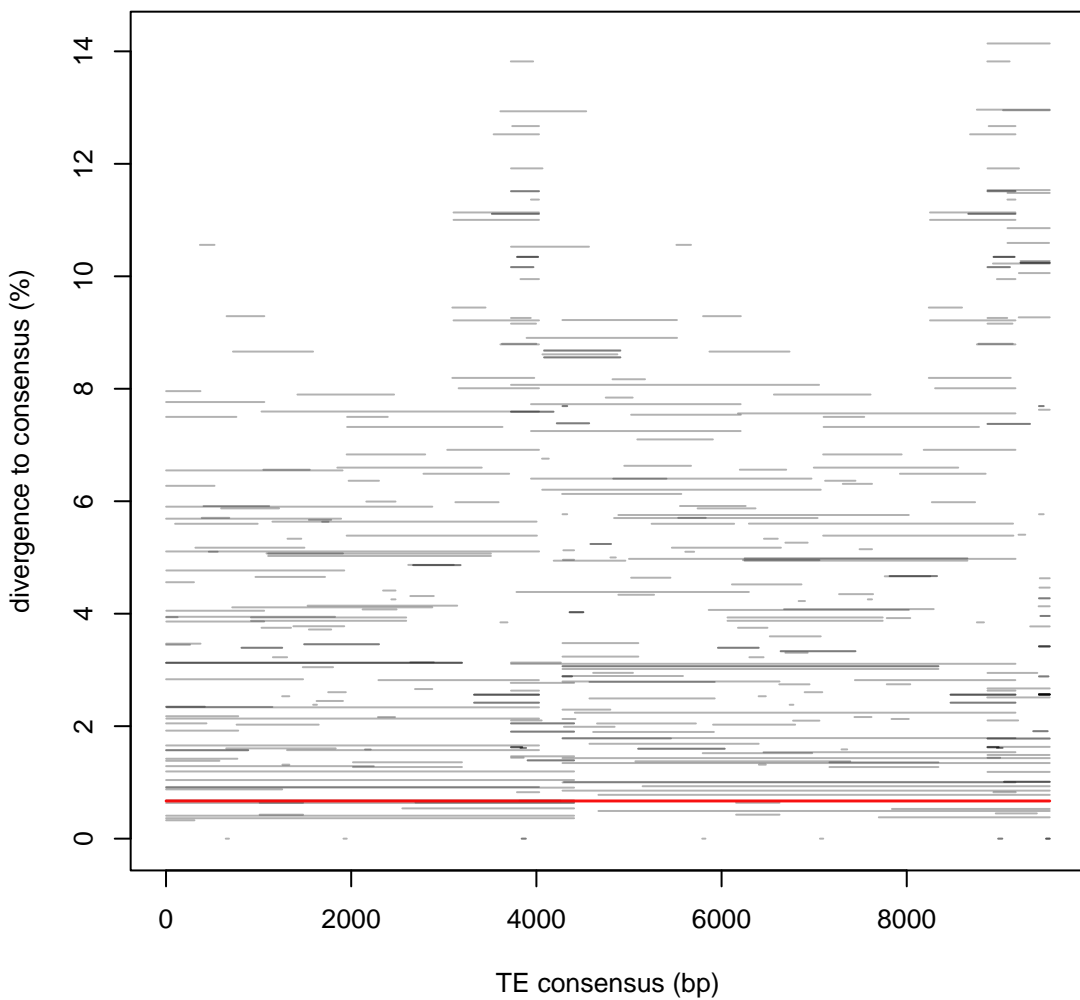
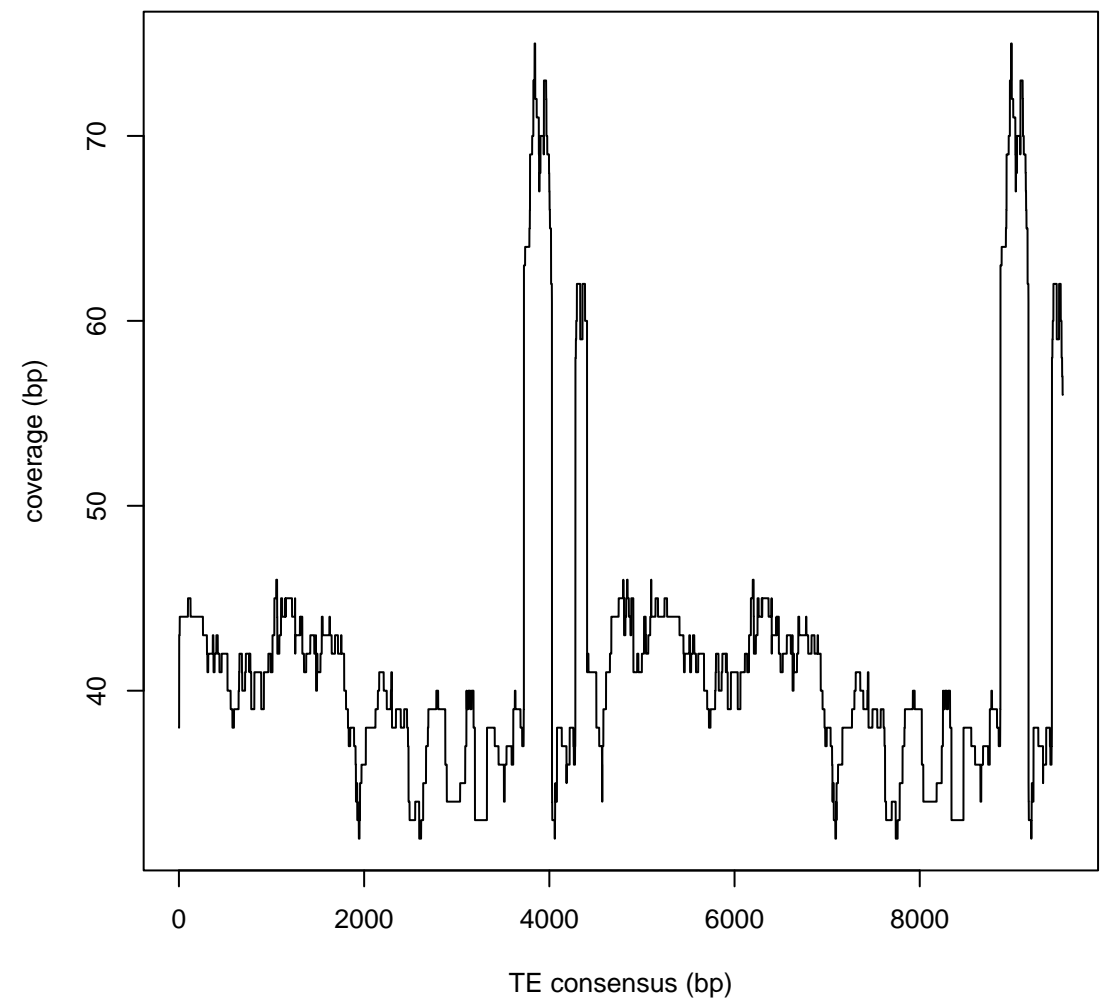


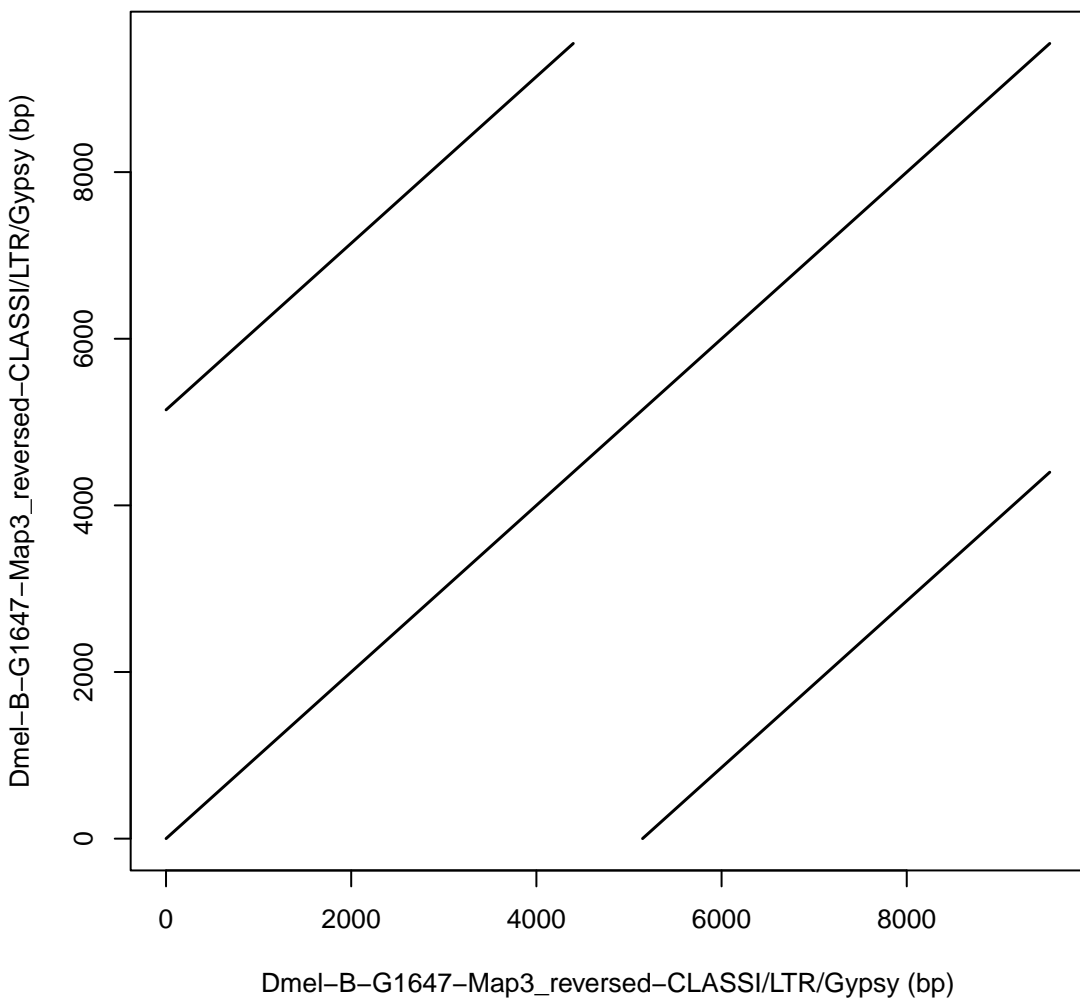
TE: Dmel-B-G1647-Map3_reversed-CLASSI/LTR/Gypsy
consensus size: 9544bp; fragments: 417; full length: 1 (>=8589.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

