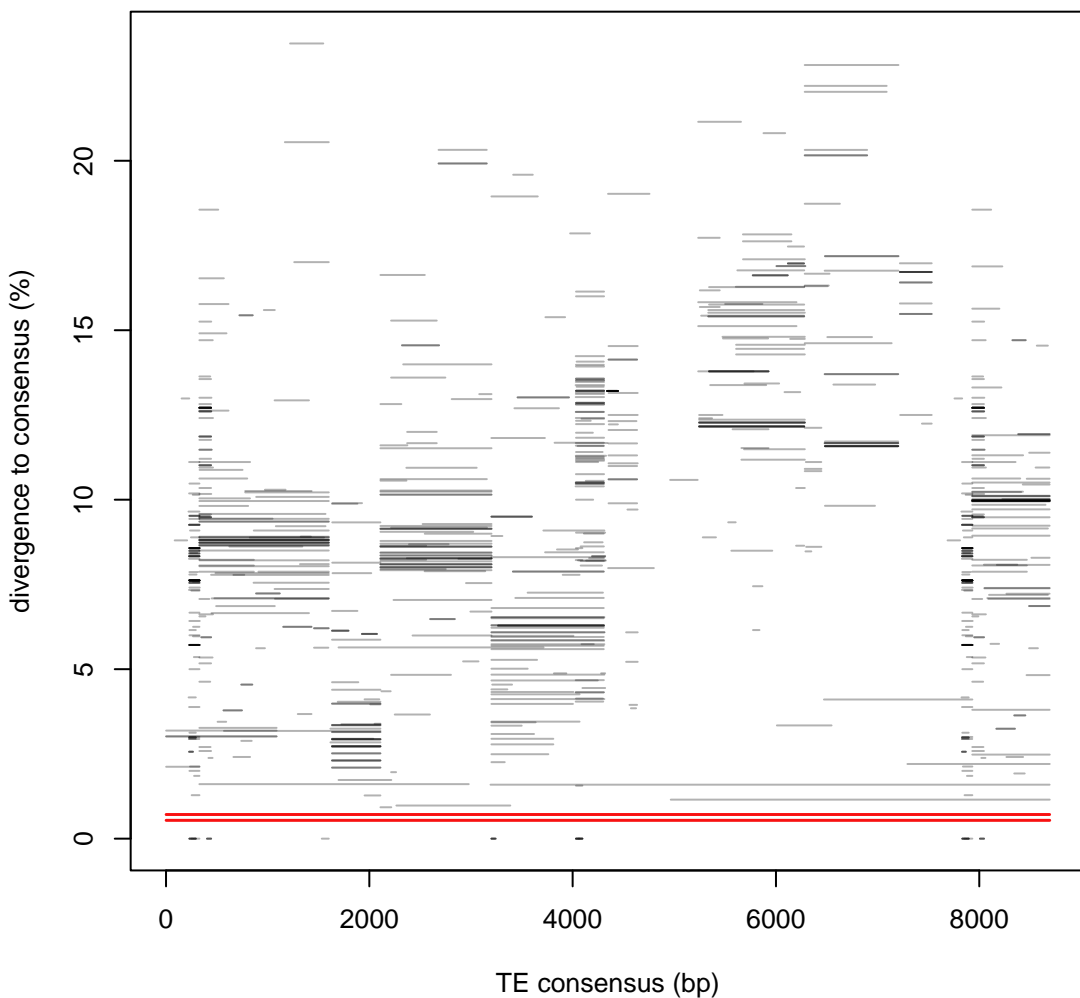
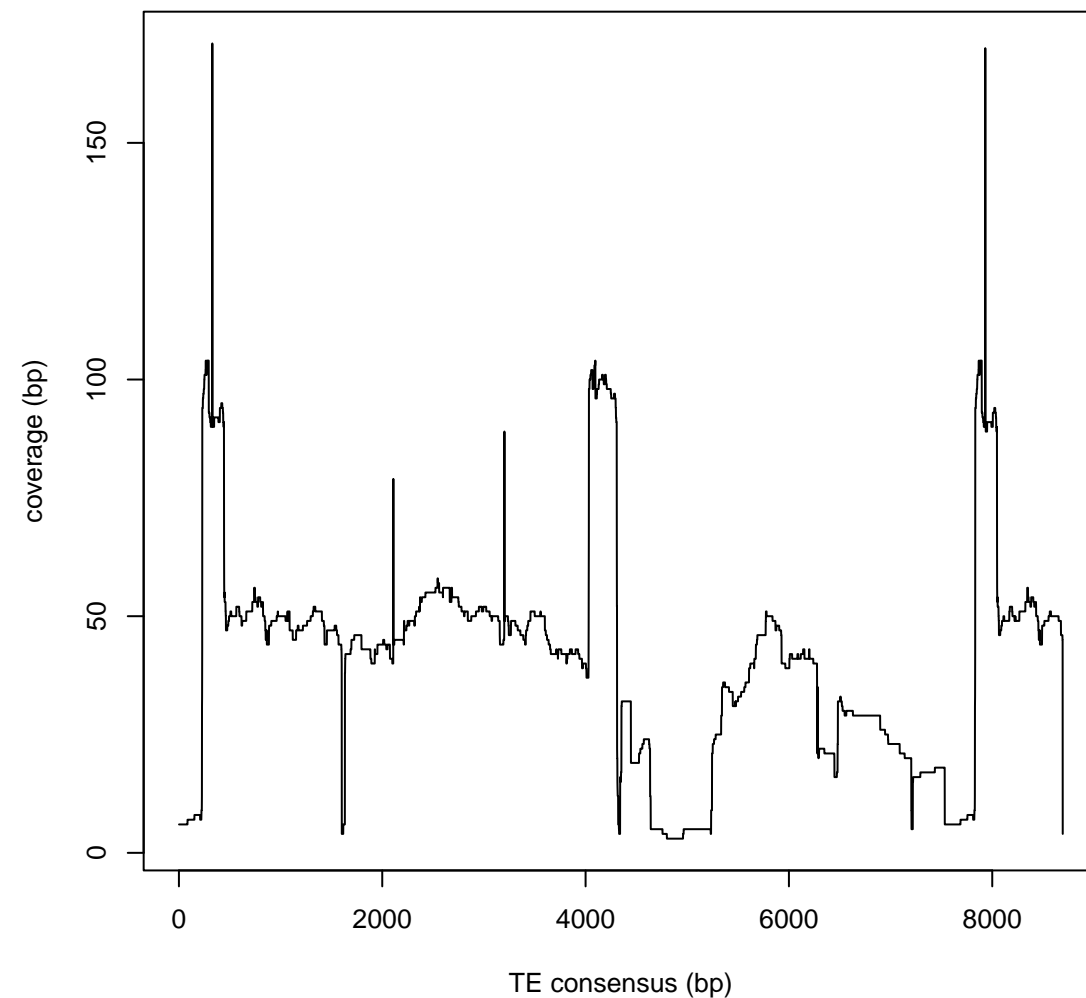


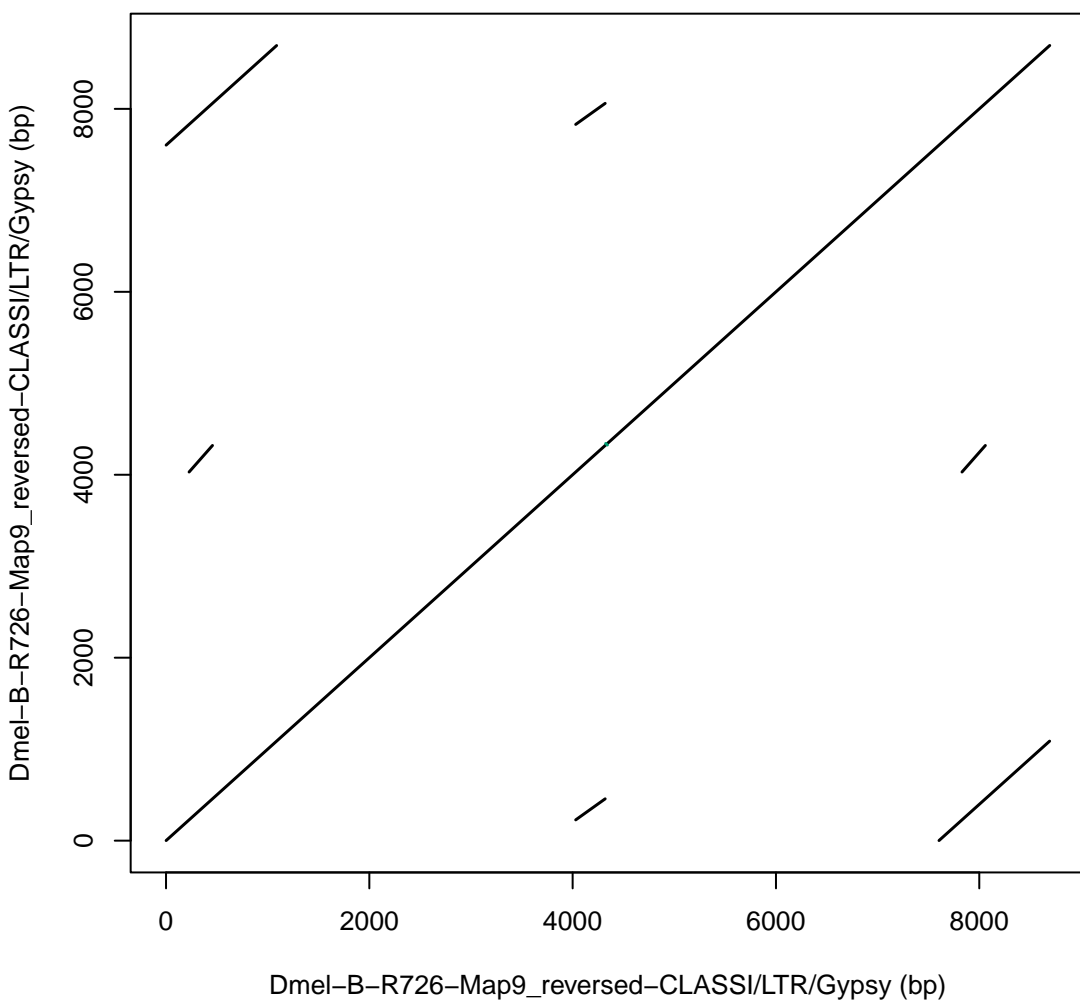
TE: Dmel-B-R726-Map9_reversed-CLASSI/LTR/Gypsy
consensus size: 8693bp; fragments: 887; full length: 2 (≥ 7823.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

