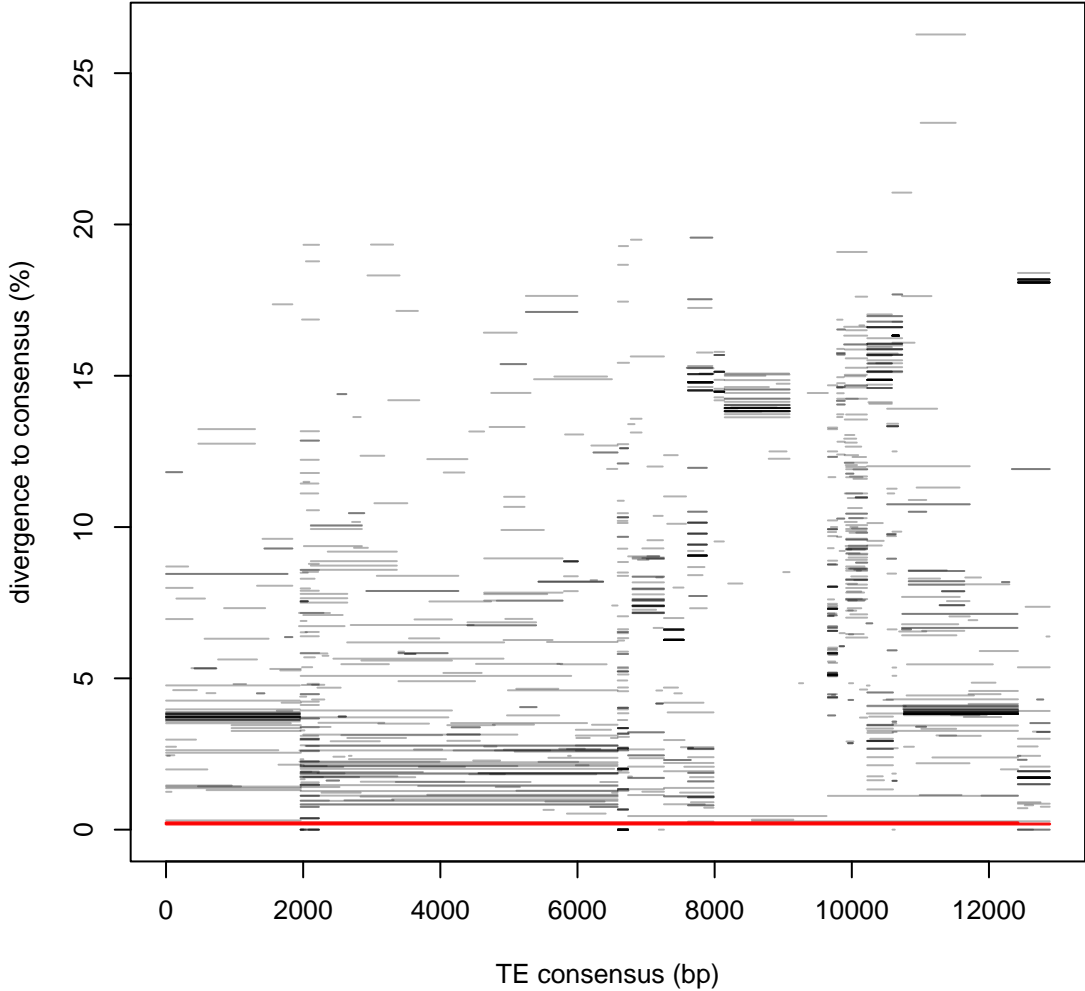
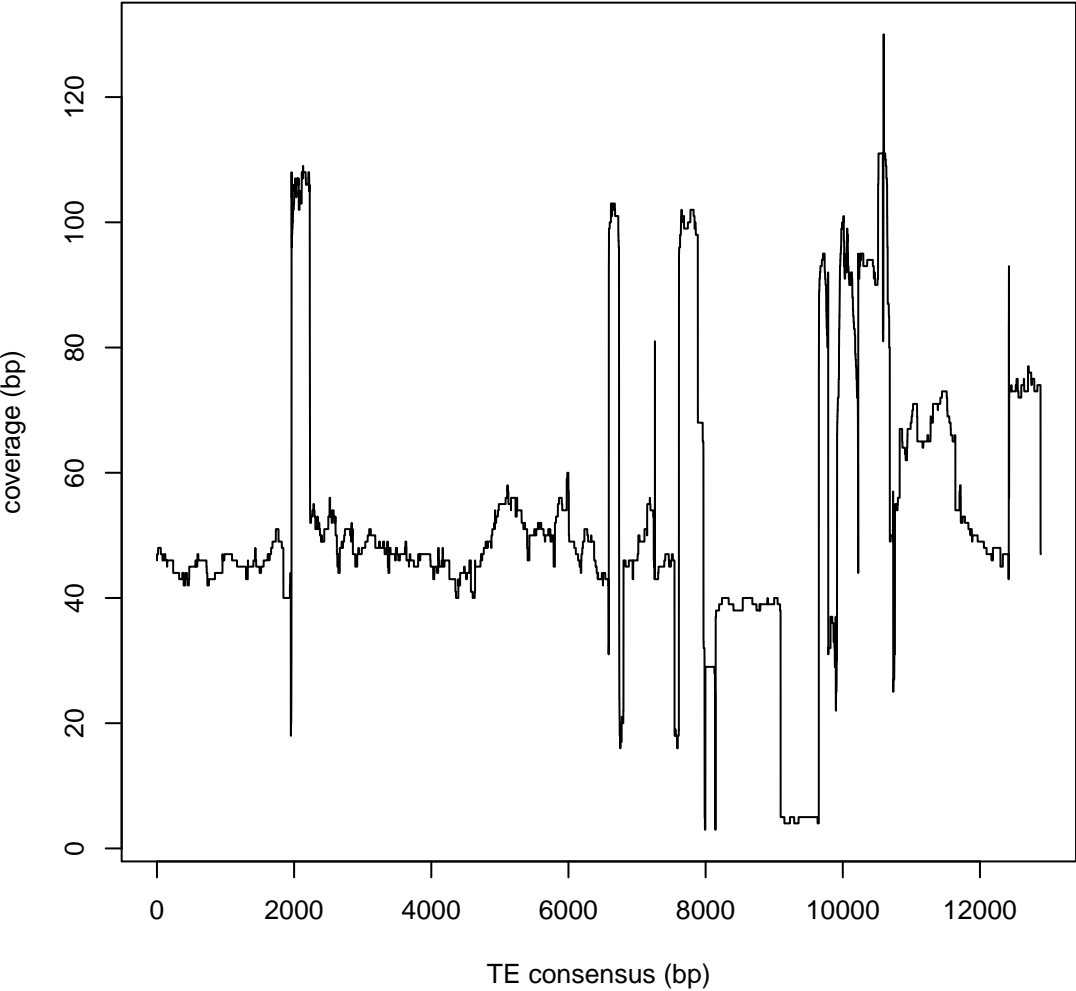


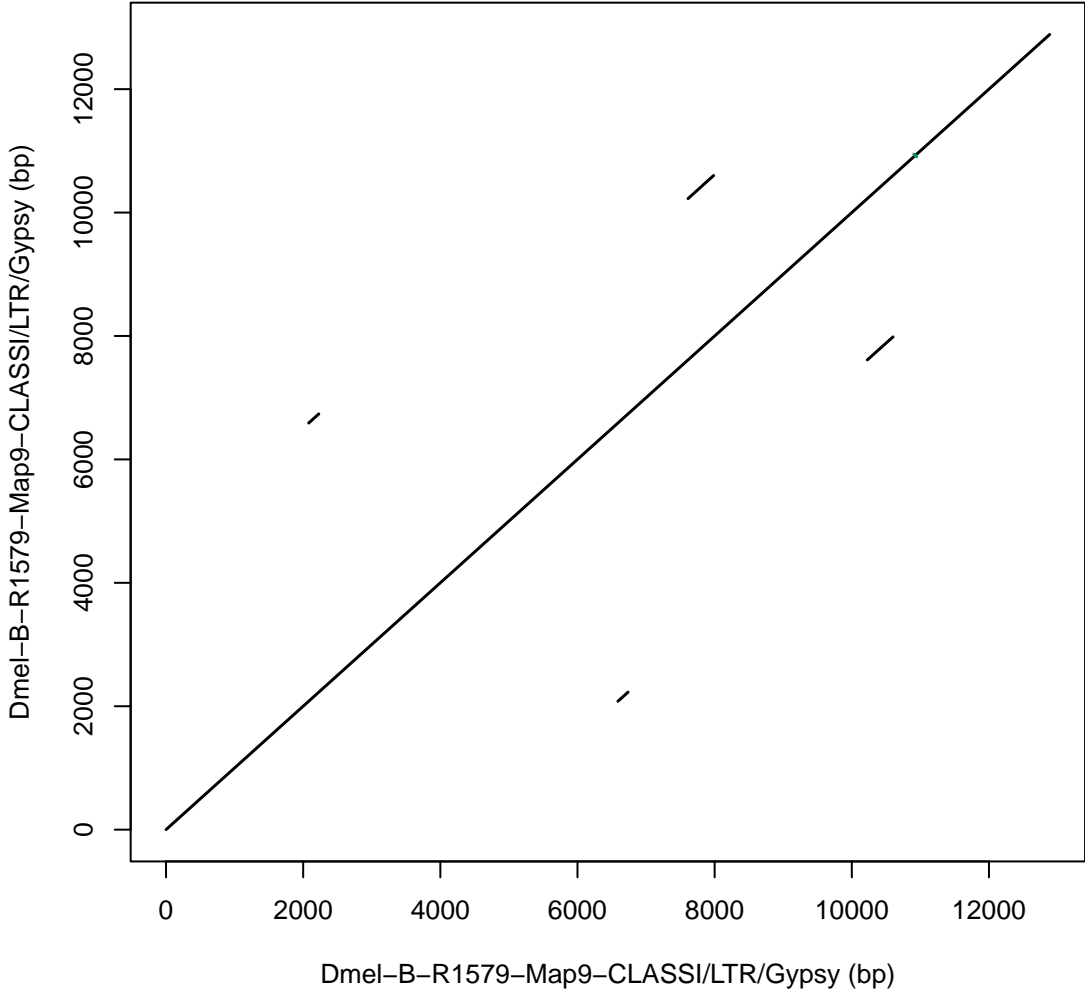
TE: Dmel-B-R1579-Map9-CLASSI/LTR/Gypsy
consensus size: 12886bp; fragments: 1296; full length: 2 (>=11597.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

