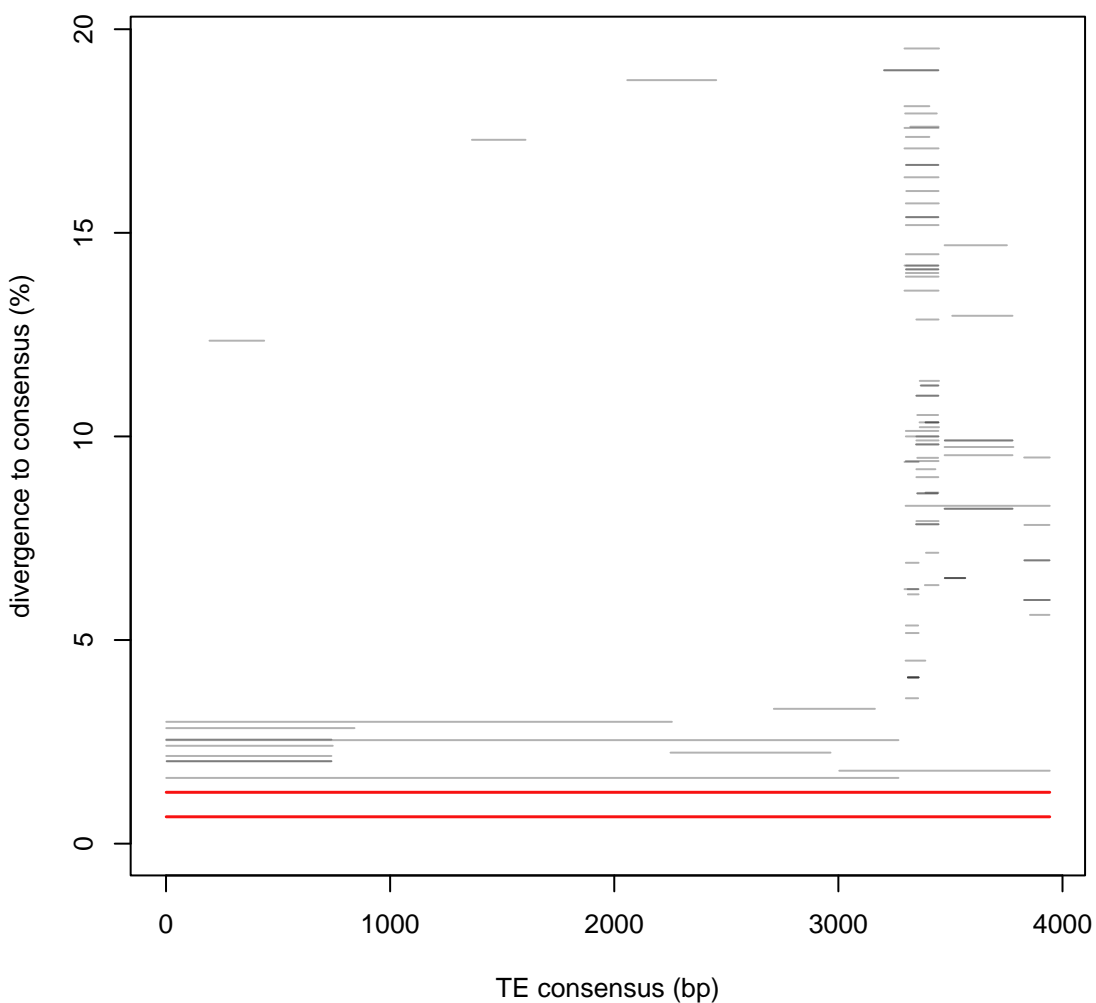
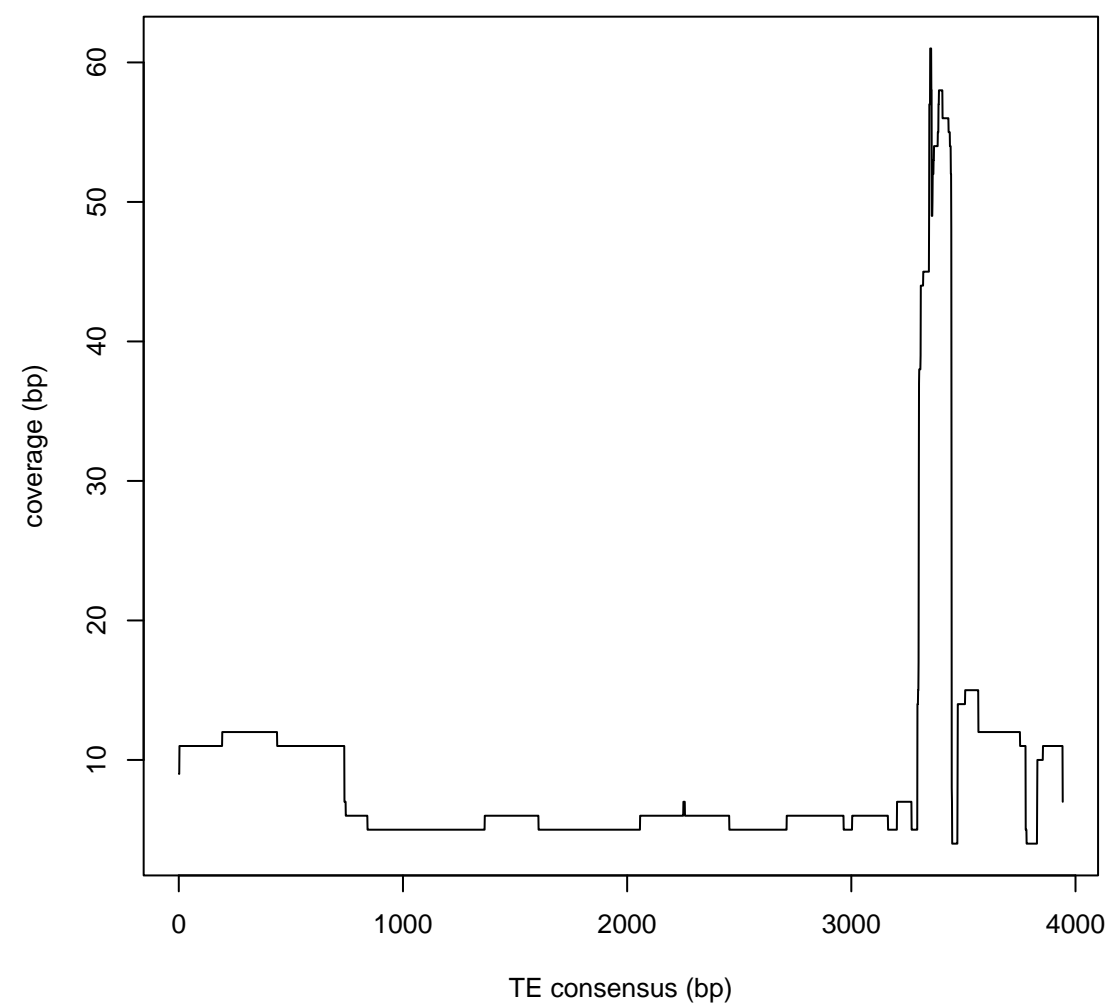


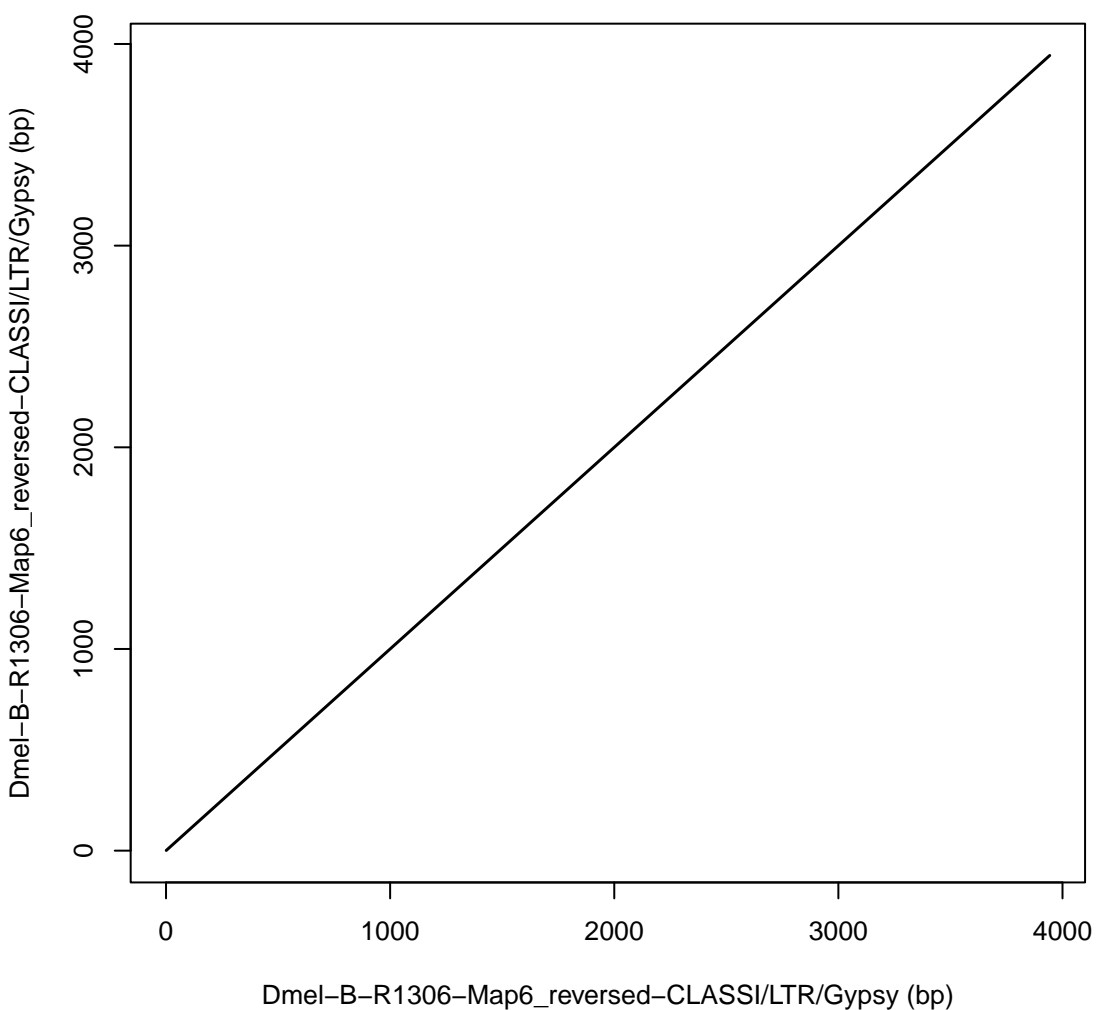
TE: Dmel-B-R1306-Map6_reversed-CLASSI/LTR/Gypsy
consensus size: 3943bp; fragments: 103; full length: 2 (>=3548.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

