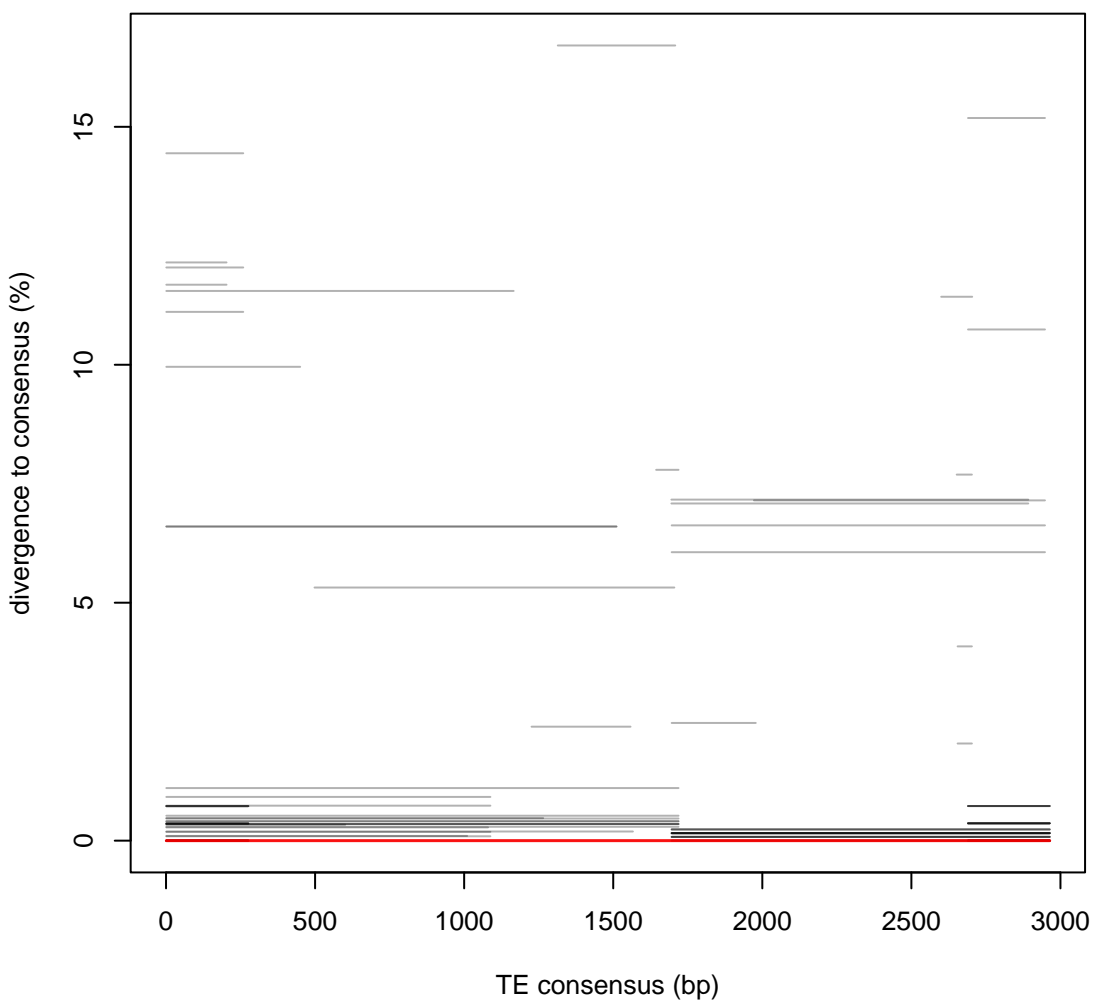
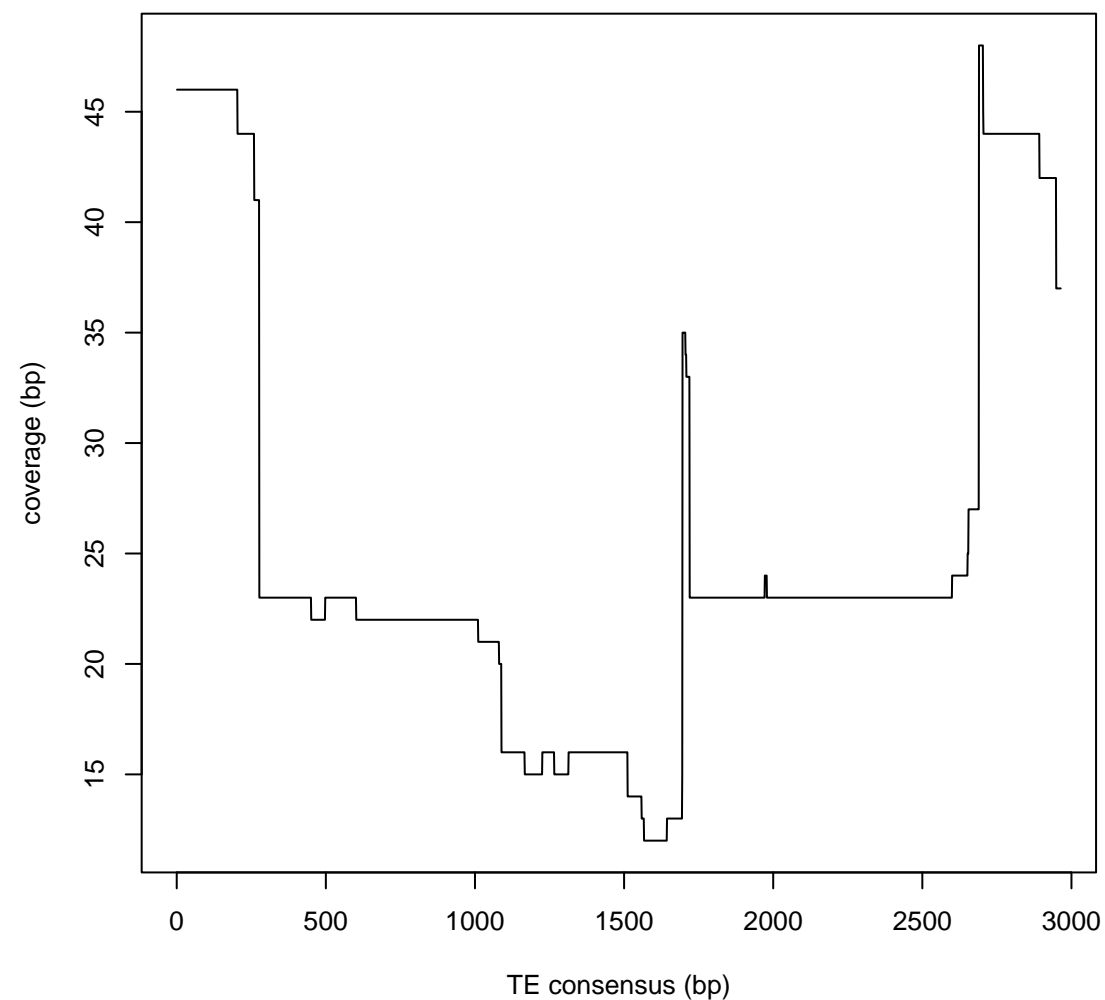


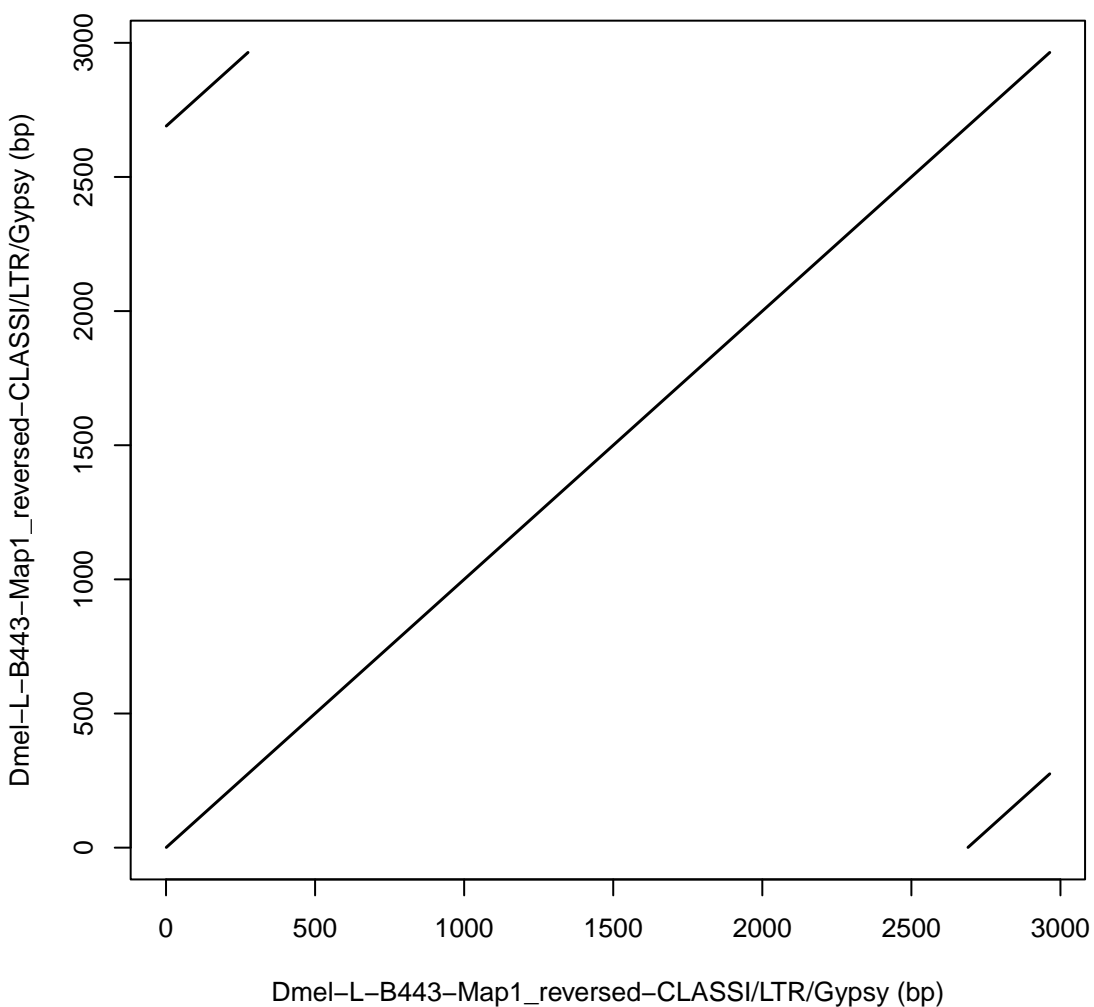
TE: Dmel-L-B443-Map1_reversed-CLASSI/LTR/Gypsy
consensus size: 2964bp; fragments: 98; full length: 1 (≥ 2667.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

