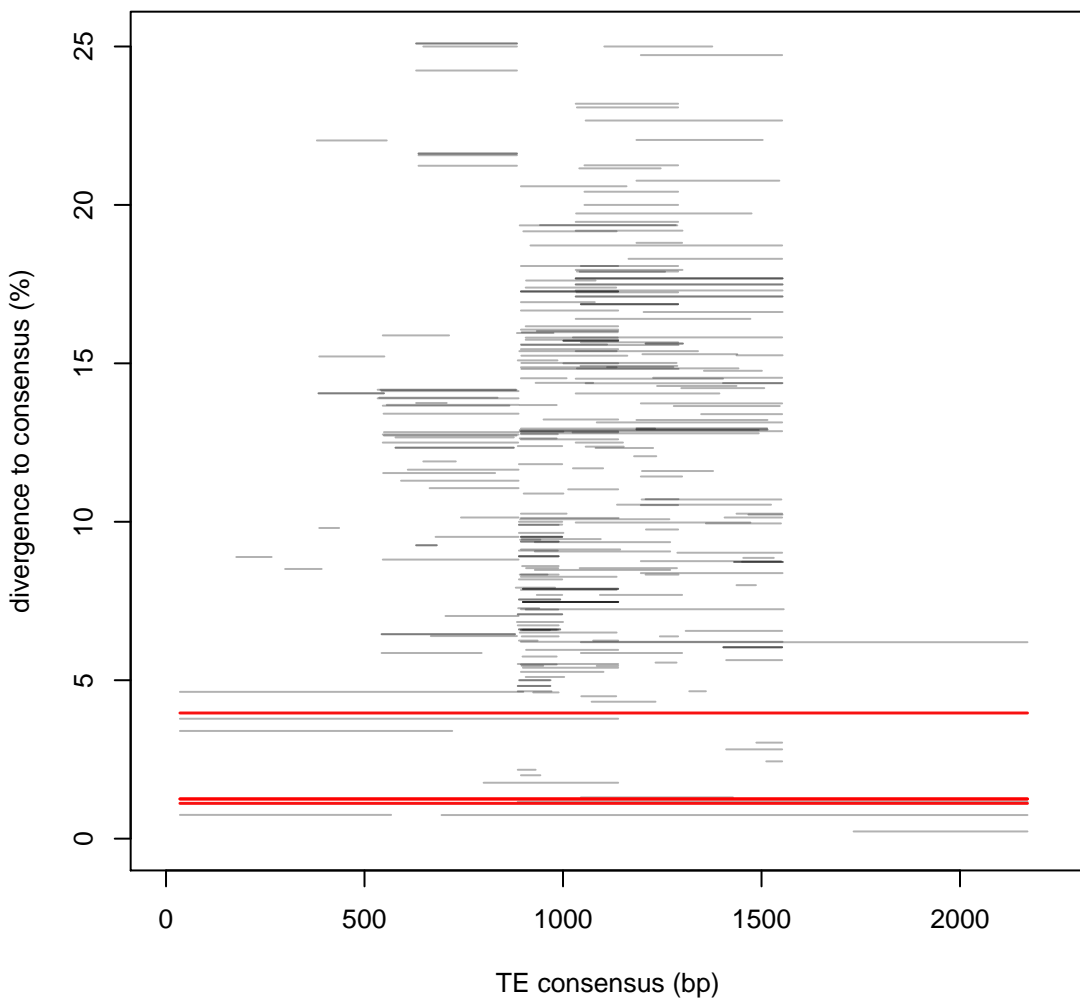
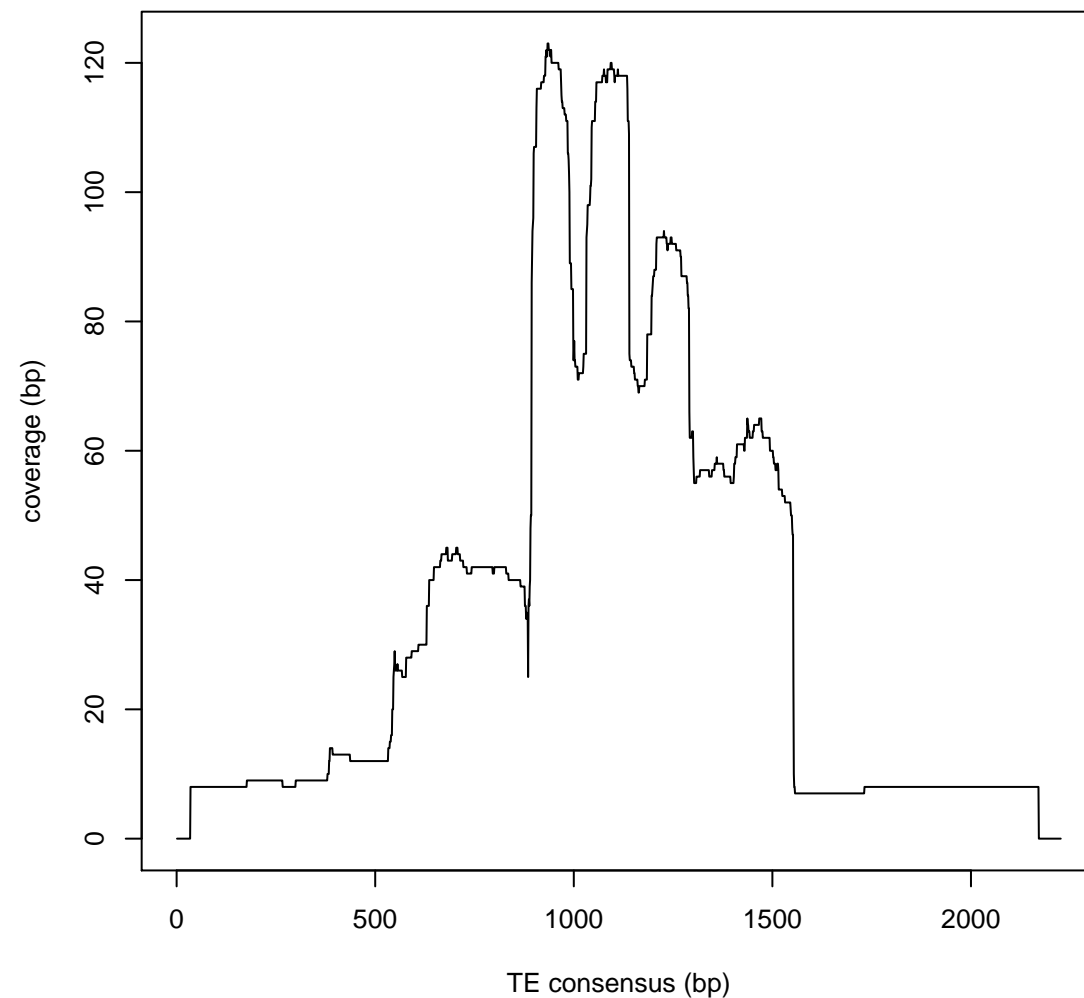


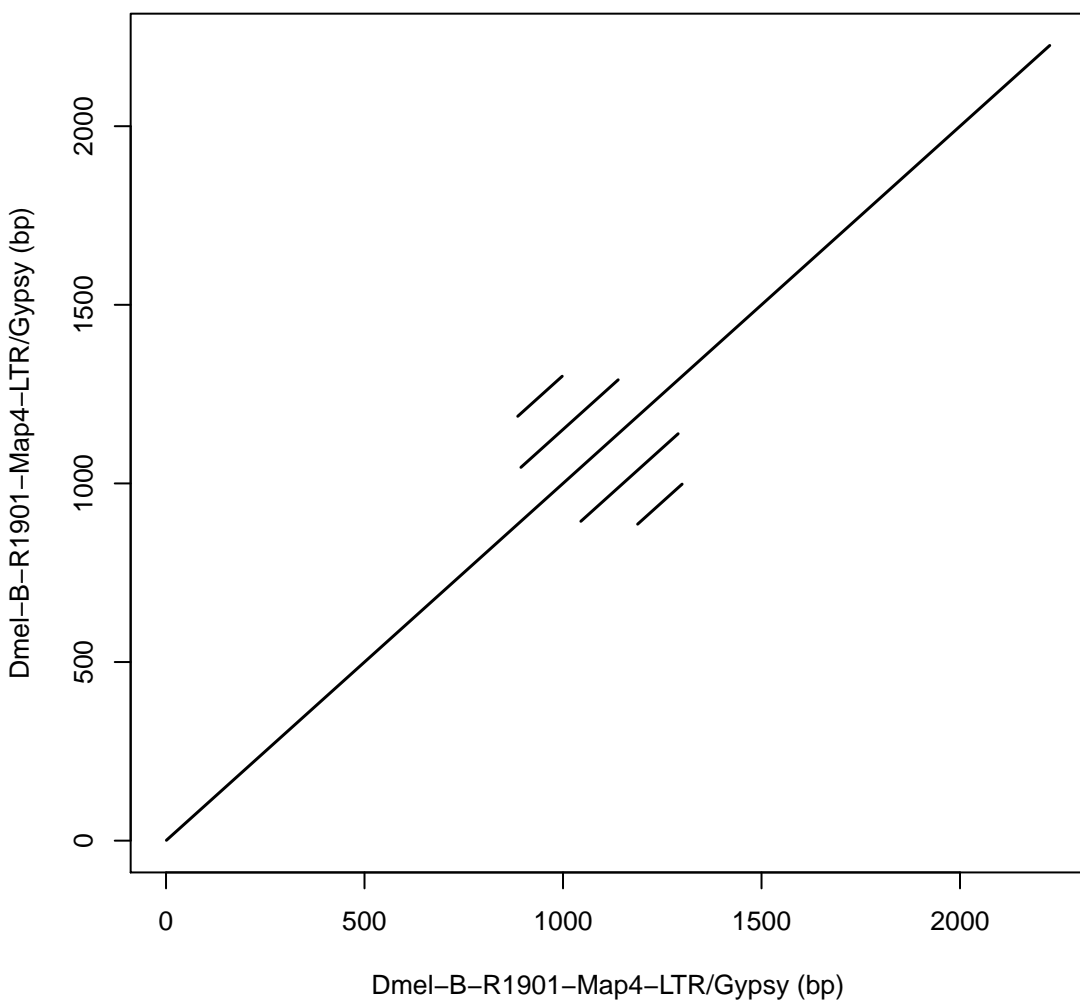
TE: Dmel-B-R1901-Map4-LTR/Gypsy
consensus size: 2226bp; fragments: 289; full length: 4 (≥ 2003.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

