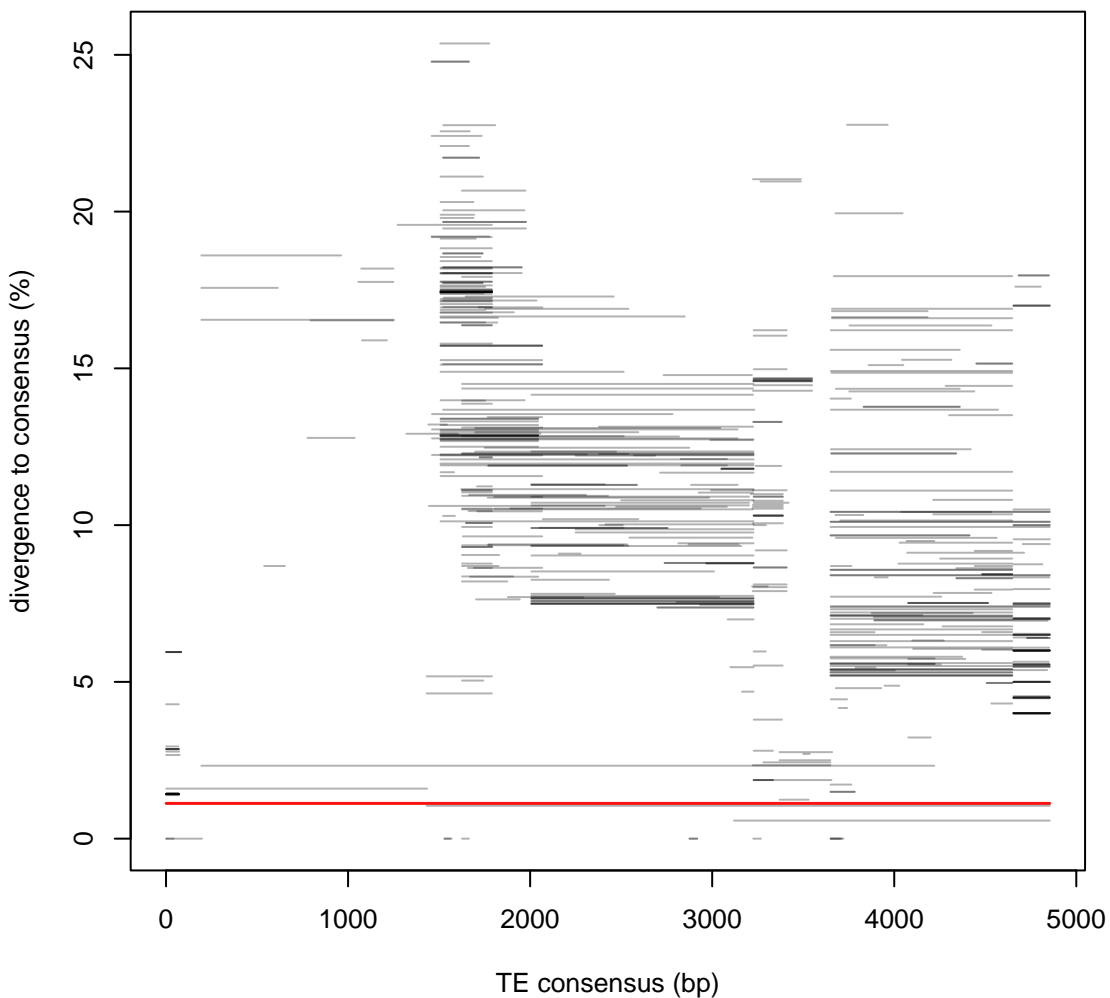
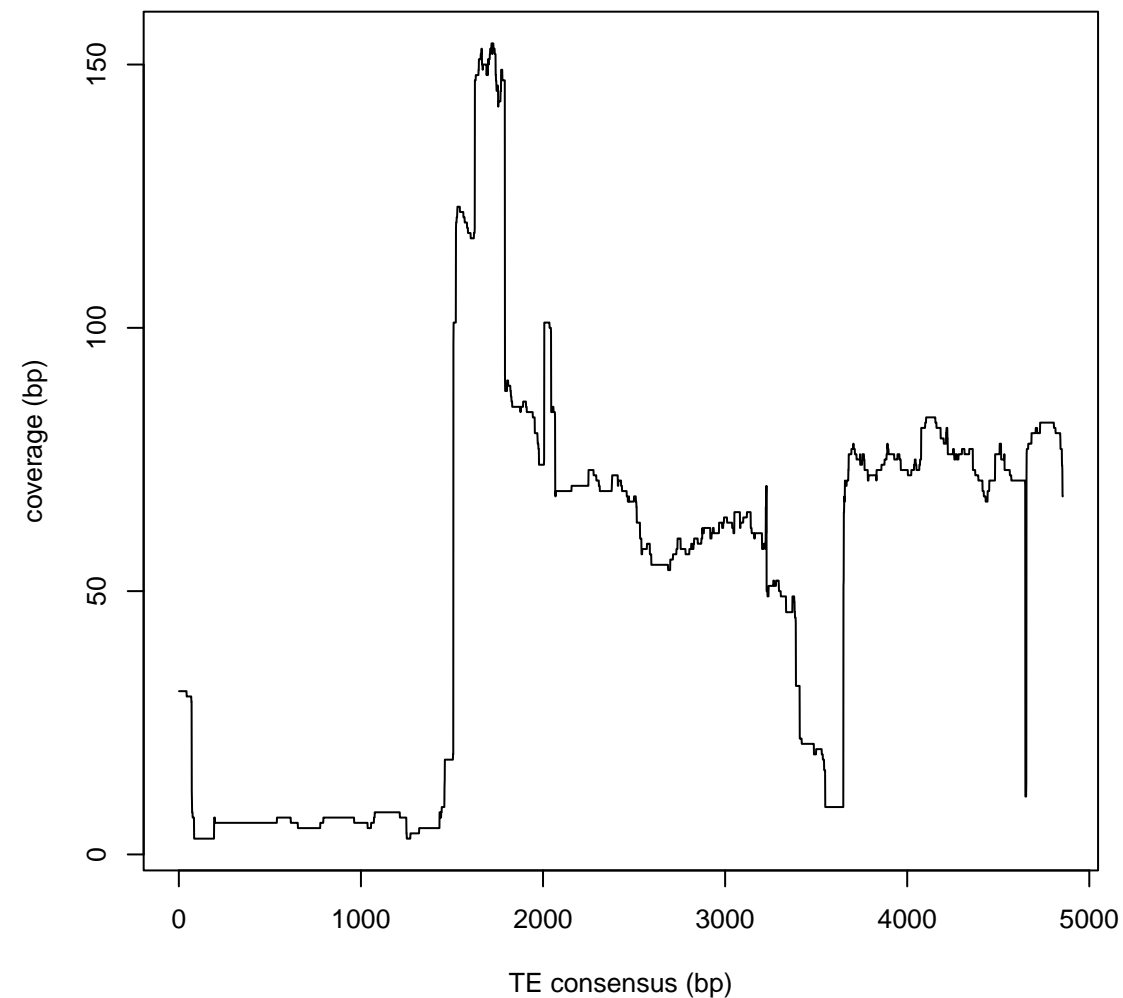


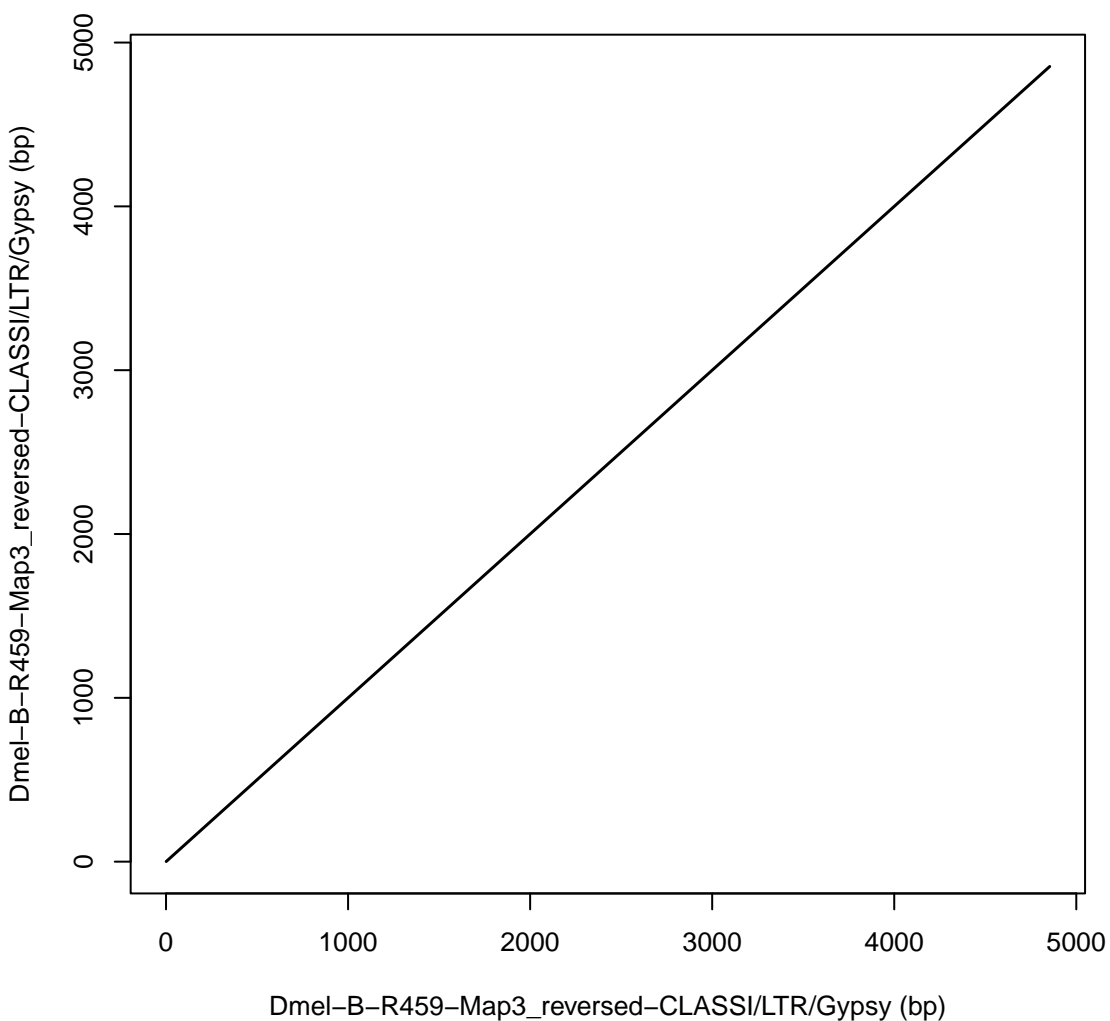
TE: Dmel-B-R459-Map3_reversed-CLASSI/LTR/Gypsy
consensus size: 4854bp; fragments: 540; full length: 1 (>=4368.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

