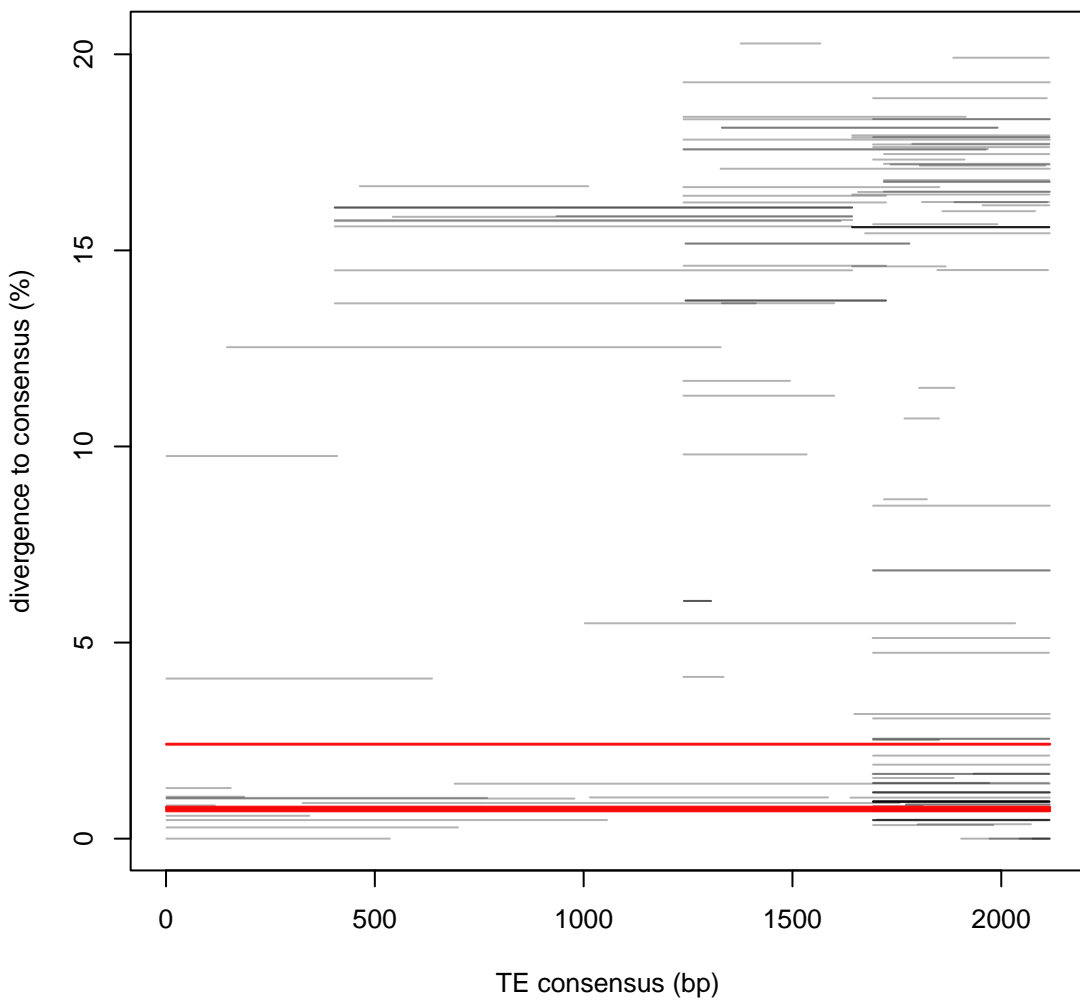
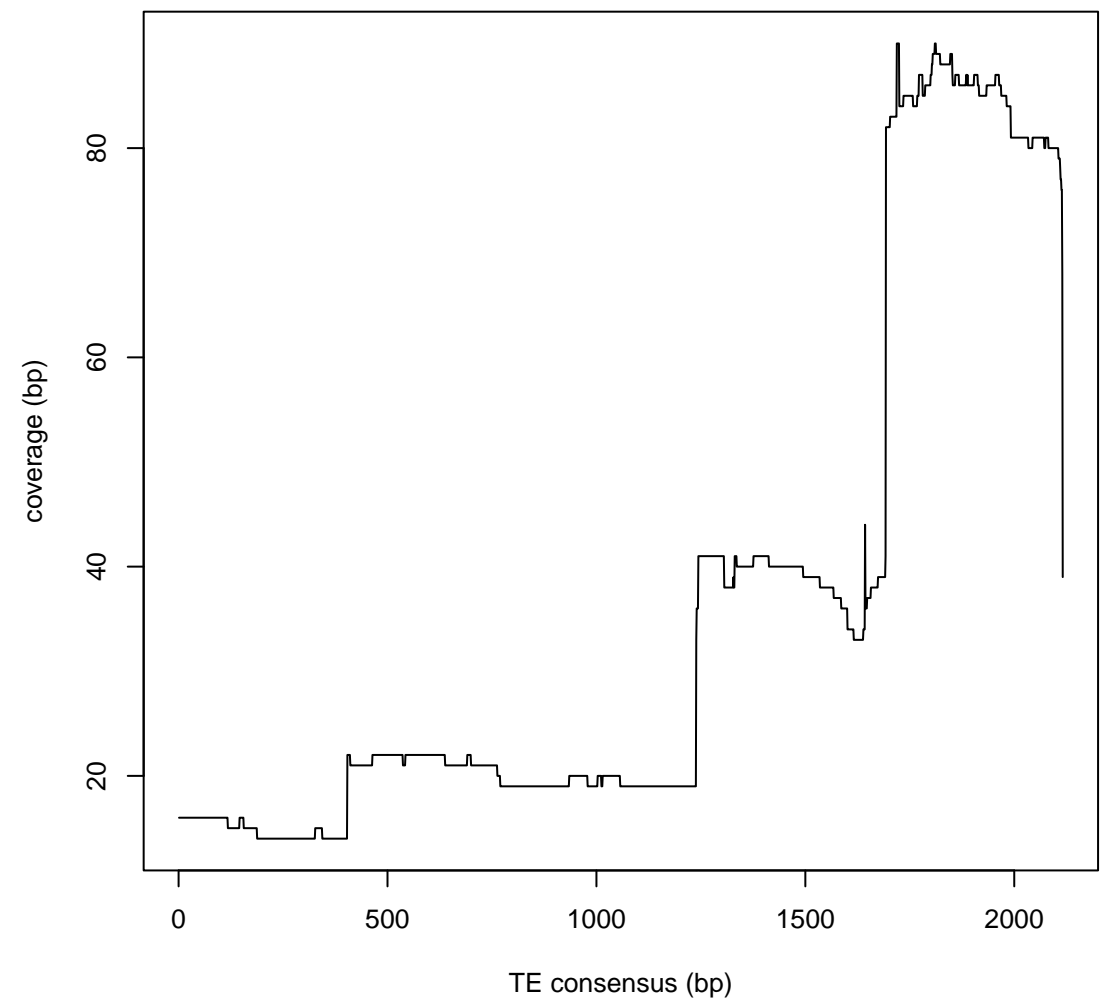


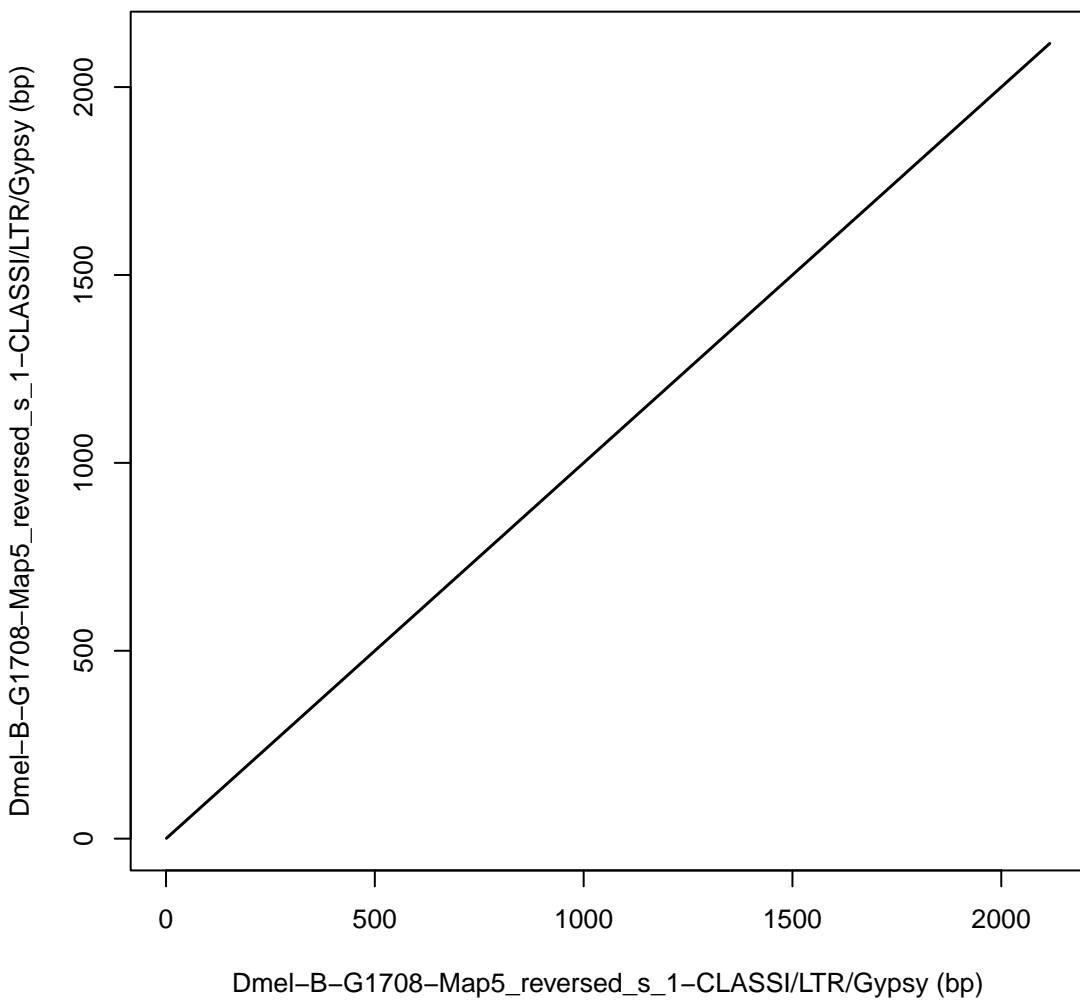
TE: Dmel-B-G1708-Map5_reversed_s_1-CLASSI/LTR/Gypsy
consensus size: 2116bp; fragments: 143; full length: 4 (>=1904.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

