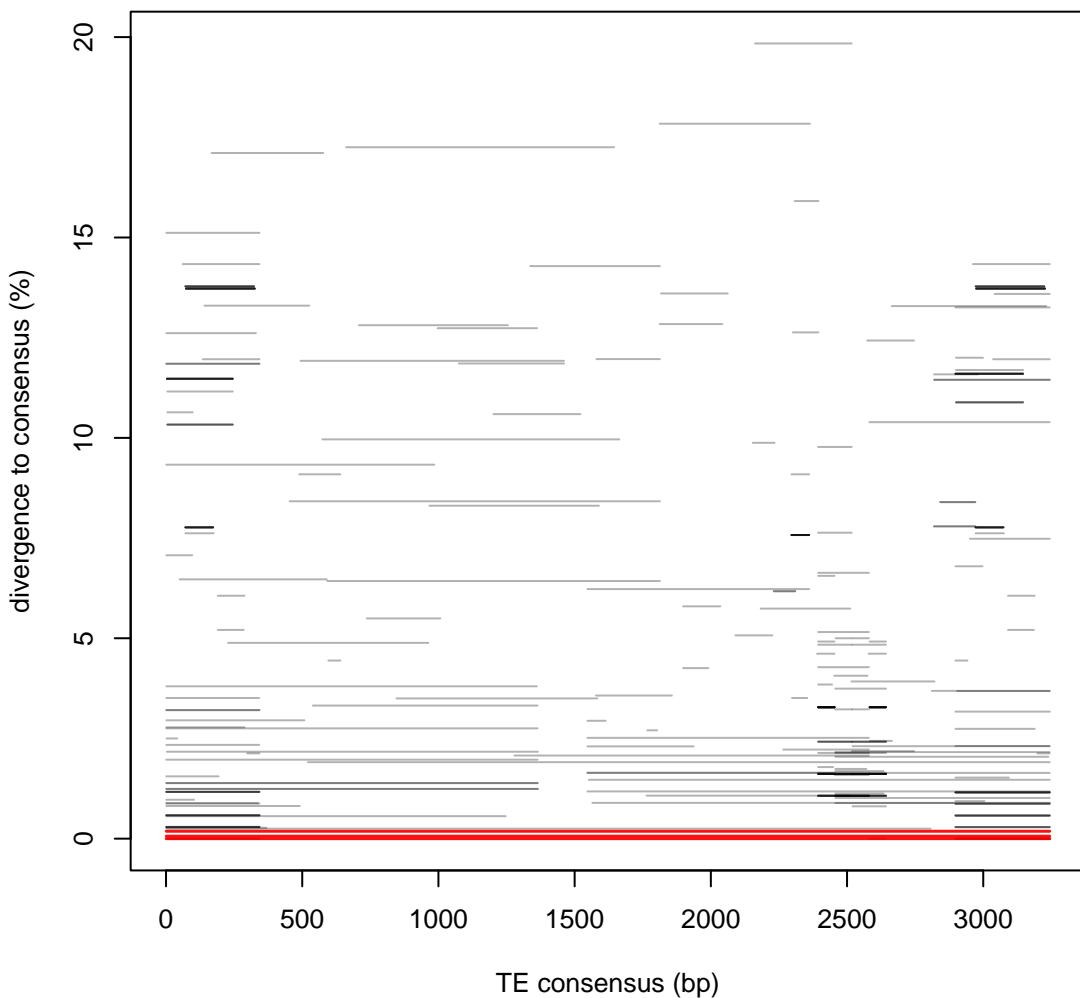
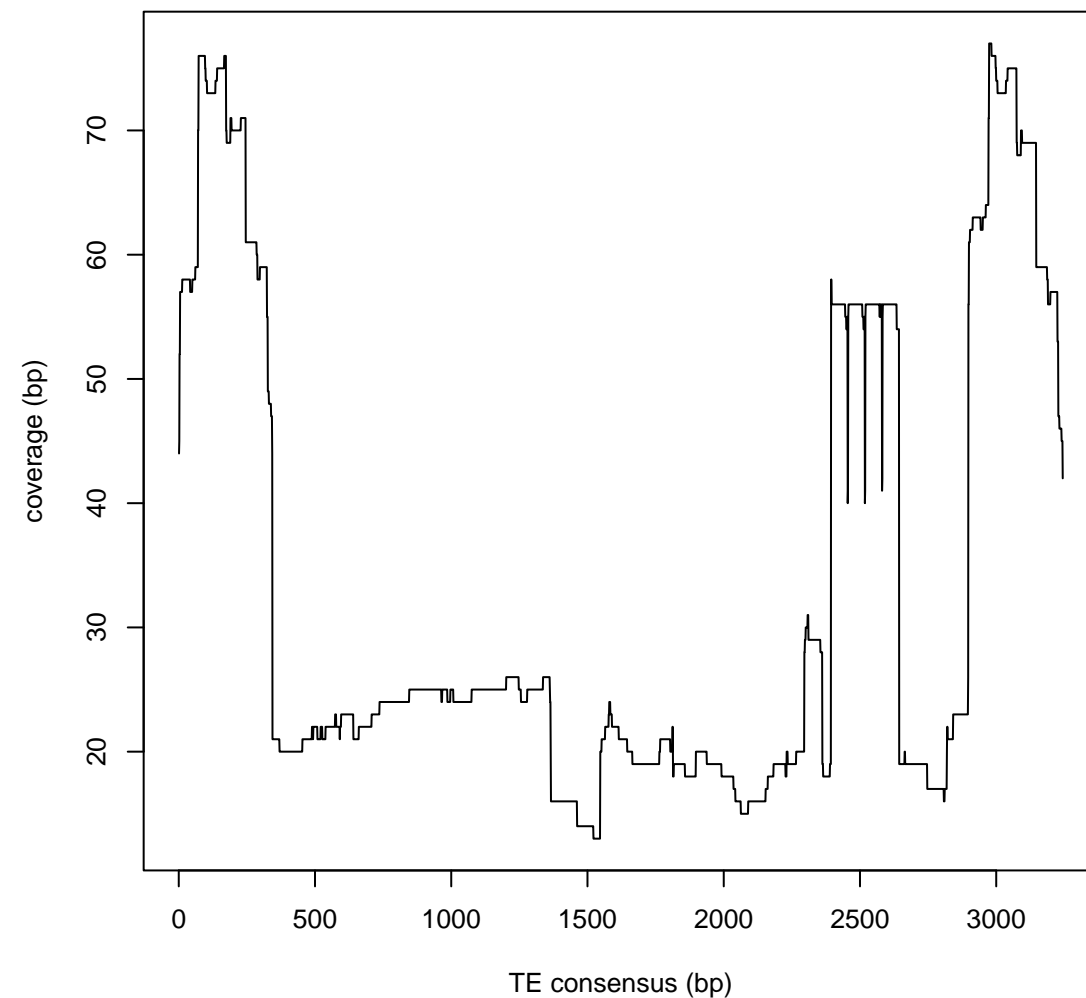


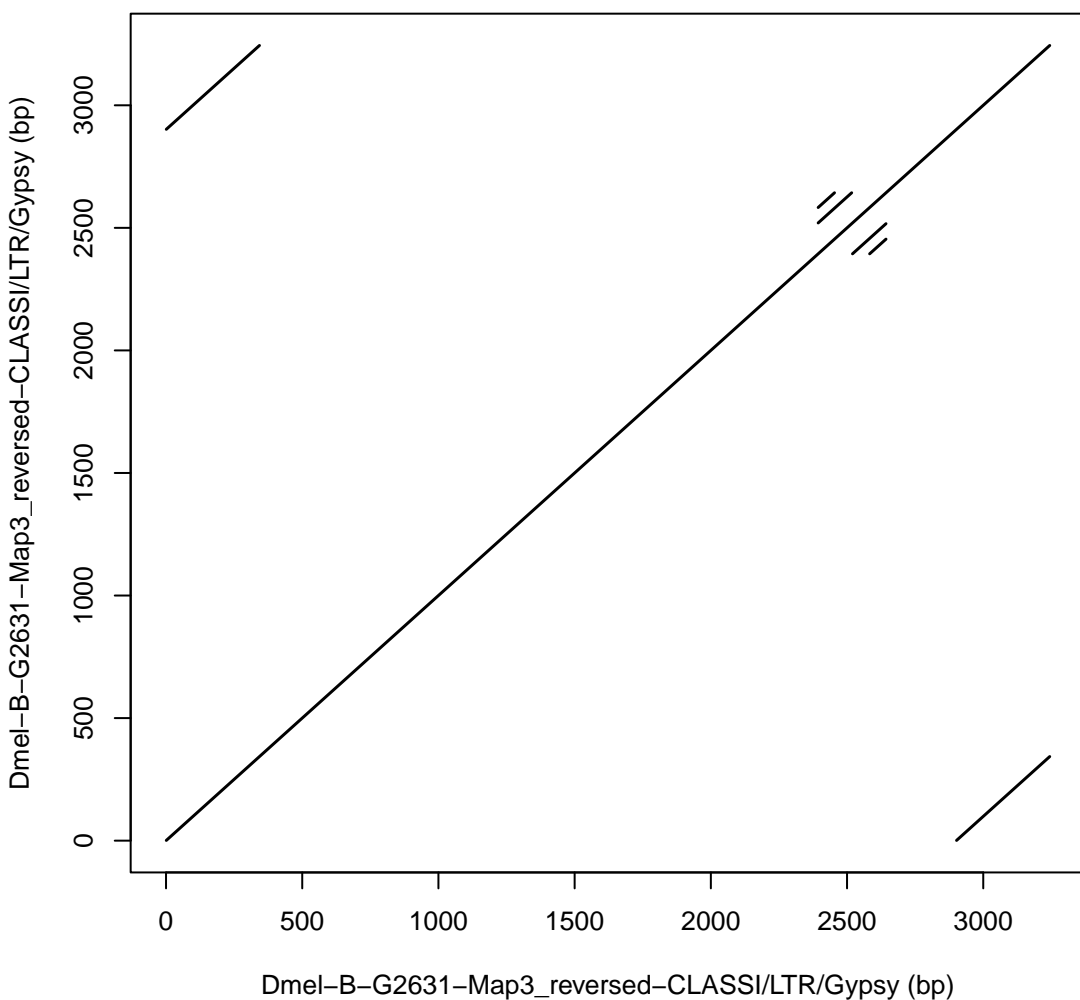
TE: Dmel-B-G2631-Map3_reversed-CLASSI/LTR/Gypsy
consensus size: 3244bp; fragments: 305; full length: 3 (>=2919.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

