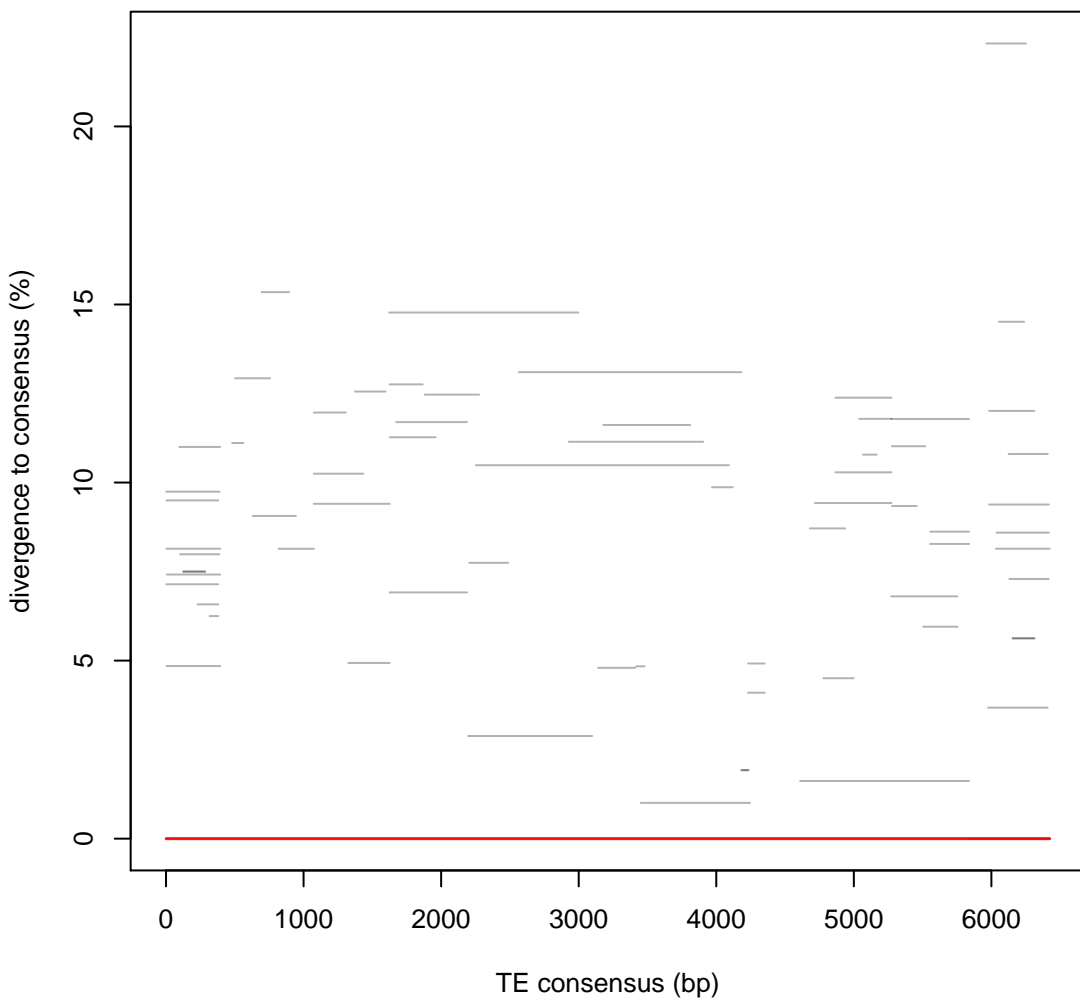
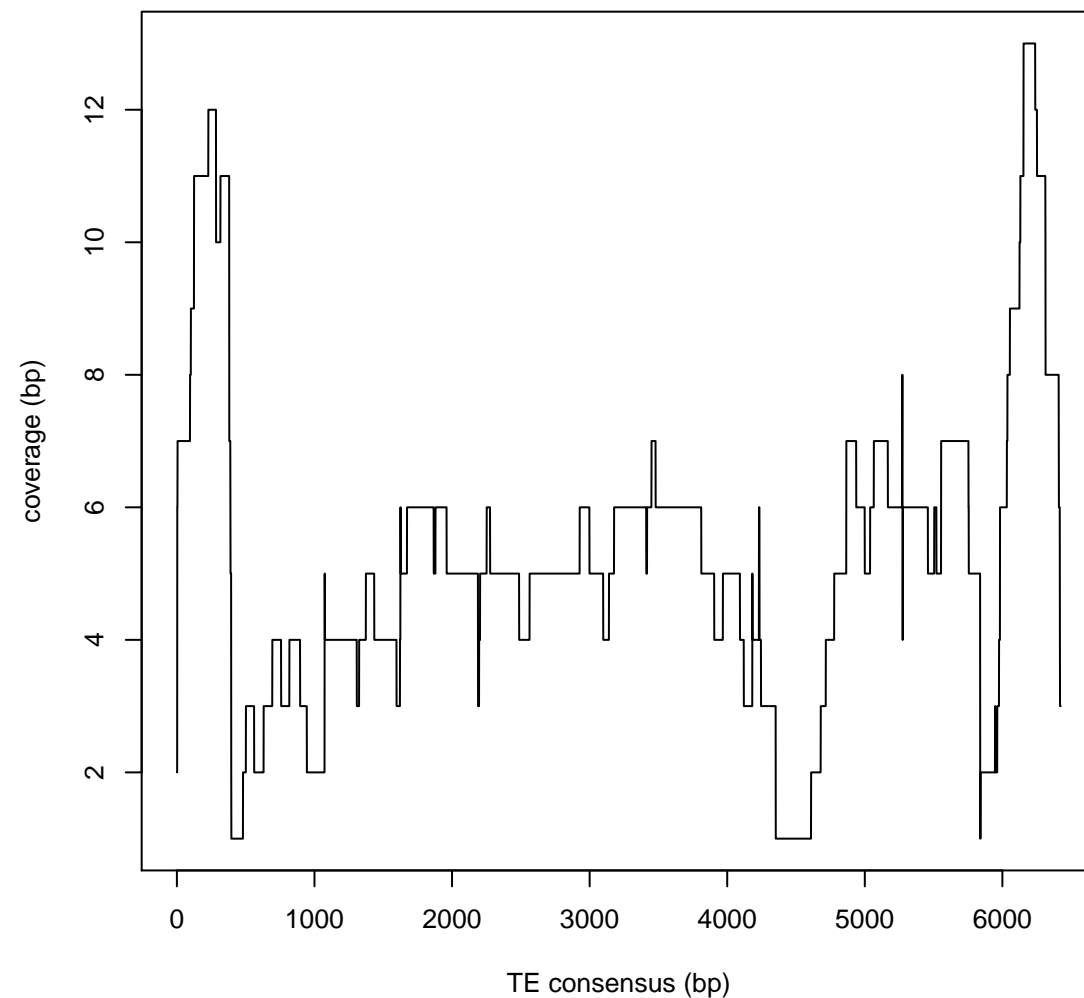


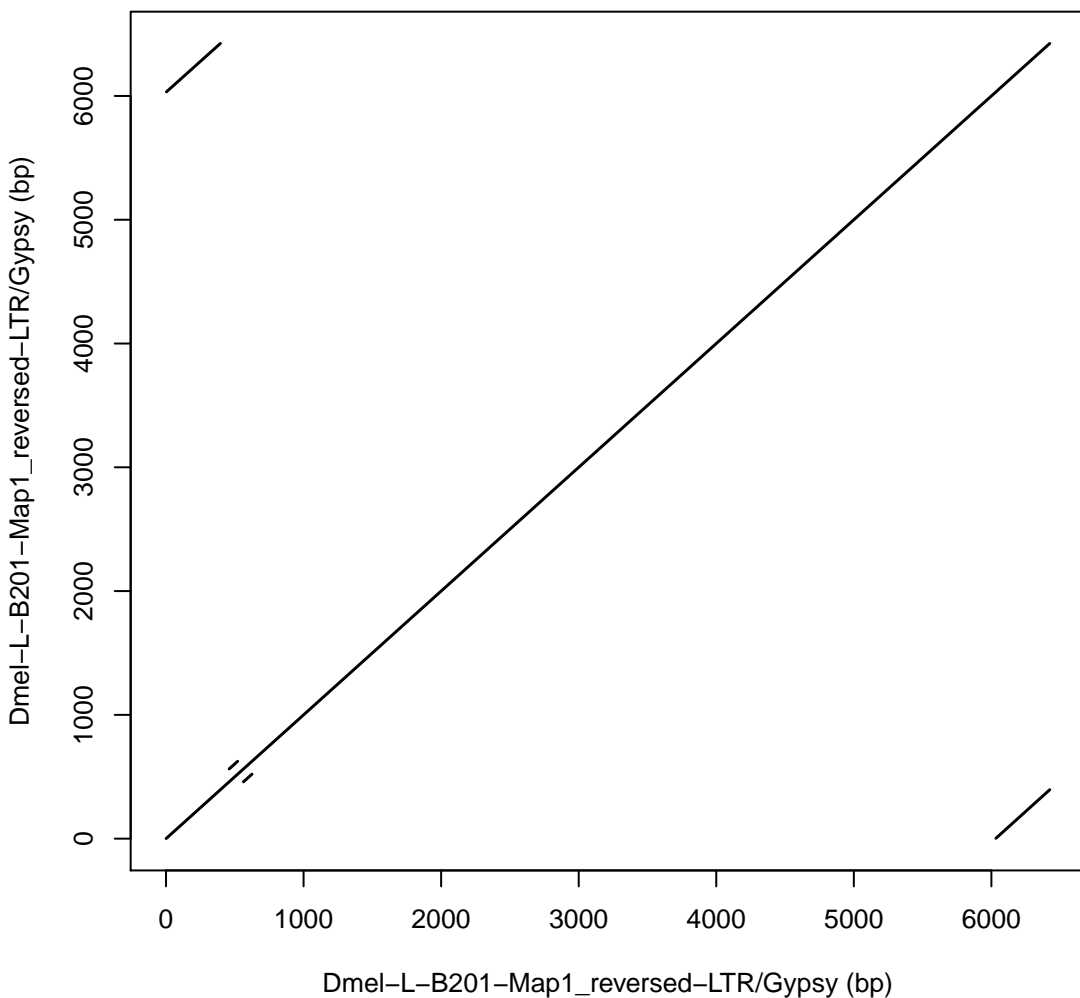
TE: Dmel-L-B201-Map1_reversed-LTR/Gypsy
consensus size: 6424bp; fragments: 71; full length: 1 (>=5781.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

