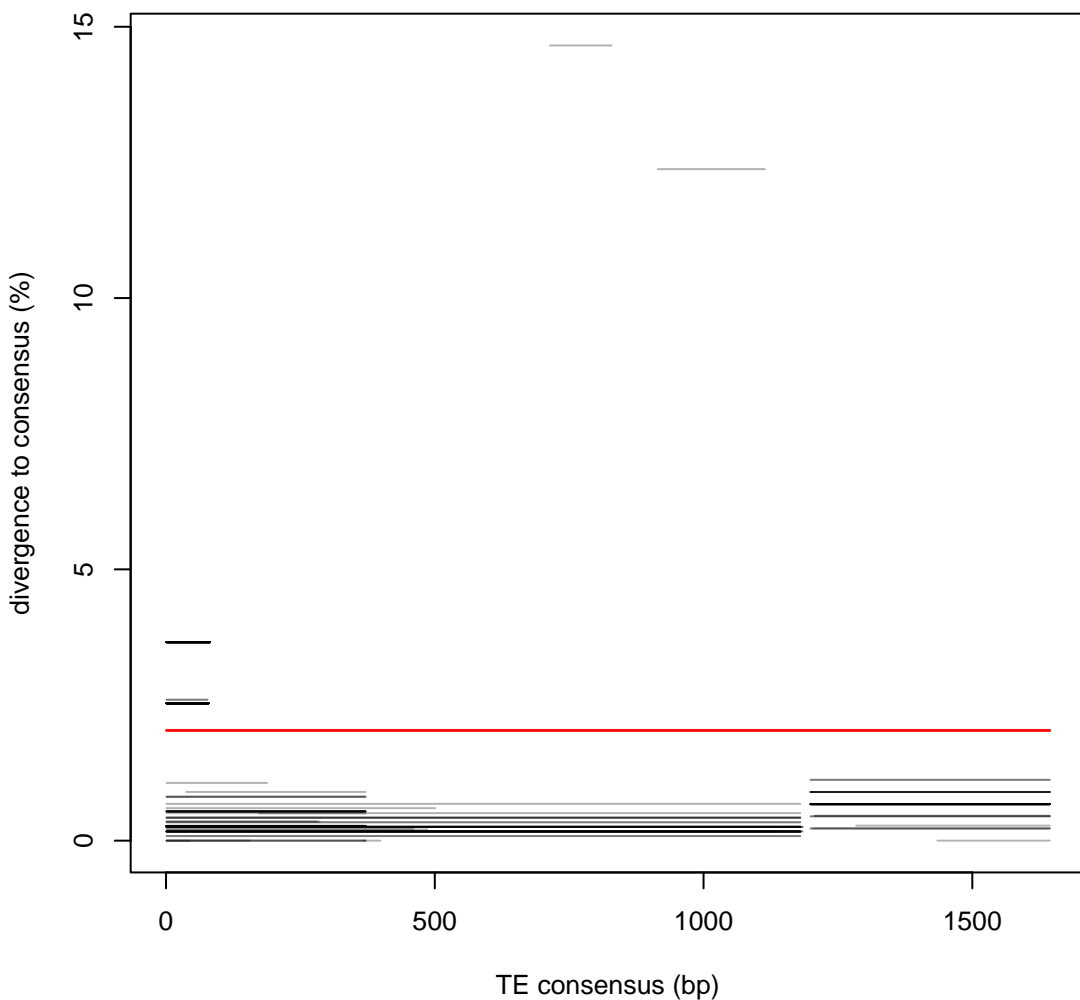
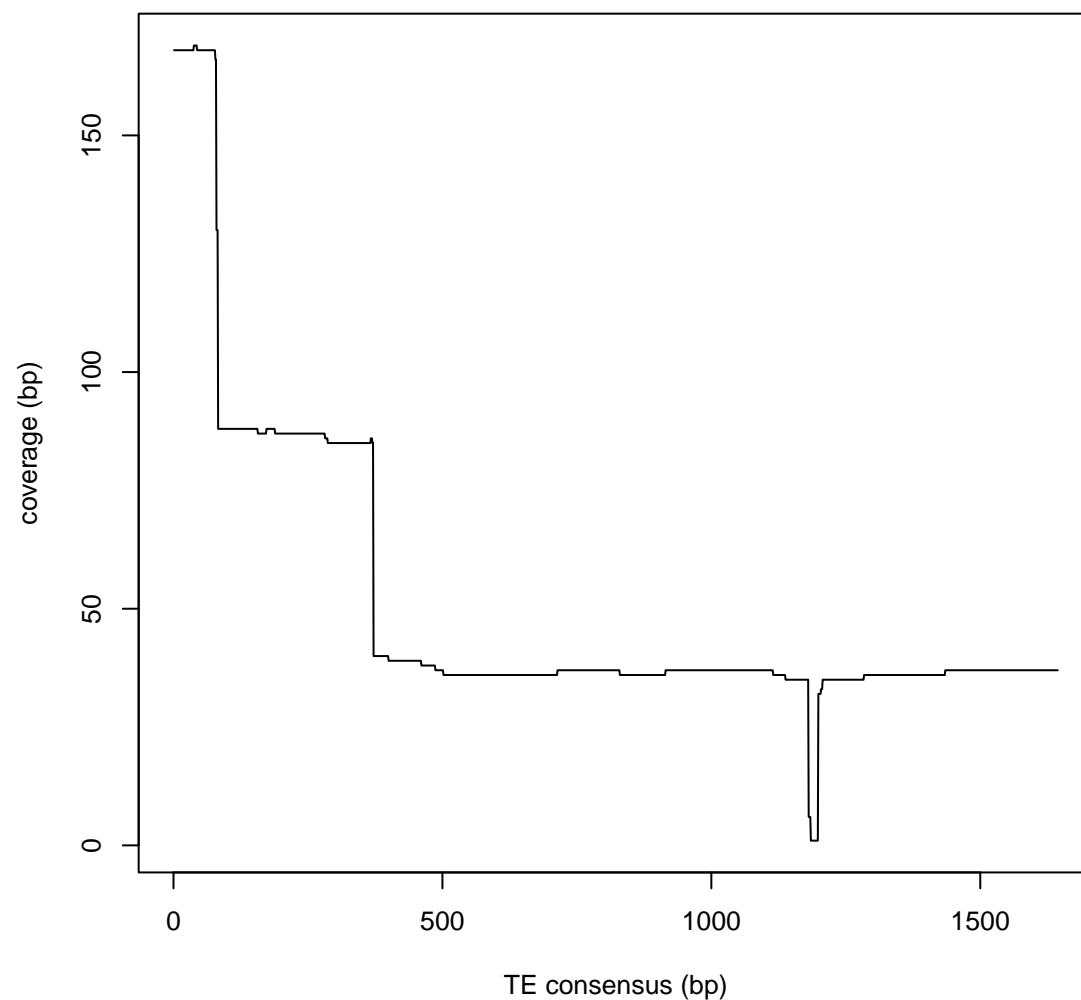


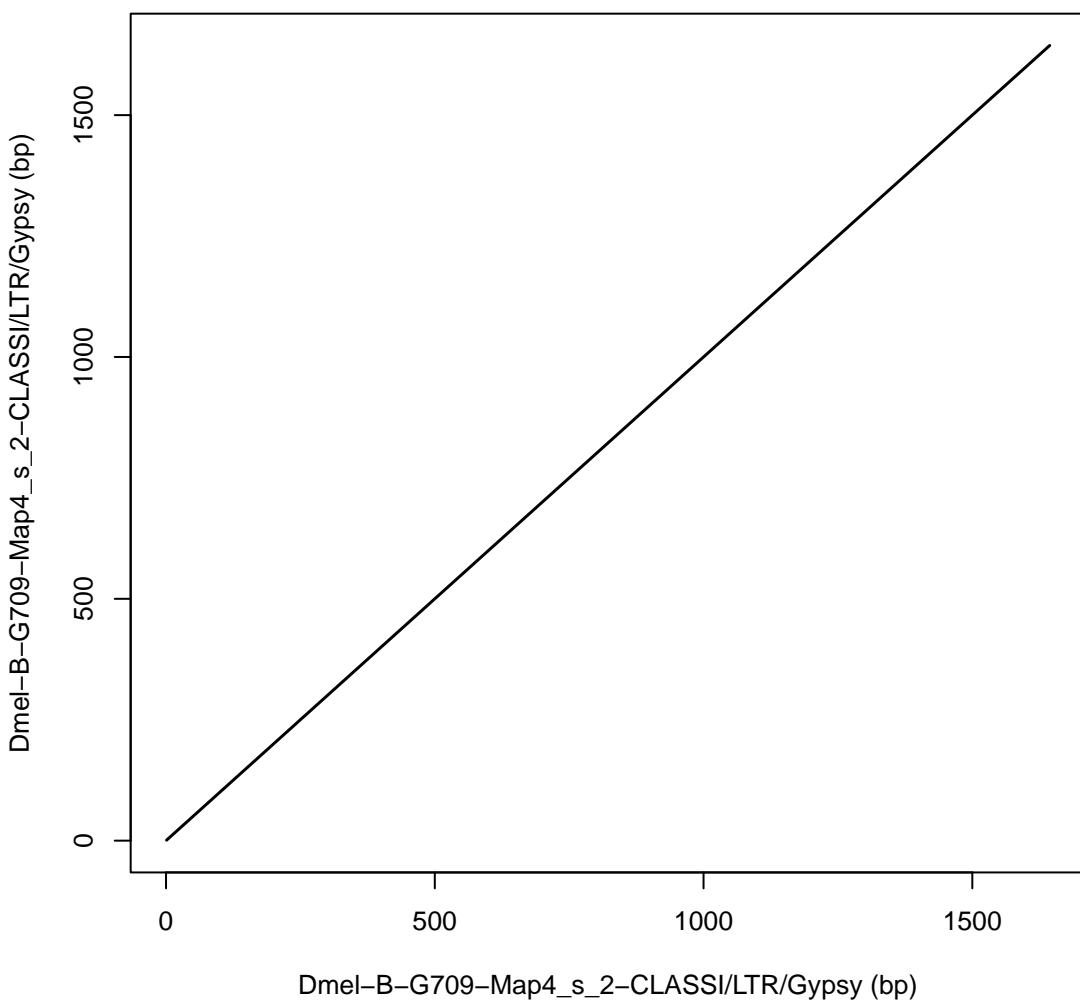
TE: Dmel-B-G709-Map4_s_2-CLASSI/LTR/Gypsy
consensus size: 1644bp; fragments: 209; full length: 1 (≥ 1479.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

