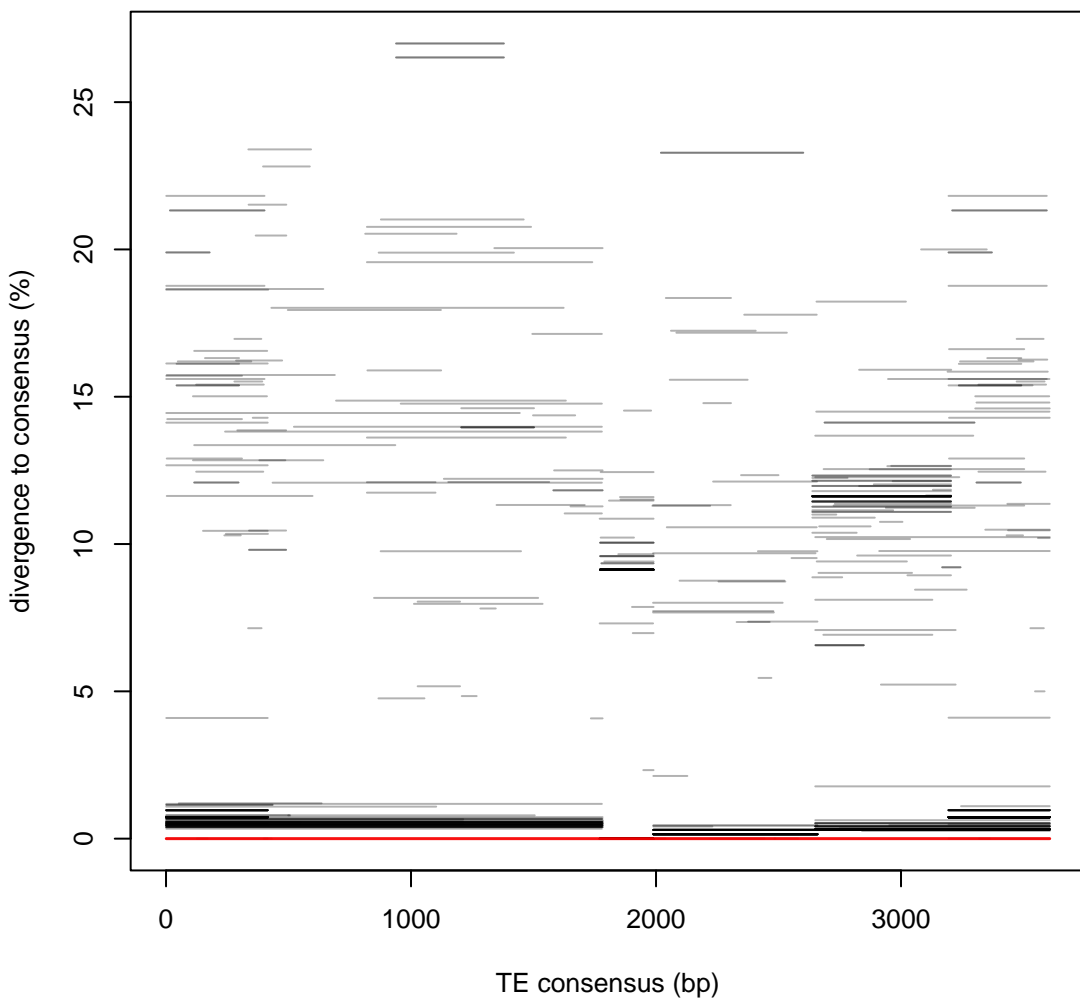
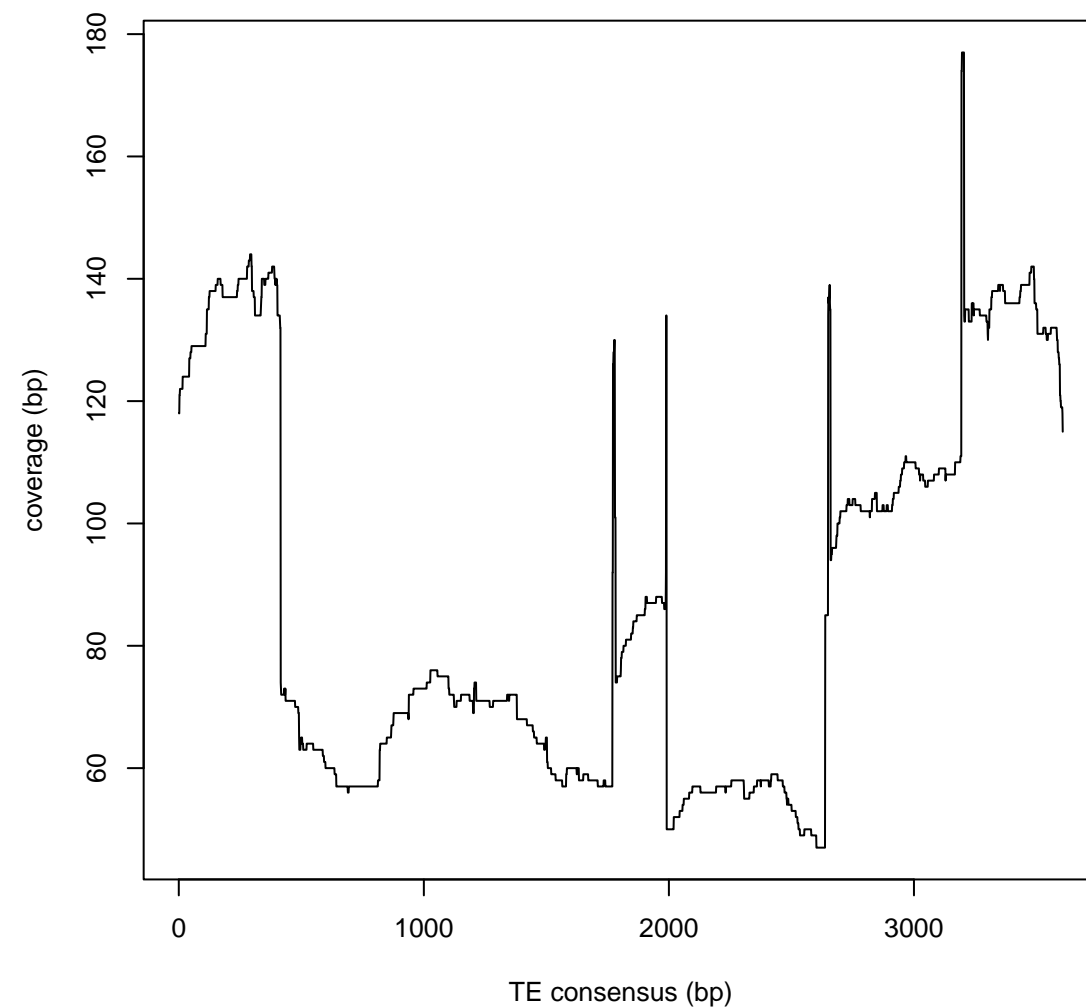


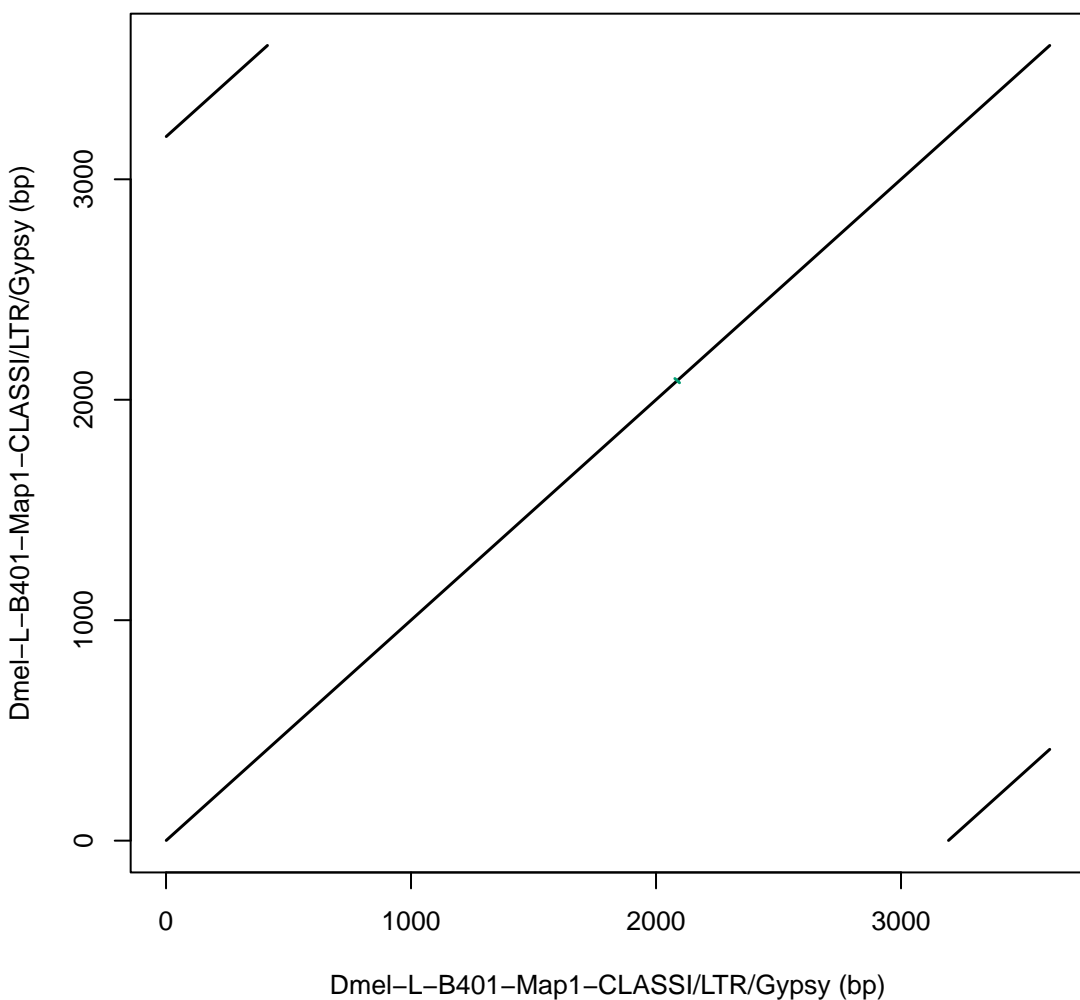
TE: Dmel-L-B401-Map1-CLASSI/LTR/Gypsy
consensus size: 3607bp; fragments: 578; full length: 1 (≥ 3246.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

