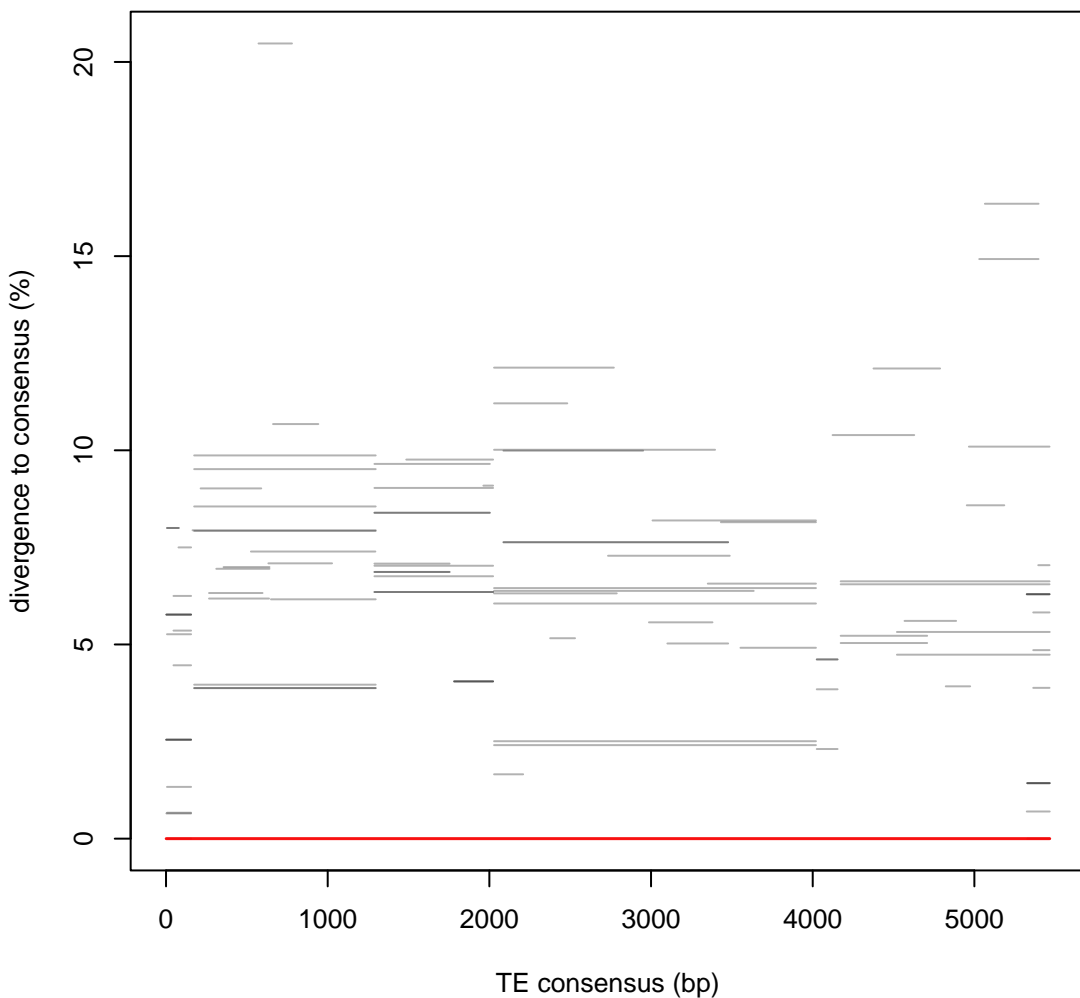
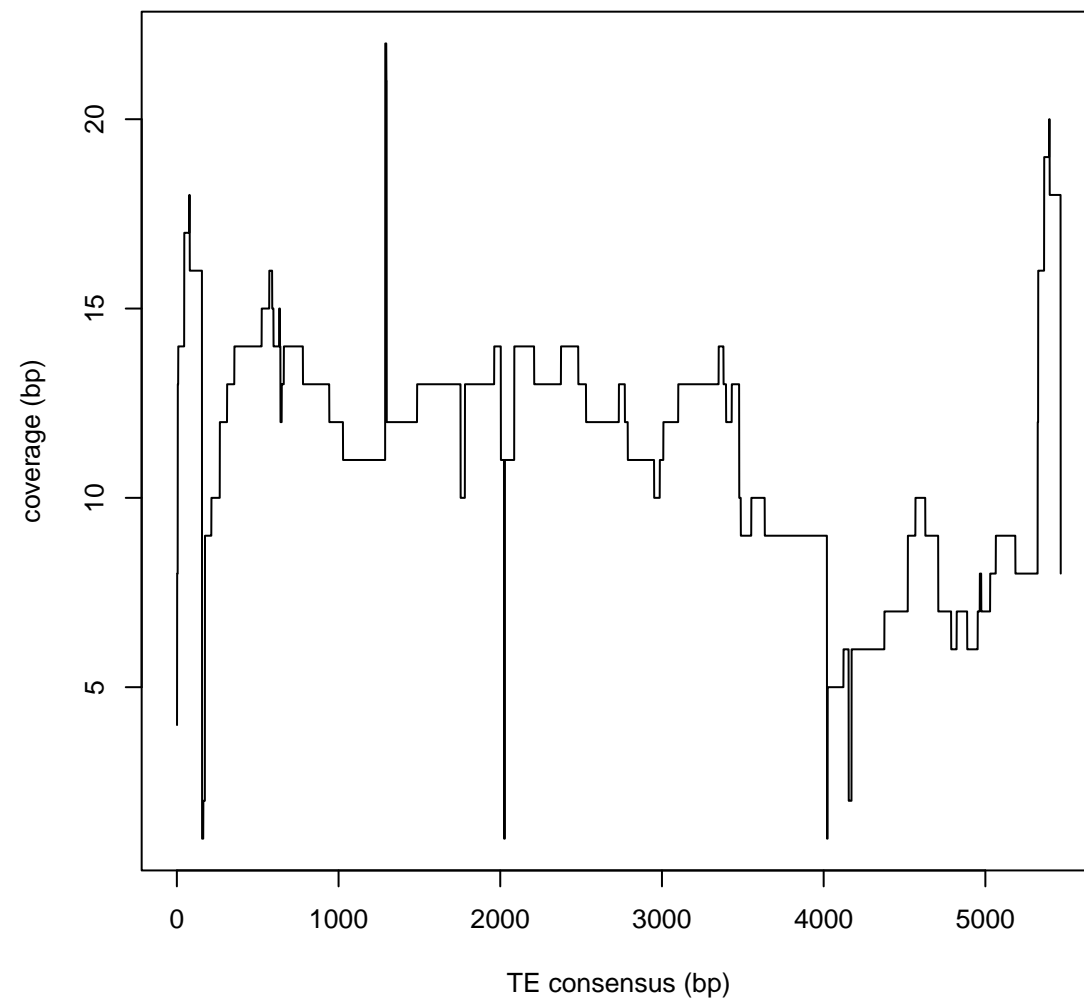


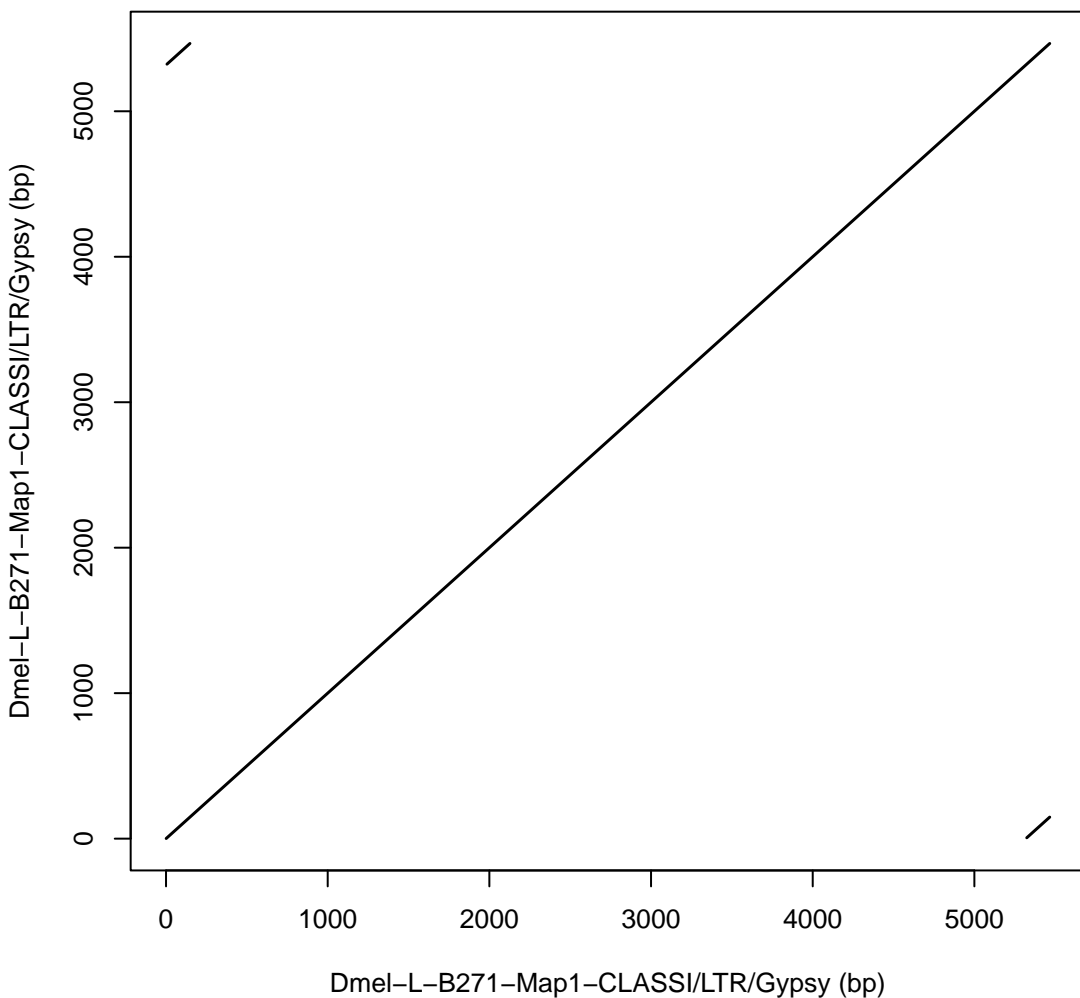
TE: Dmel-L-B271-Map1-CLASSI/LTR/Gypsy
consensus size: 5466bp; fragments: 103; full length: 1 (>=4919.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

