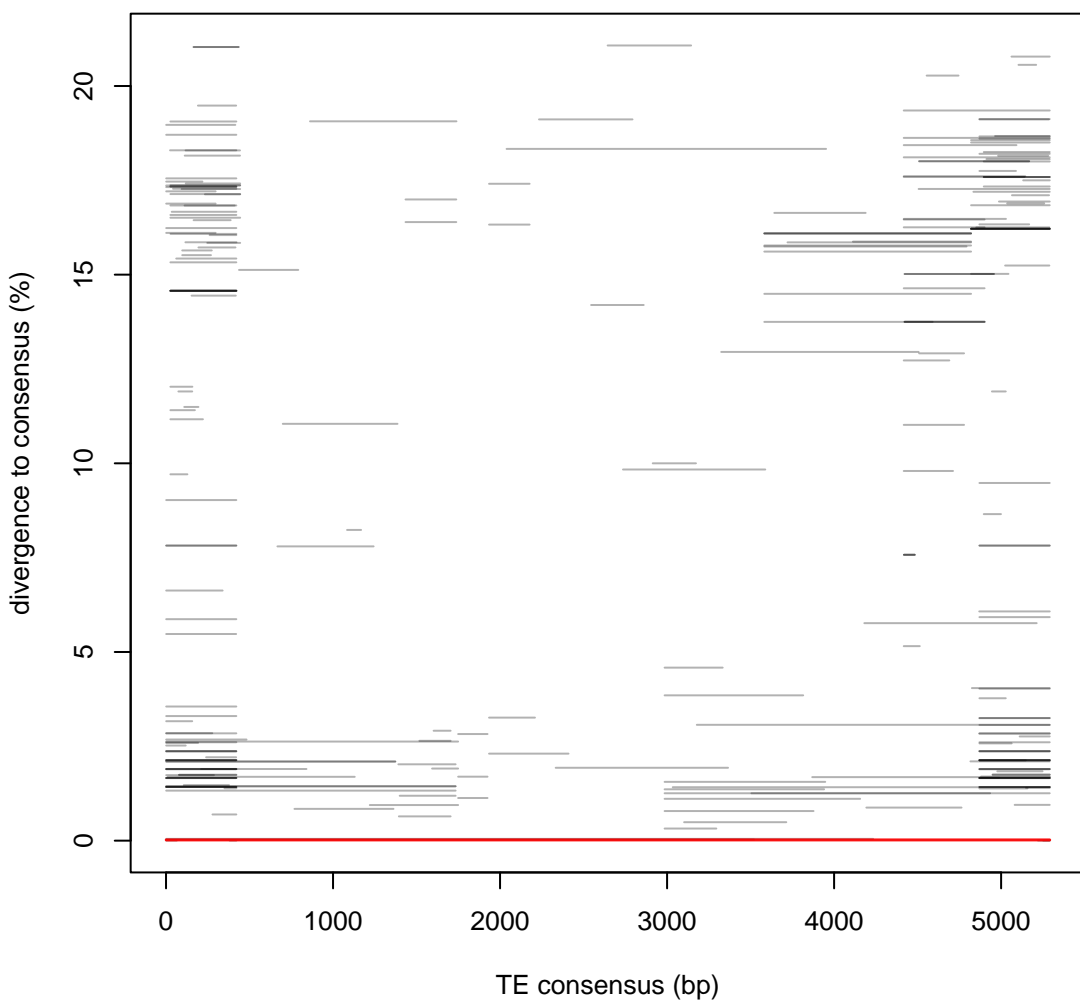
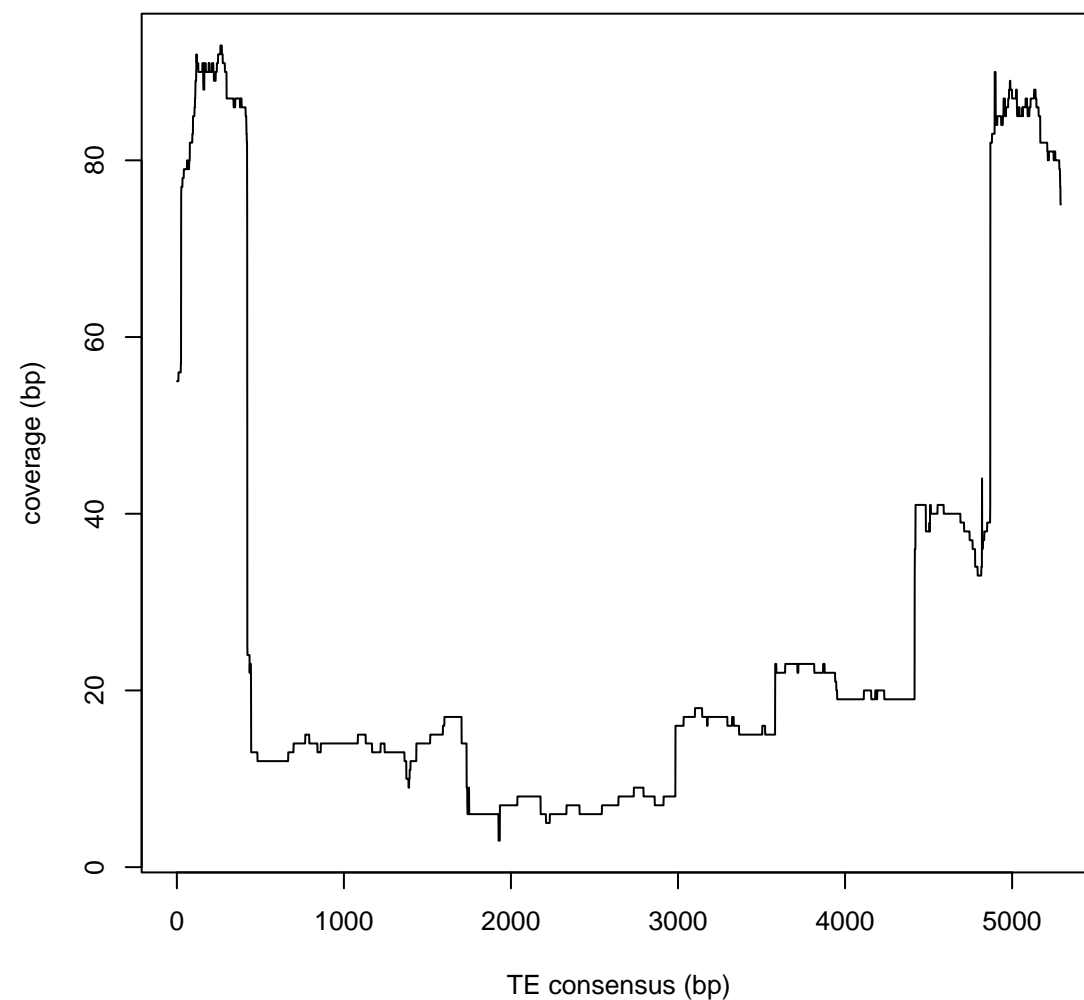


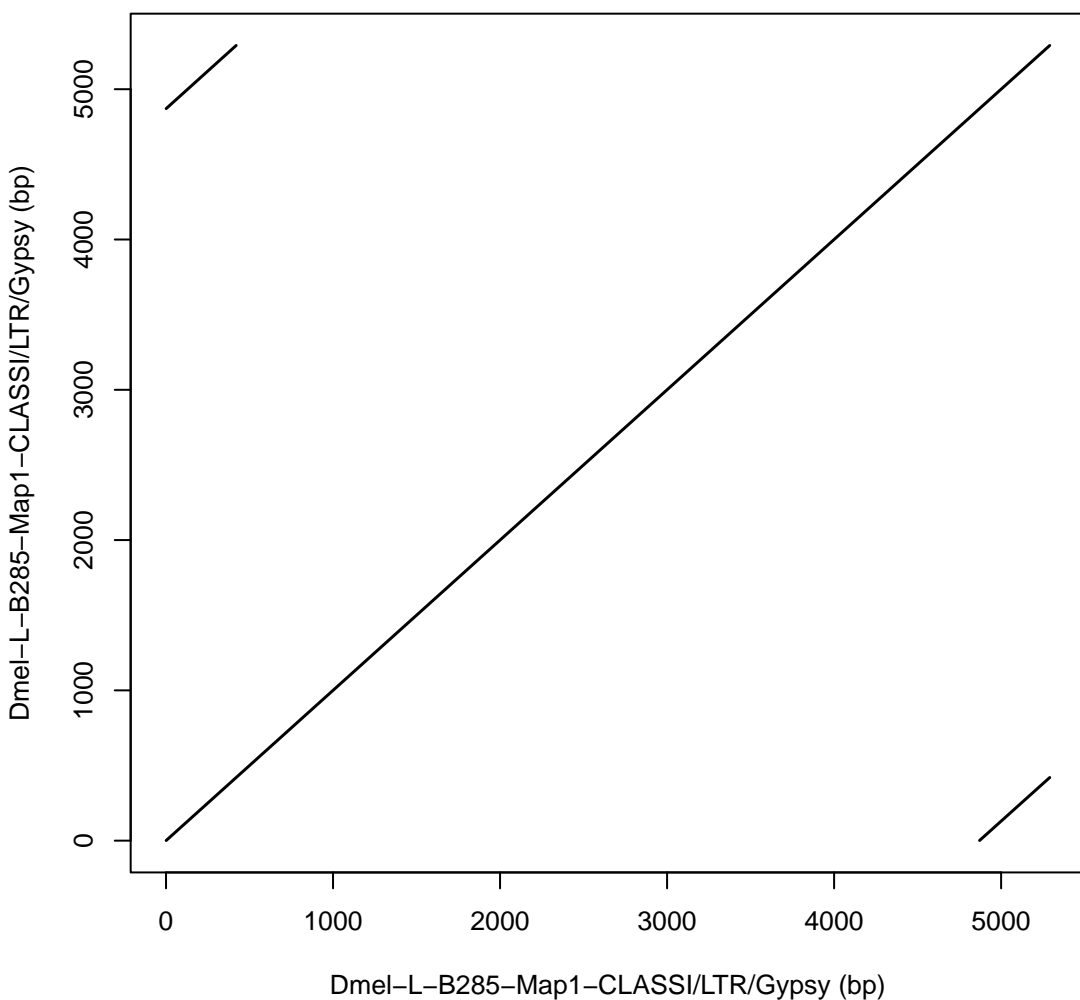
TE: Dmel-L-B285-Map1-CLASSI/LTR/Gypsy
consensus size: 5291bp; fragments: 274; full length: 1 (≥ 4761.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

