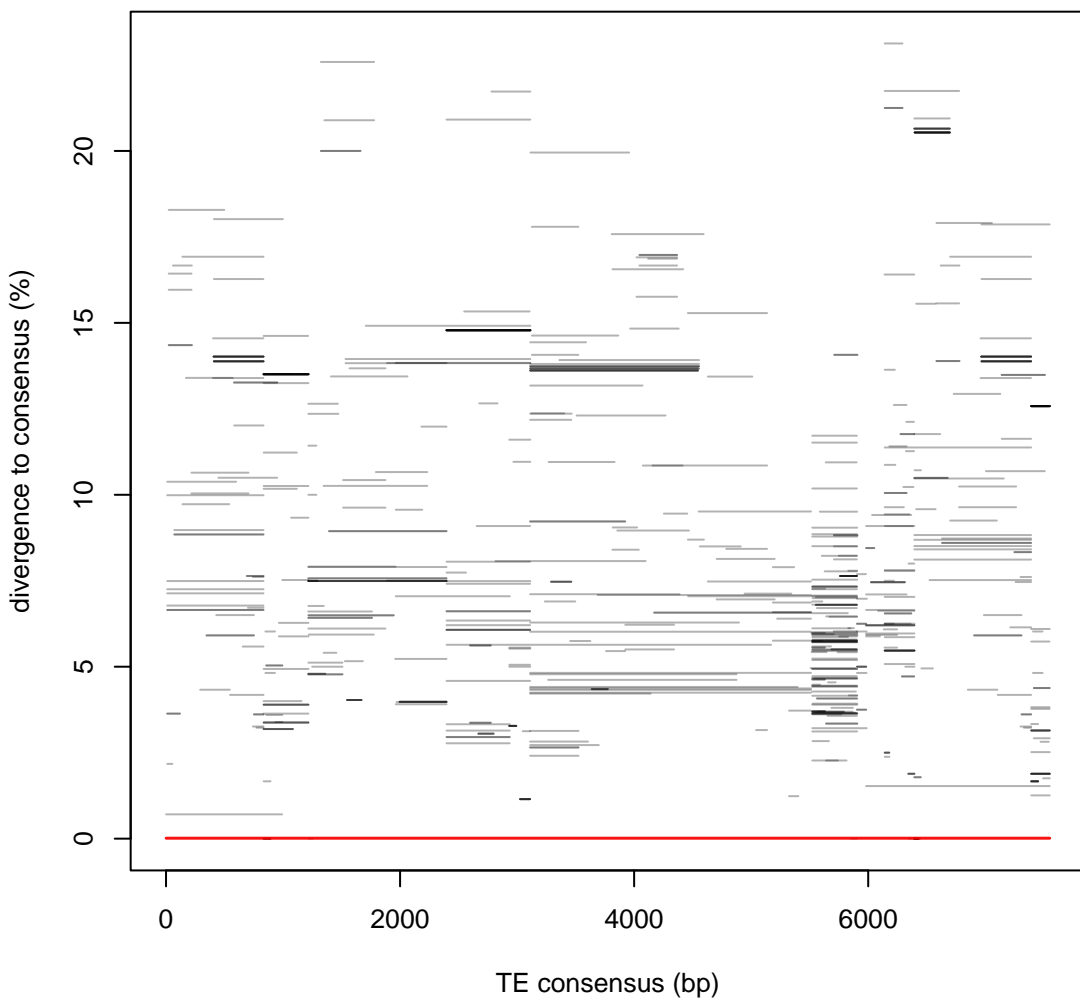
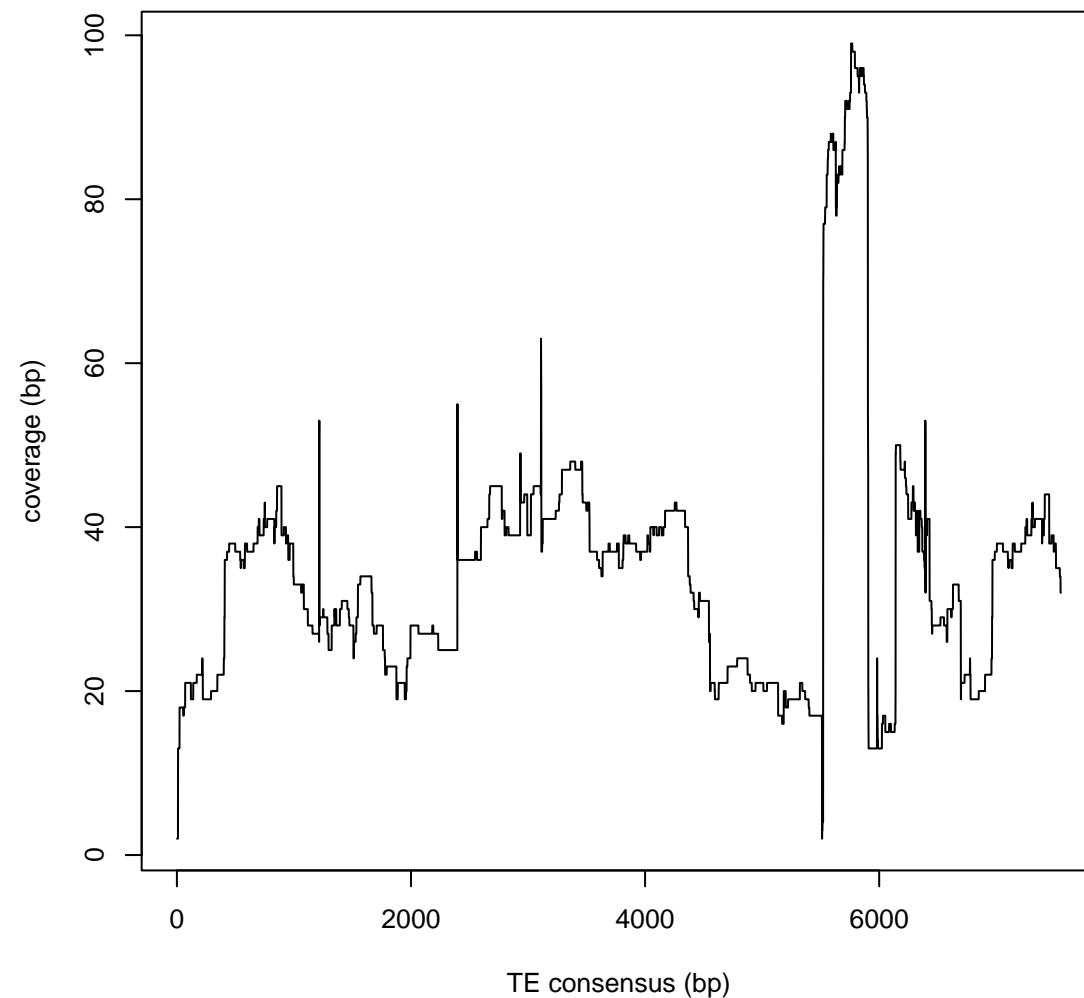


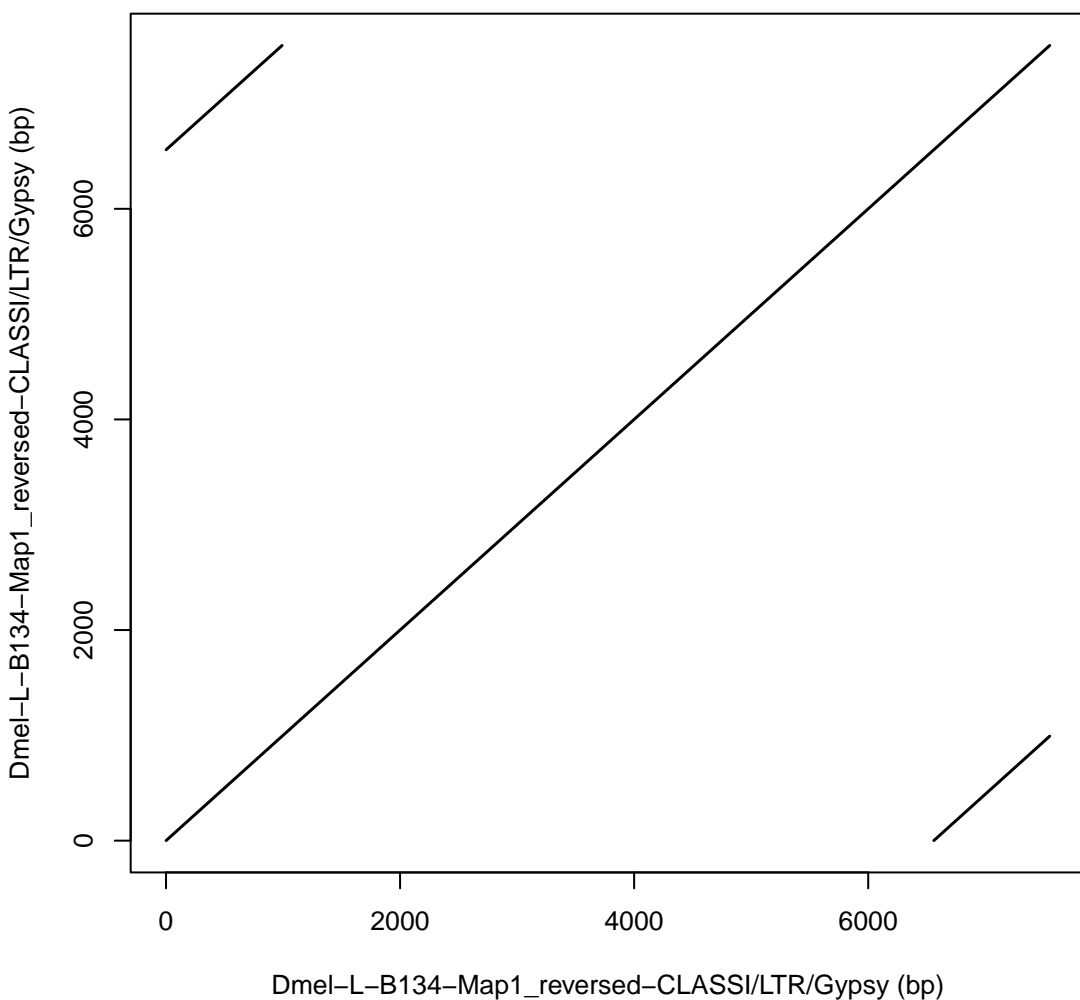
TE: Dmel-L-B134-Map1_reversed-CLASSI/LTR/Gypsy
consensus size: 7551bp; fragments: 638; full length: 1 (>=6795.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

