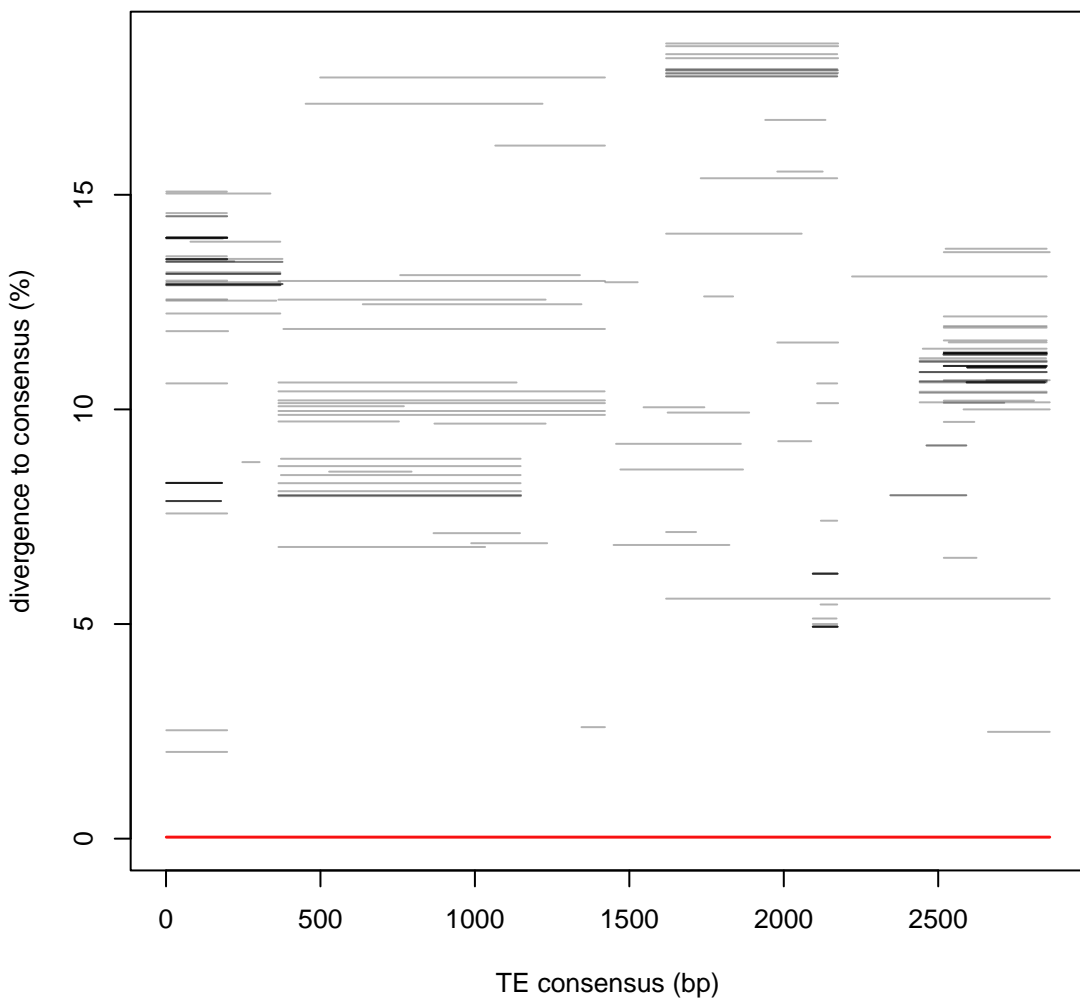
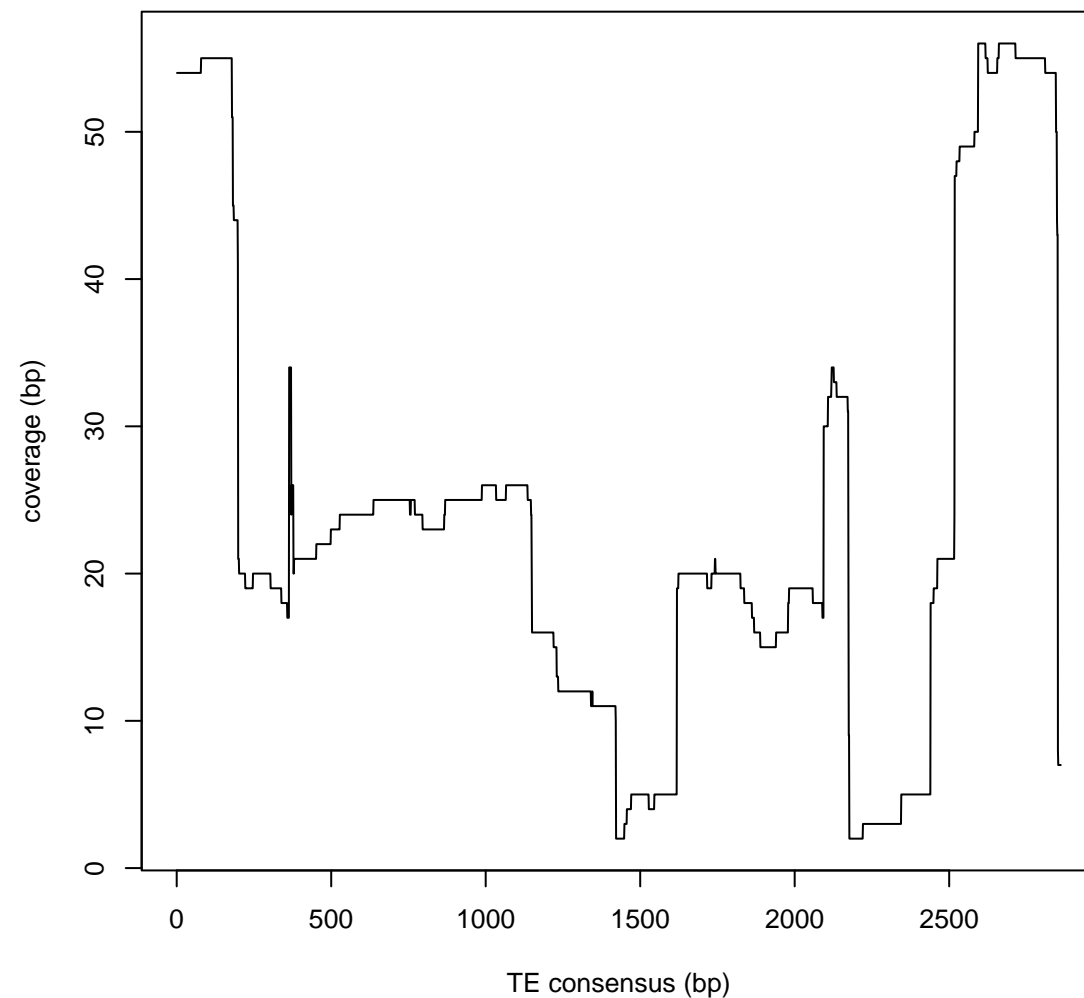


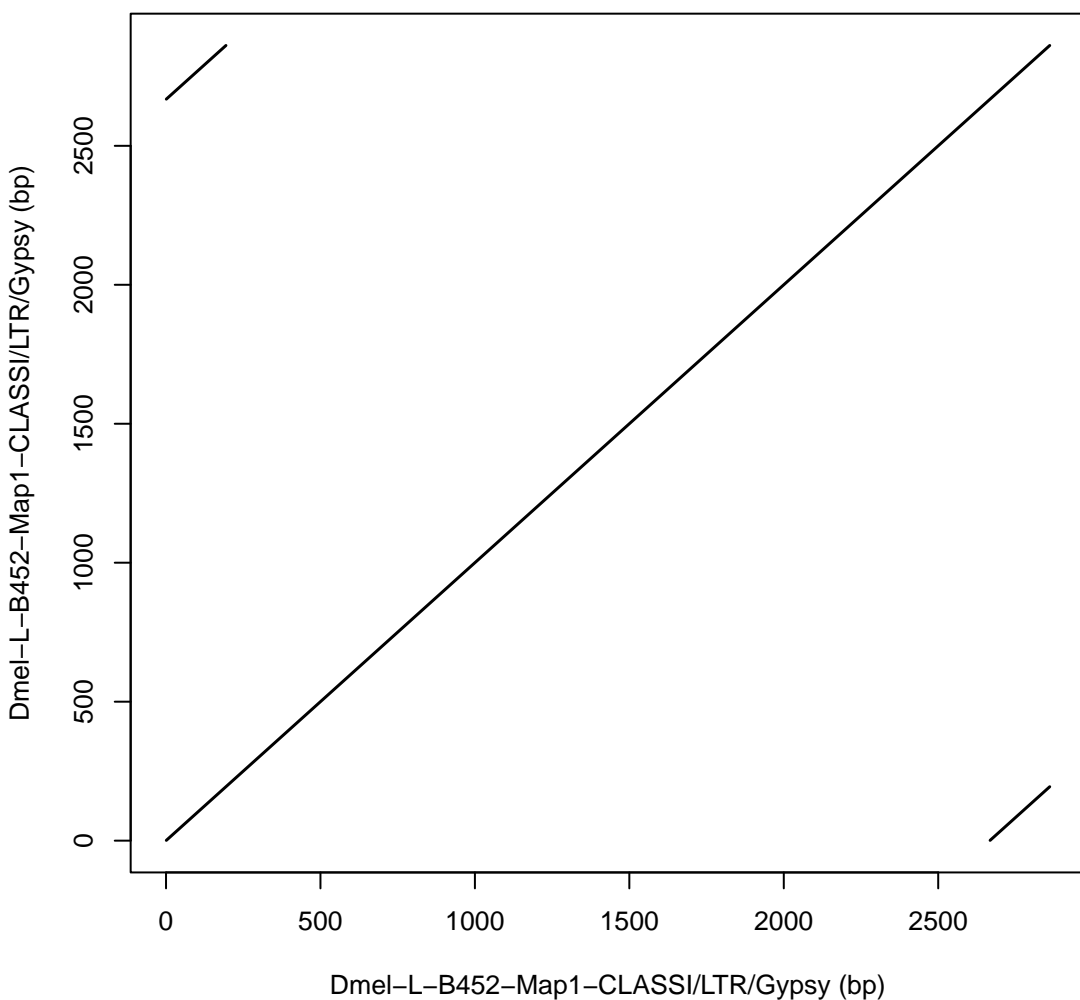
TE: Dmel-L-B452-Map1-CLASSI/LTR/Gypsy
consensus size: 2861bp; fragments: 189; full length: 1 (>=2574.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

