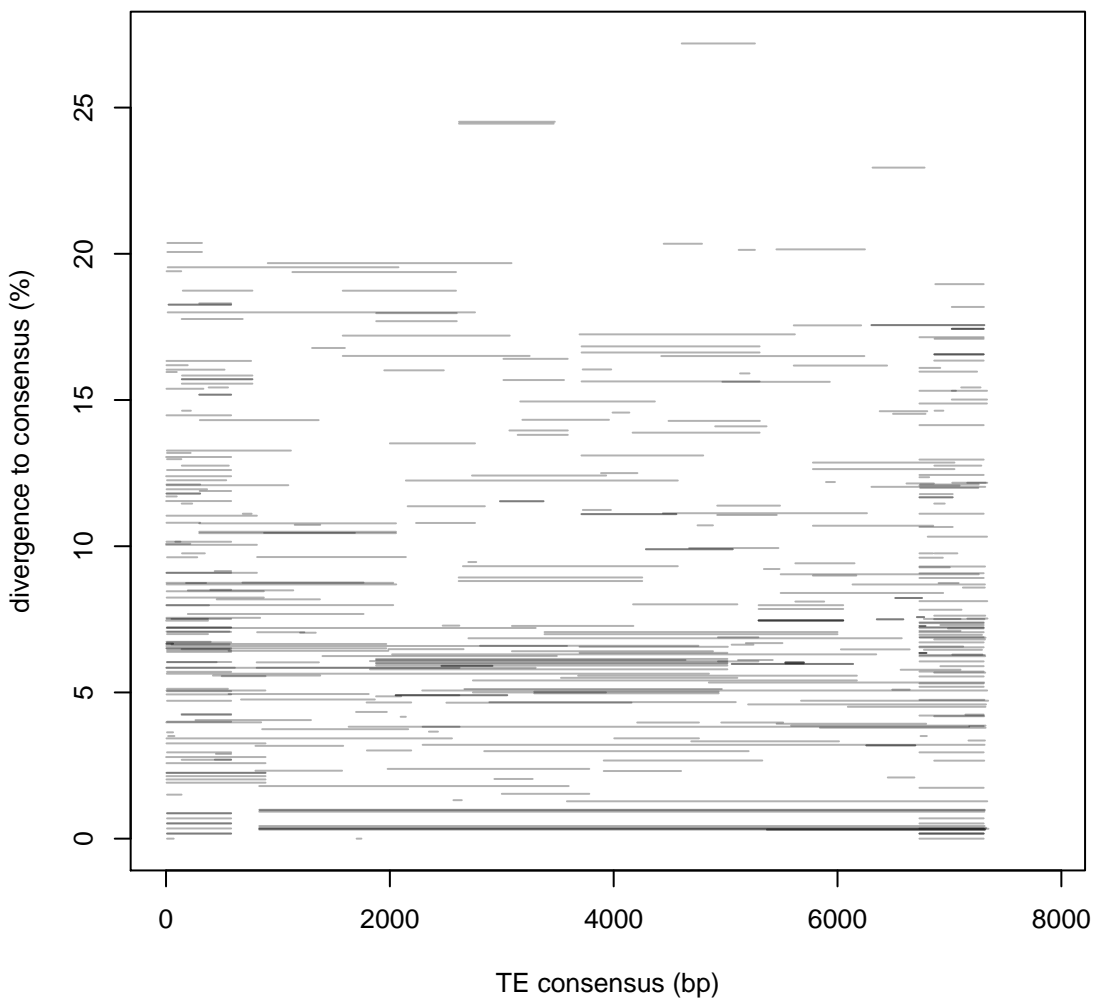
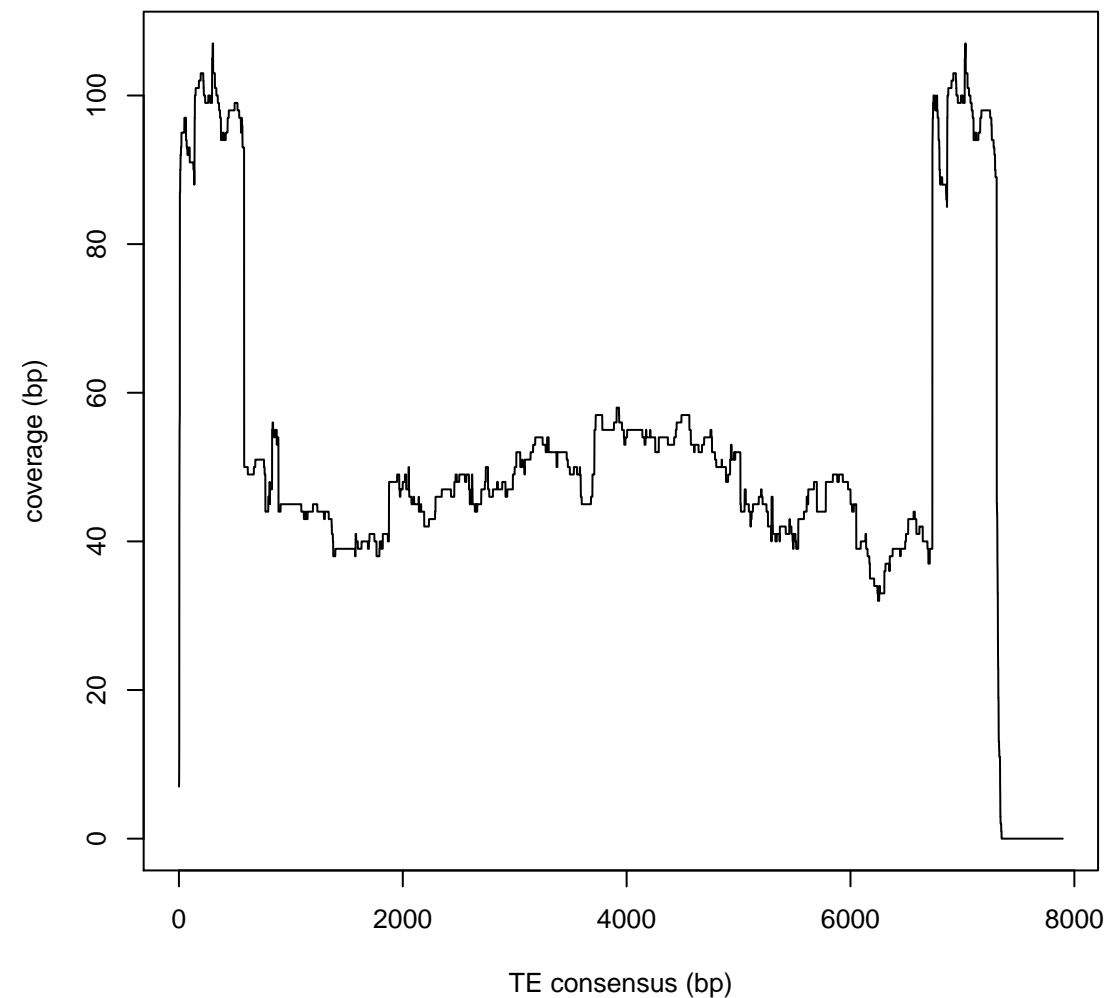


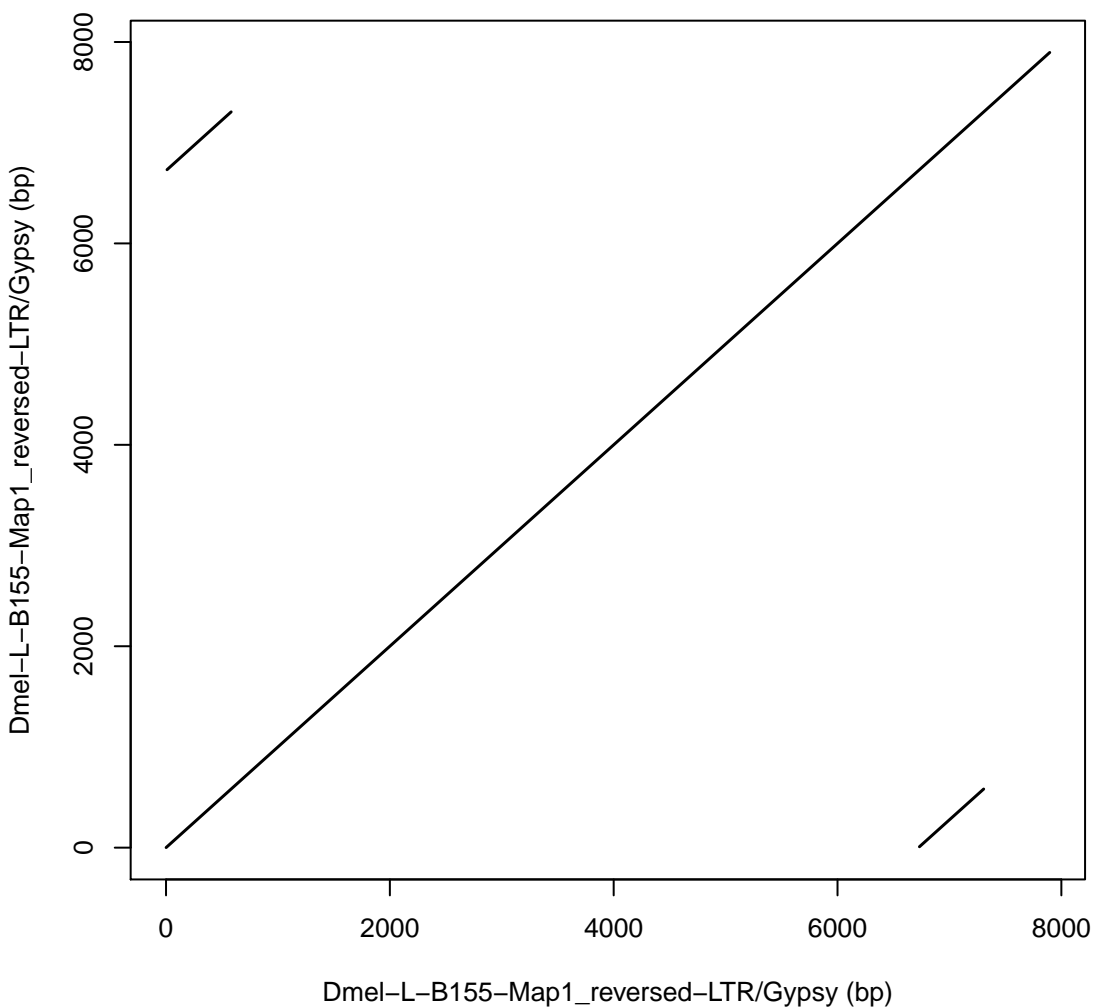
TE: Dmel-L-B155-Map1_reversed-LTR/Gypsy
consensus size: 7896bp; fragments: 426; full length: 0 (≥ 7106.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

