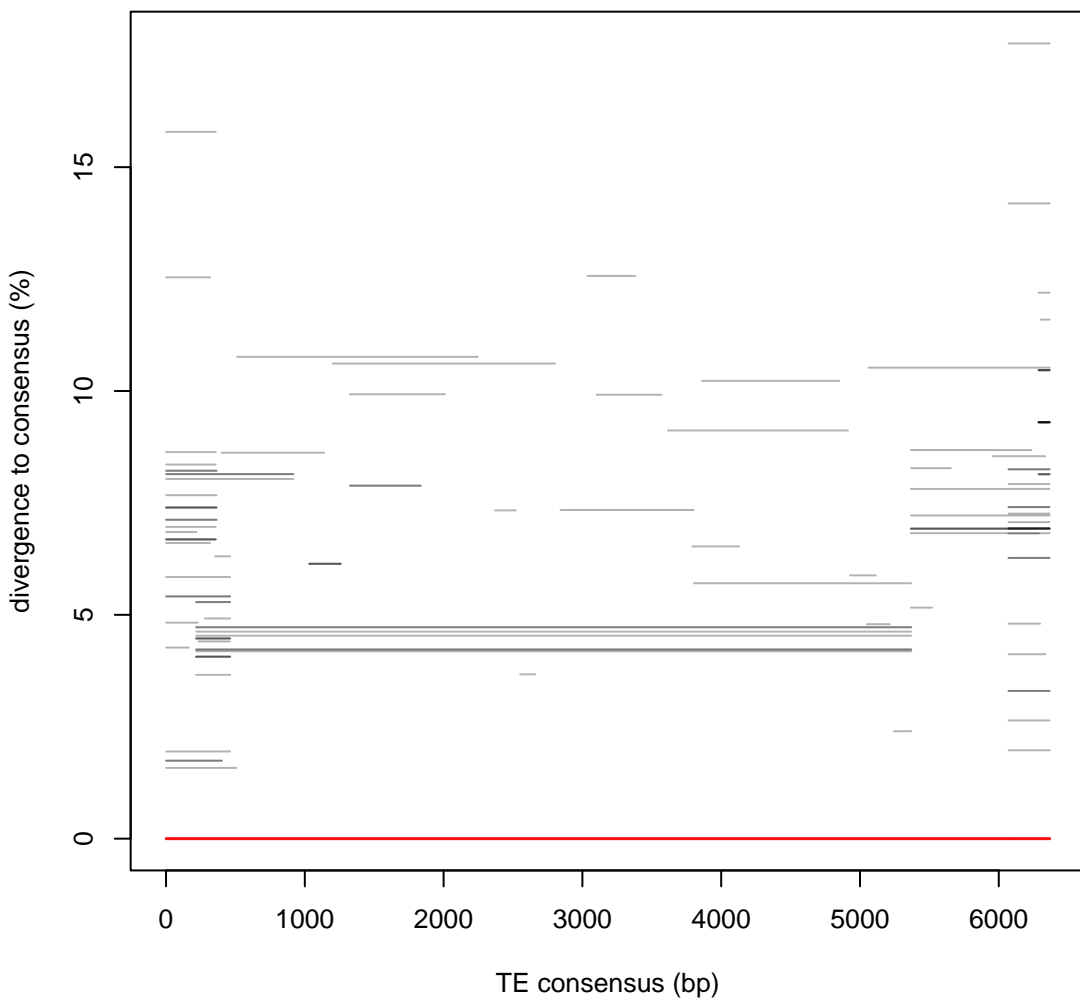
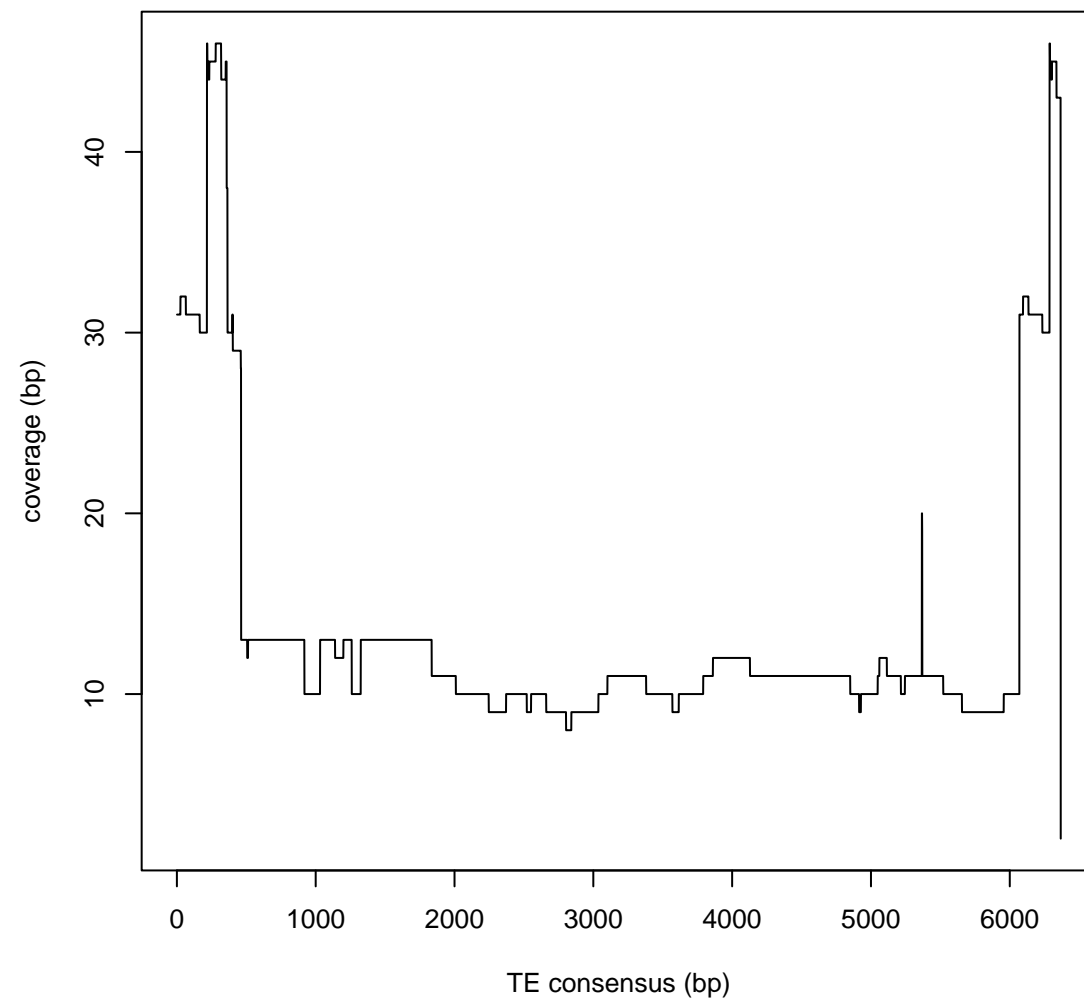


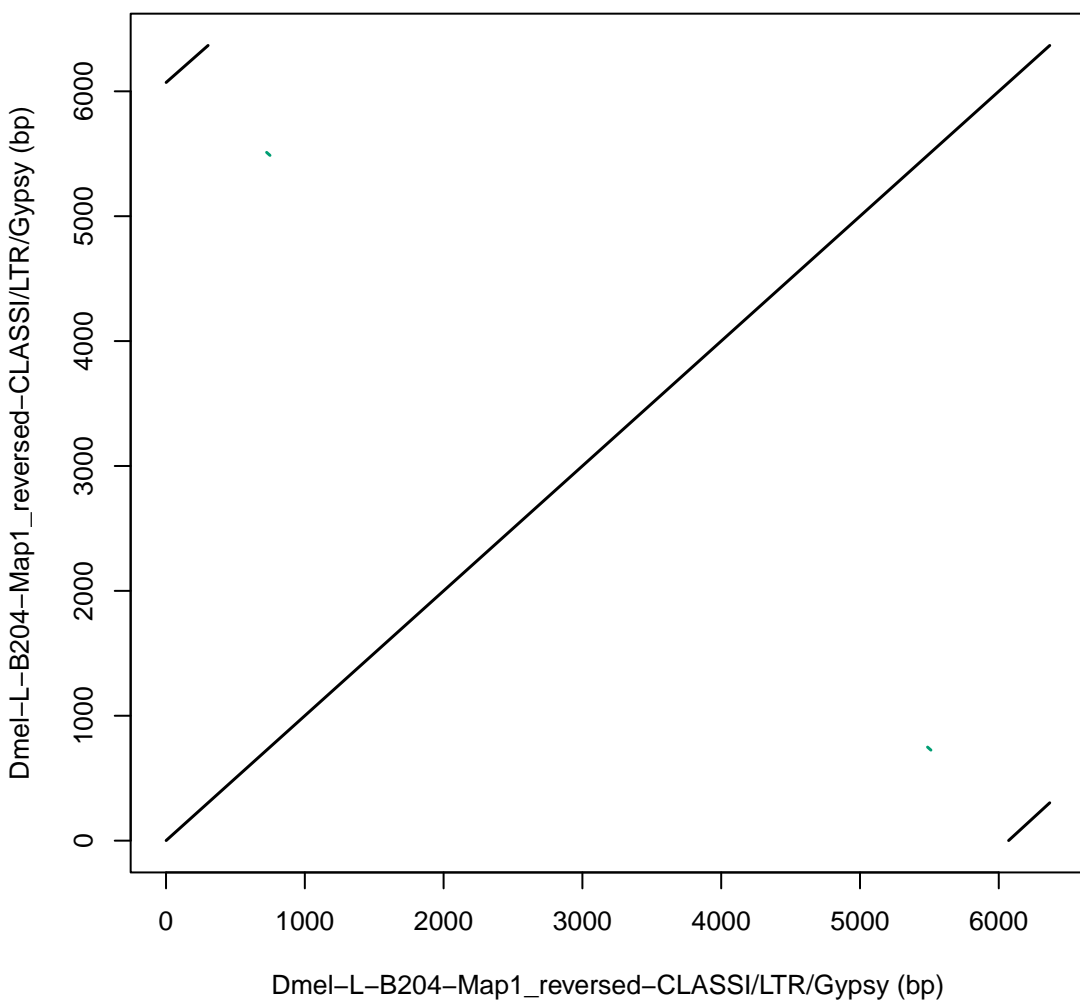
TE: Dmel-L-B204-Map1_reversed-CLASSI/LTR/Gypsy
consensus size: 6367bp; fragments: 122; full length: 1 (>=5730.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

