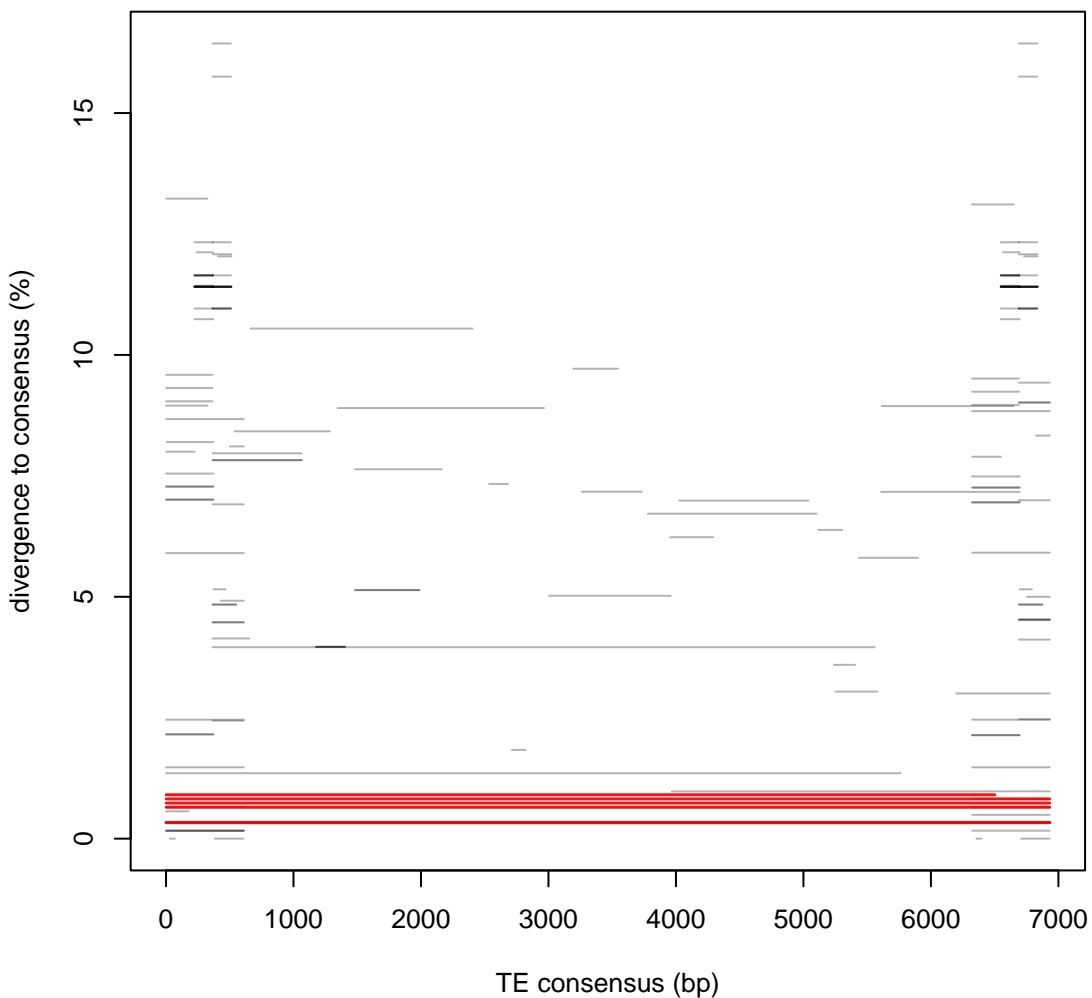
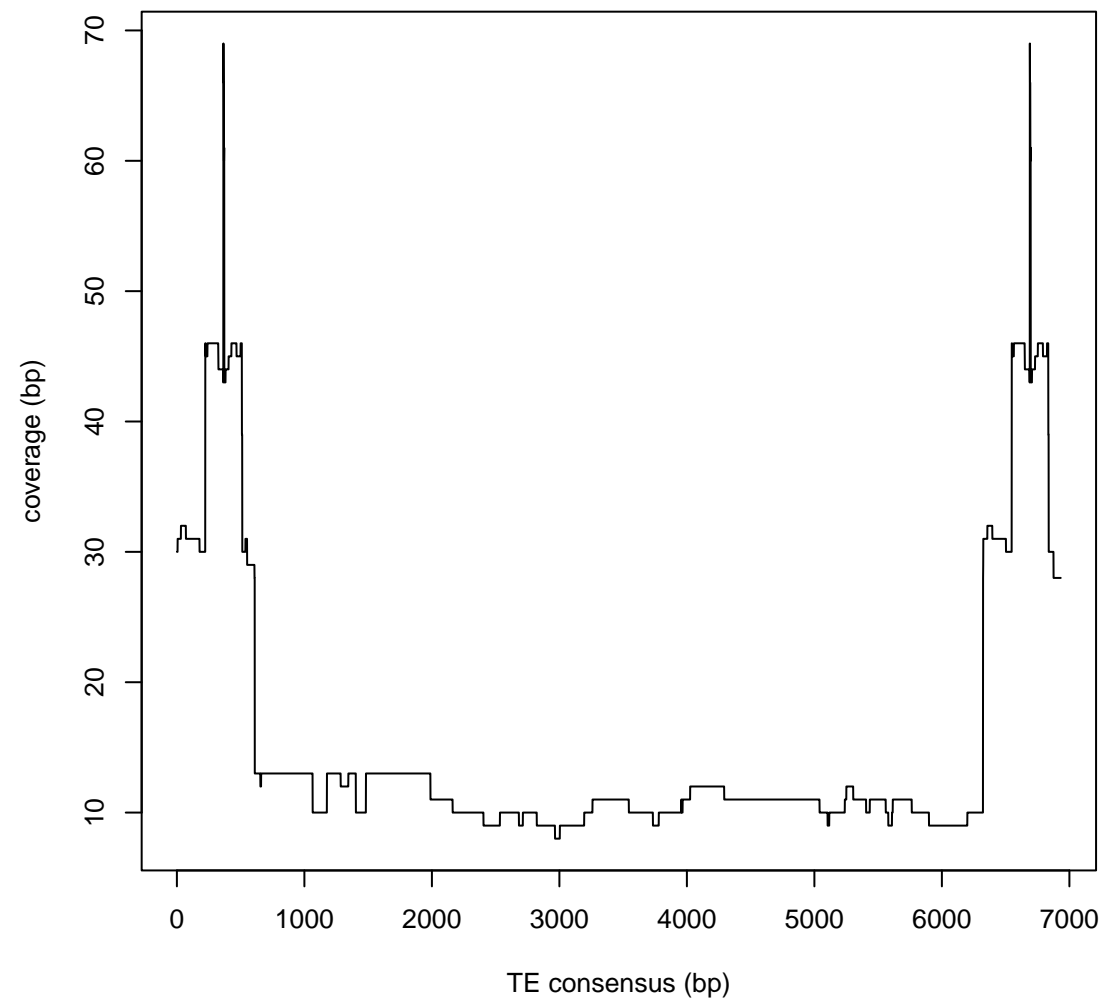


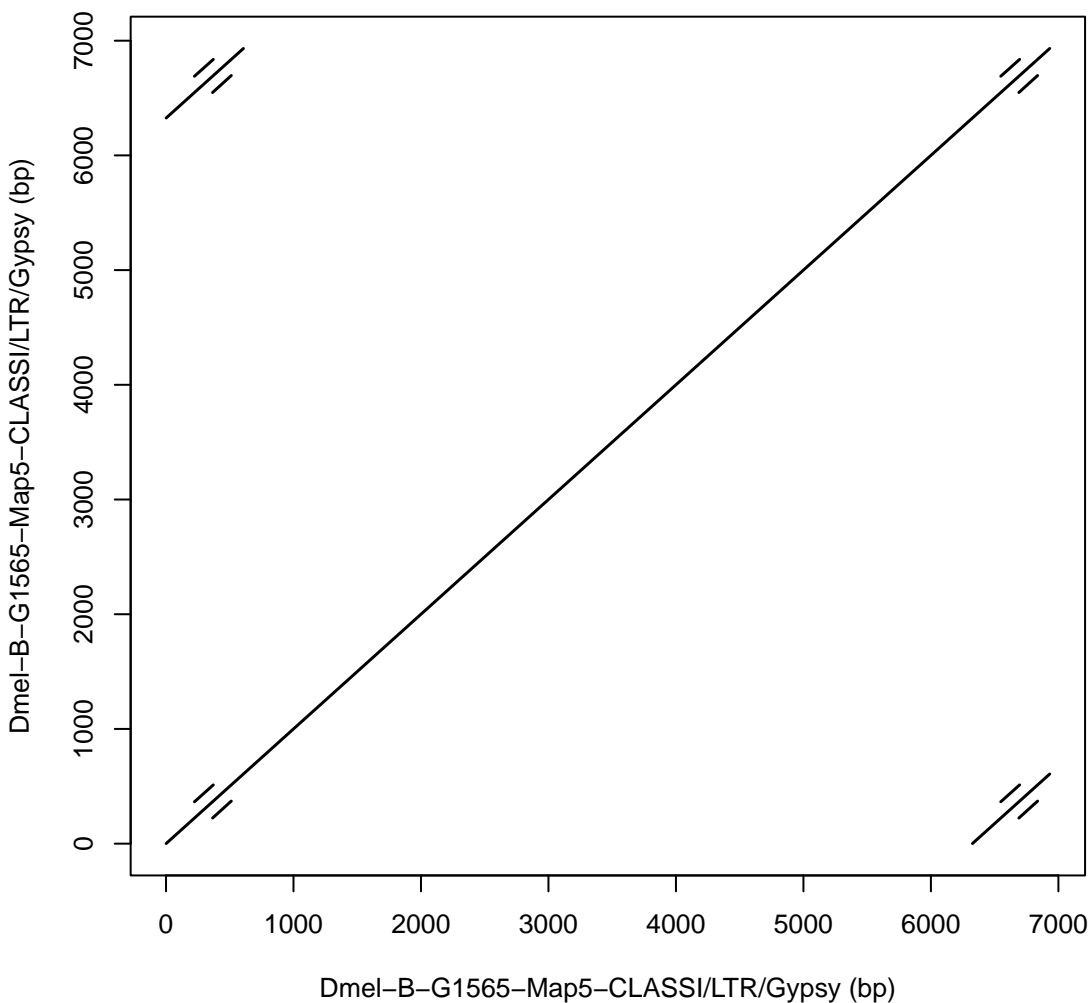
TE: Dmel-B-G1565-Map5-CLASSI/LTR/Gypsy
consensus size: 6932bp; fragments: 175; full length: 6 (≥ 6238.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

