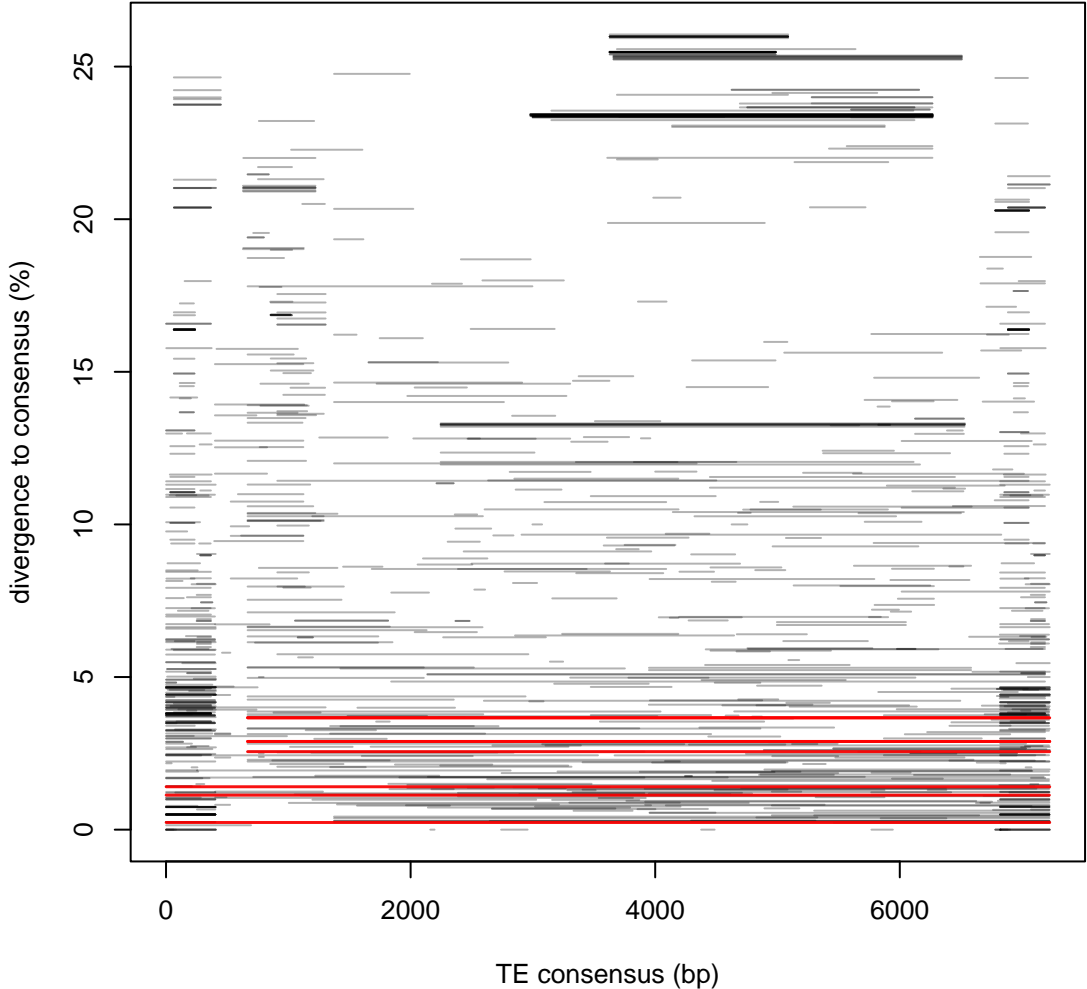
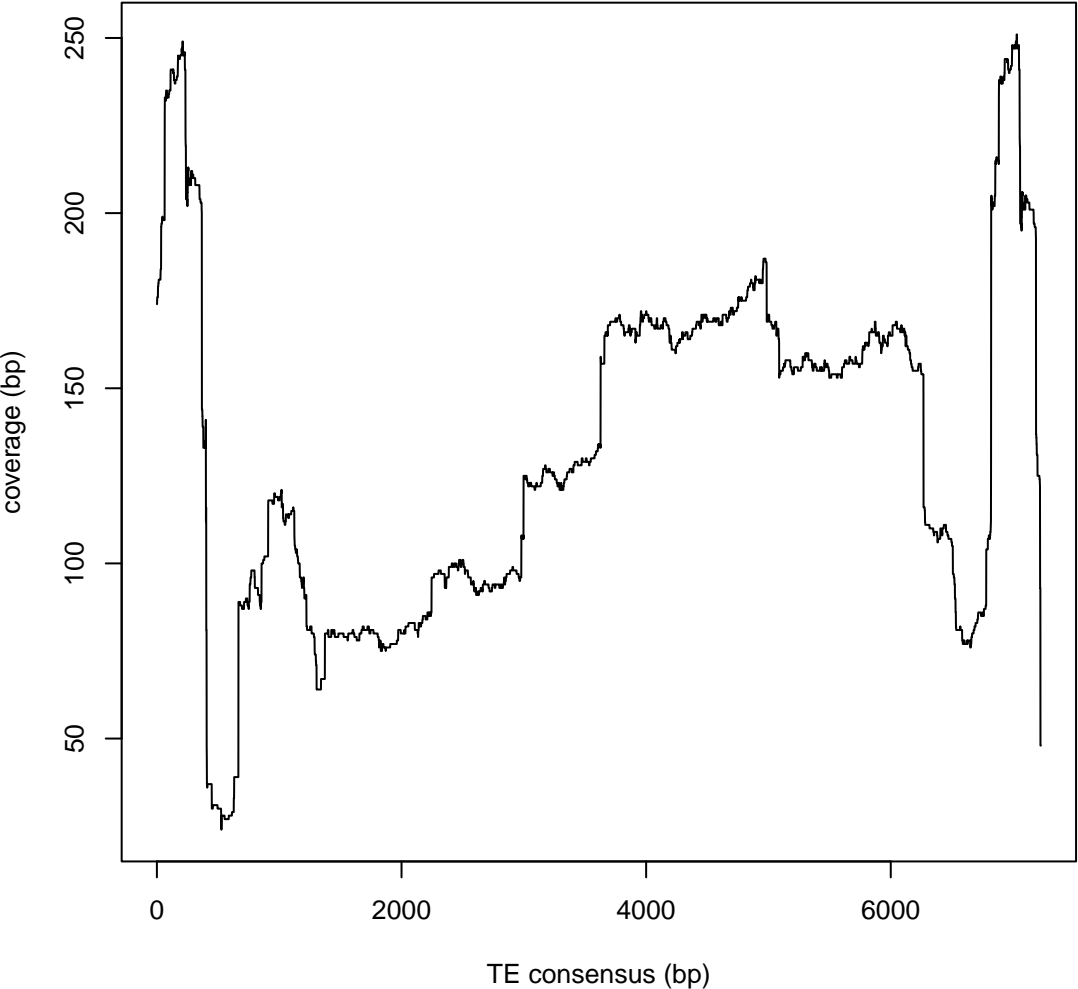


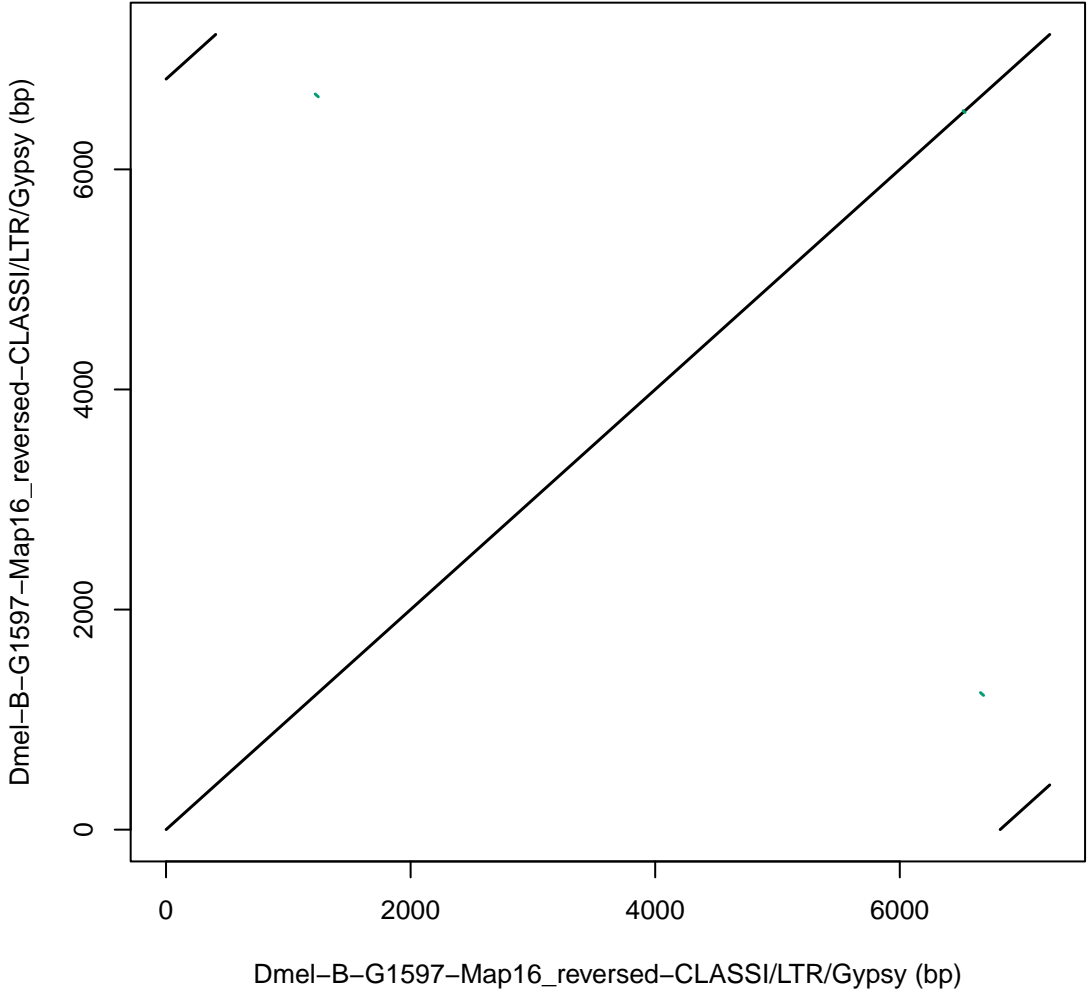
TE: Dmel-B-G1597-Map16_reversed-CLASSI/LTR/Gypsy
consensus size: 7226bp; fragments: 1065; full length: 8 (>=6503.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

