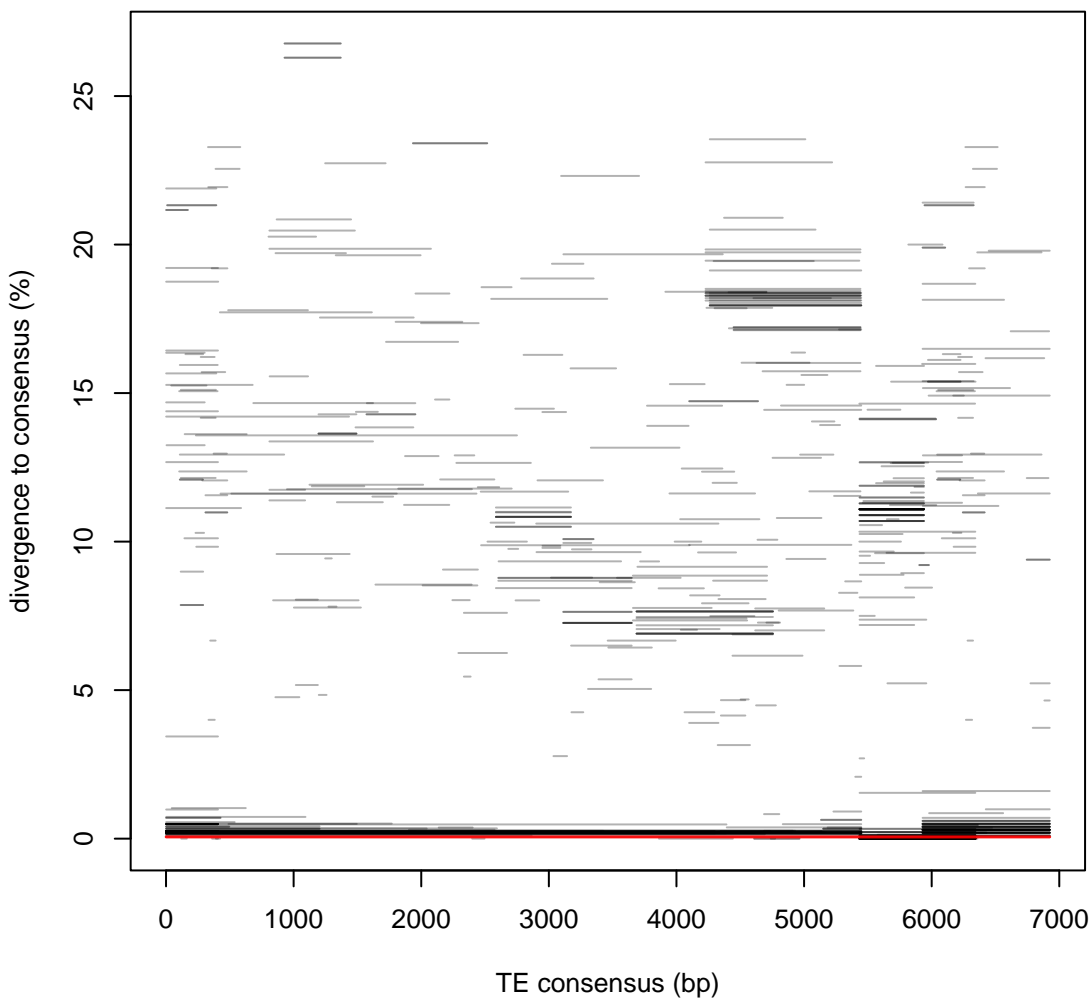
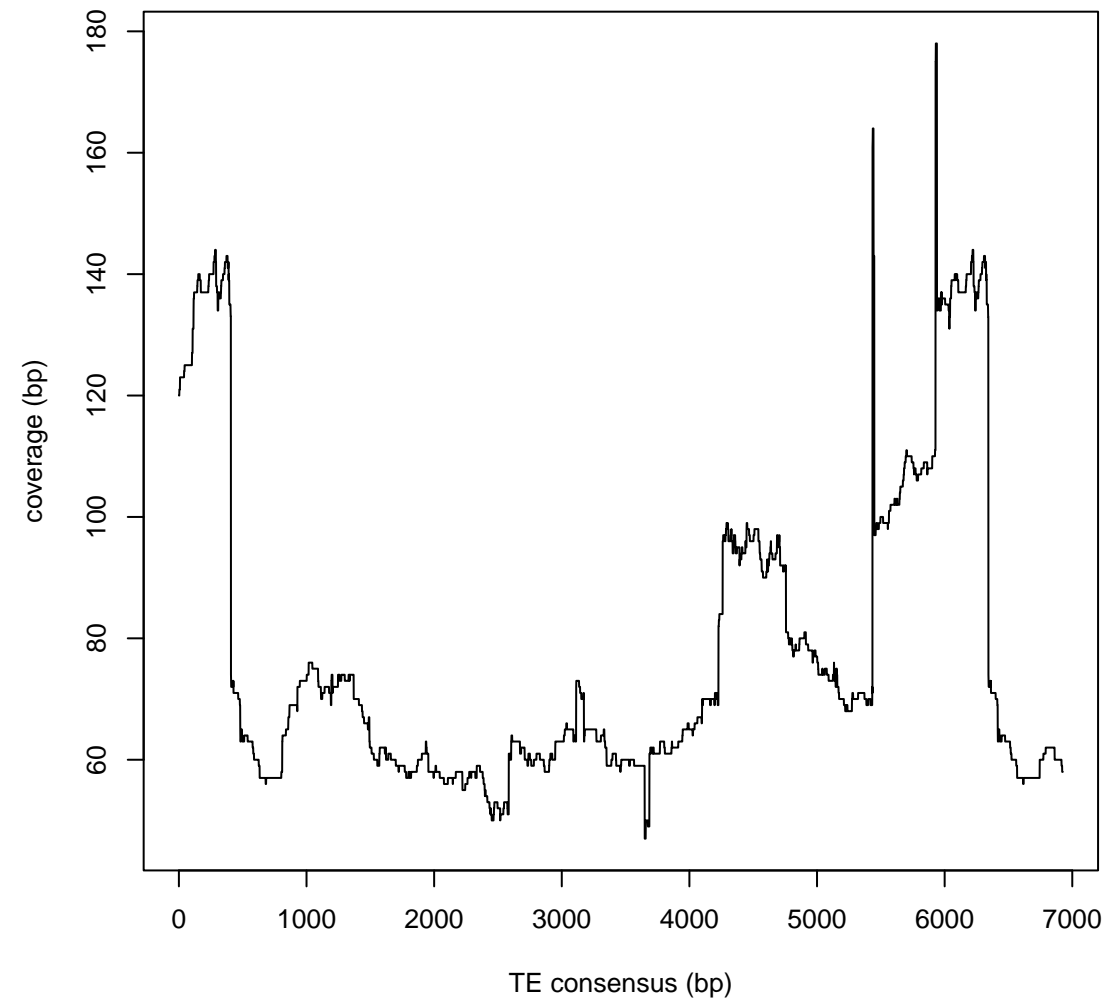


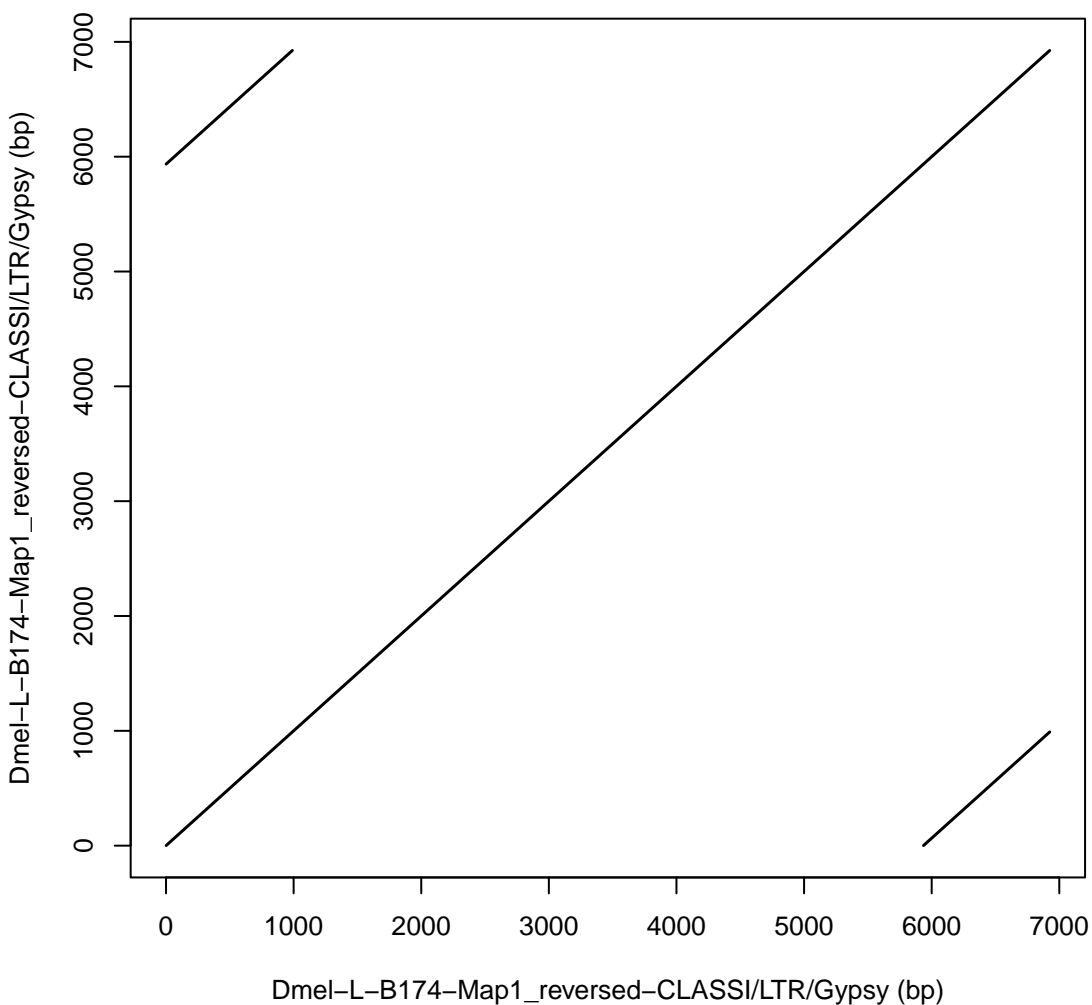
TE: Dmel-L-B174-Map1_reversed-CLASSI/LTR/Gypsy
consensus size: 6925bp; fragments: 650; full length: 1 (>=6232.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

