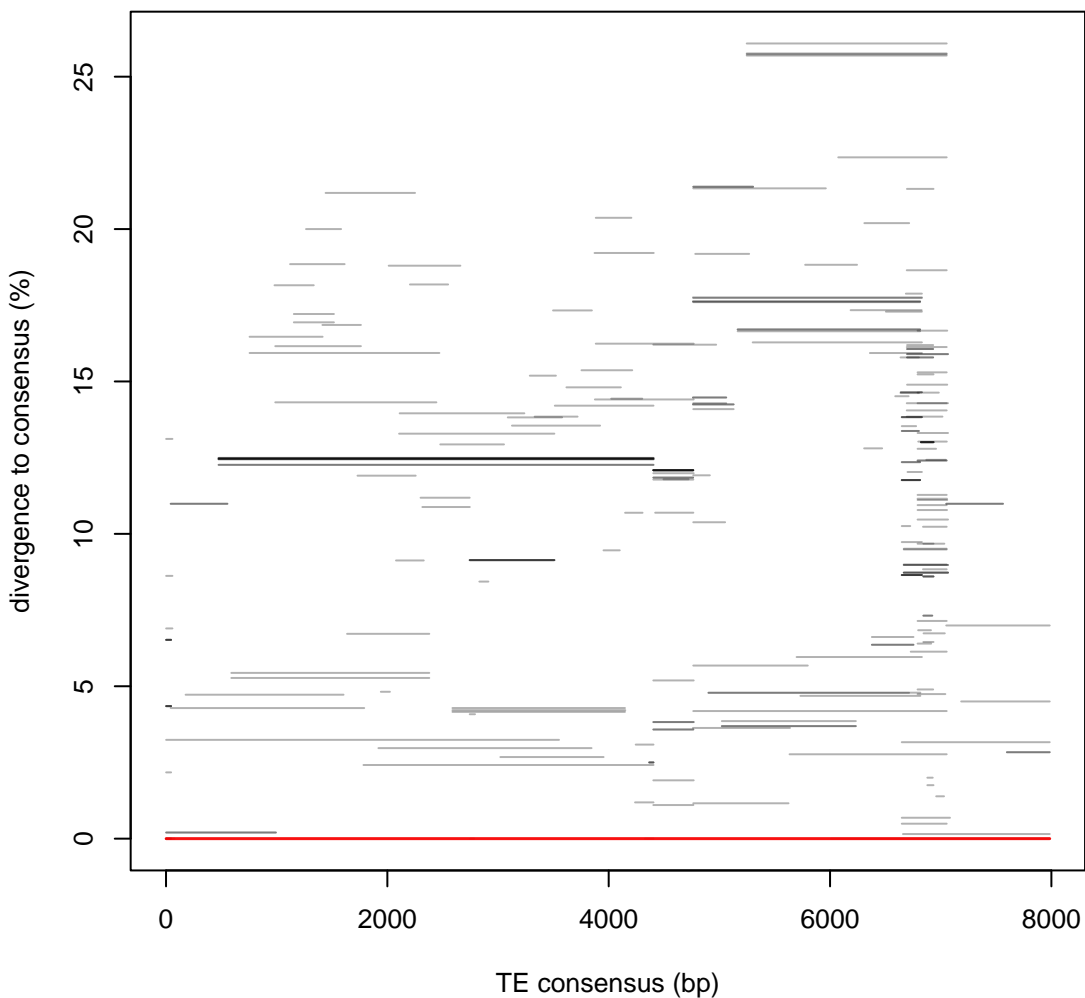
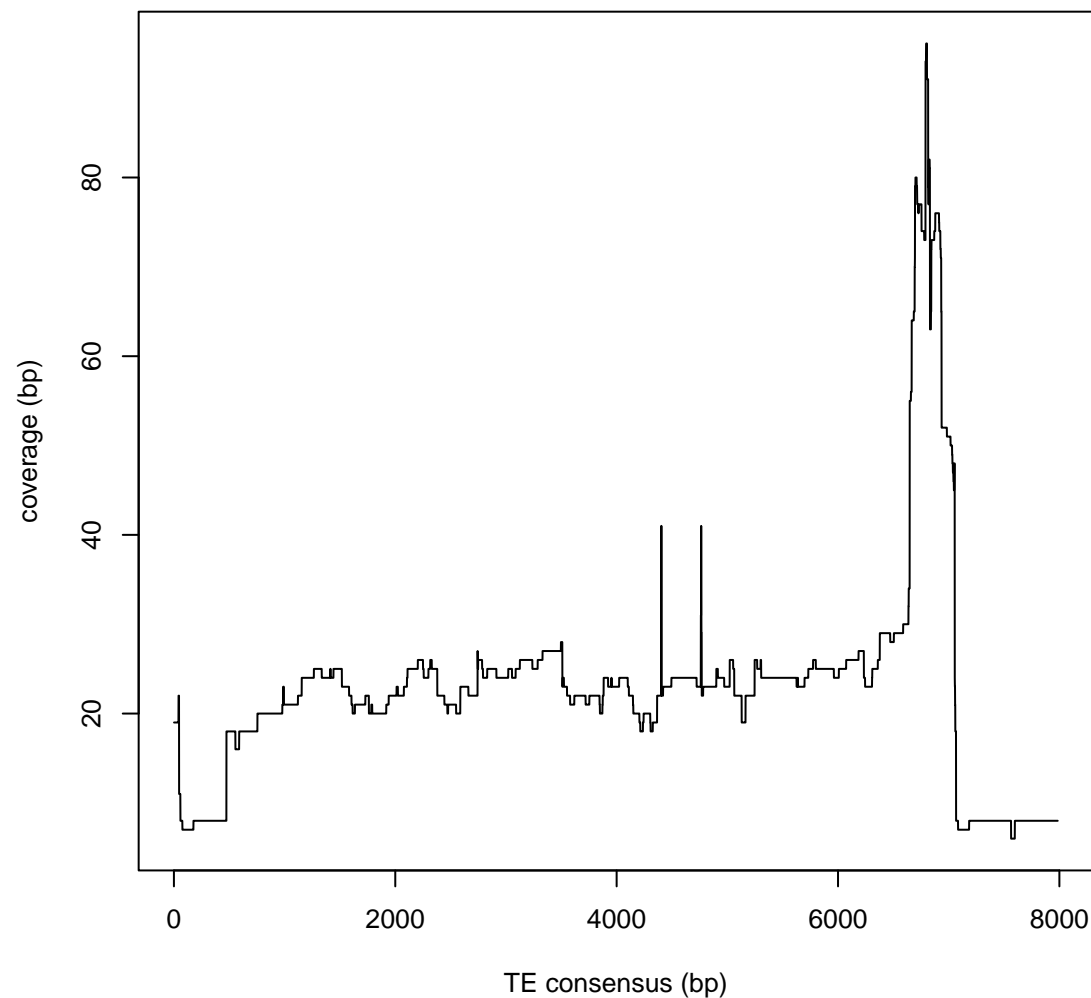


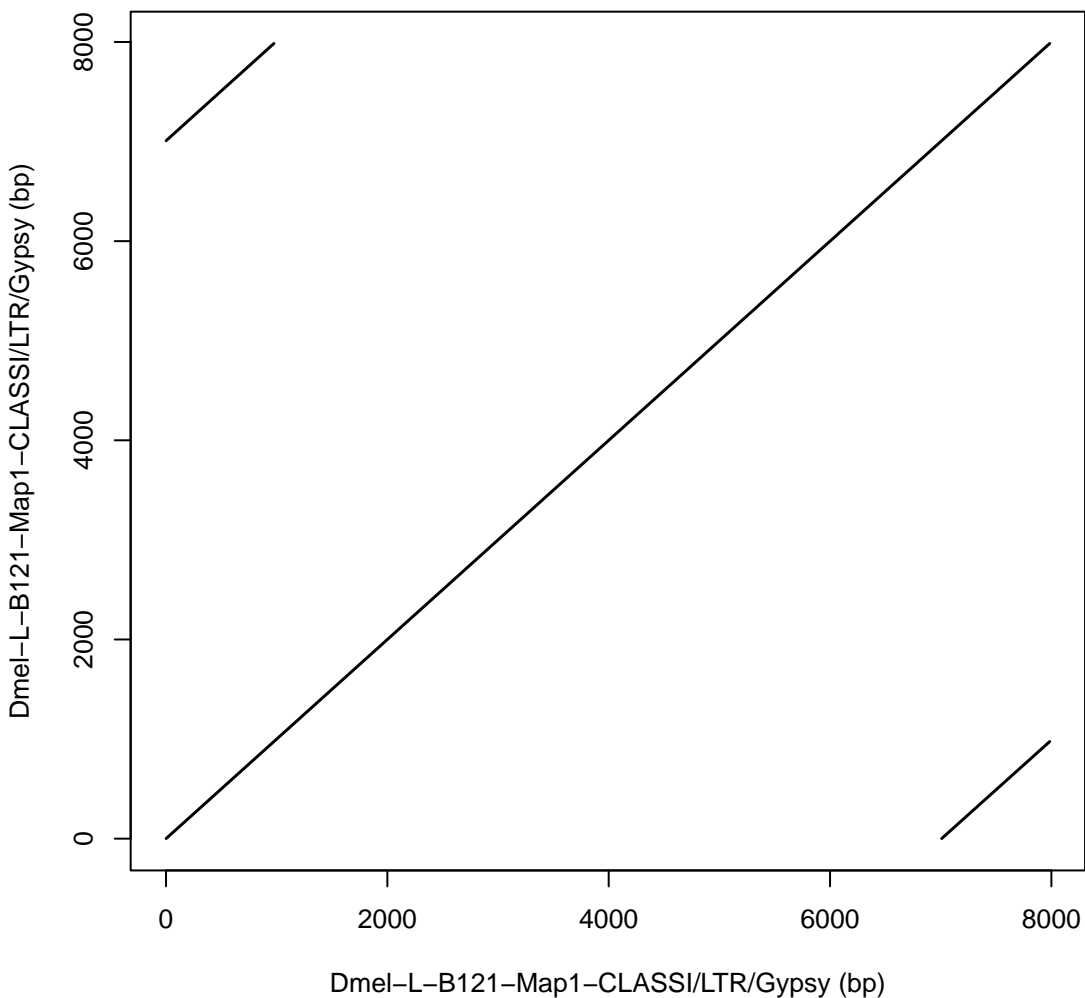
TE: Dmel-L-B121-Map1-CLASSI/LTR/Gypsy
consensus size: 7985bp; fragments: 259; full length: 1 (>=7186.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

