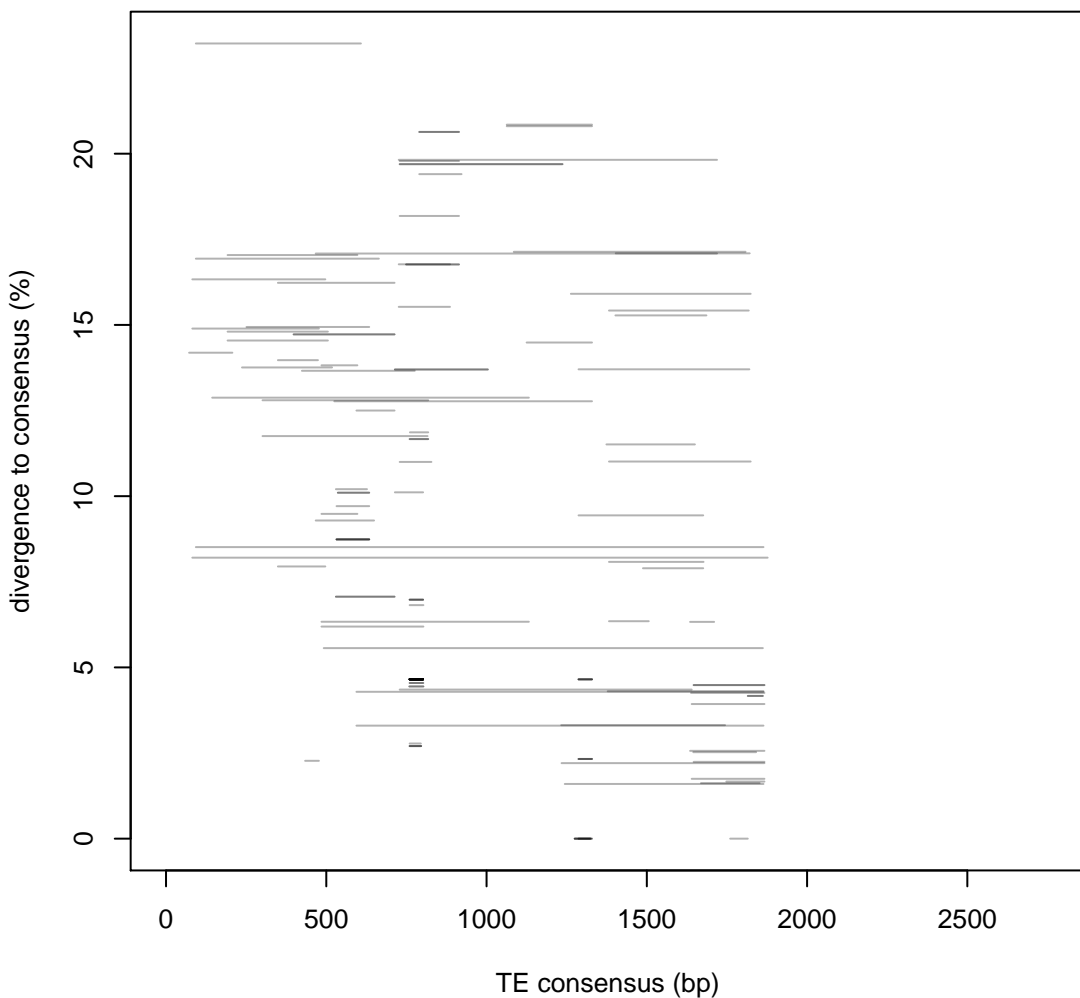
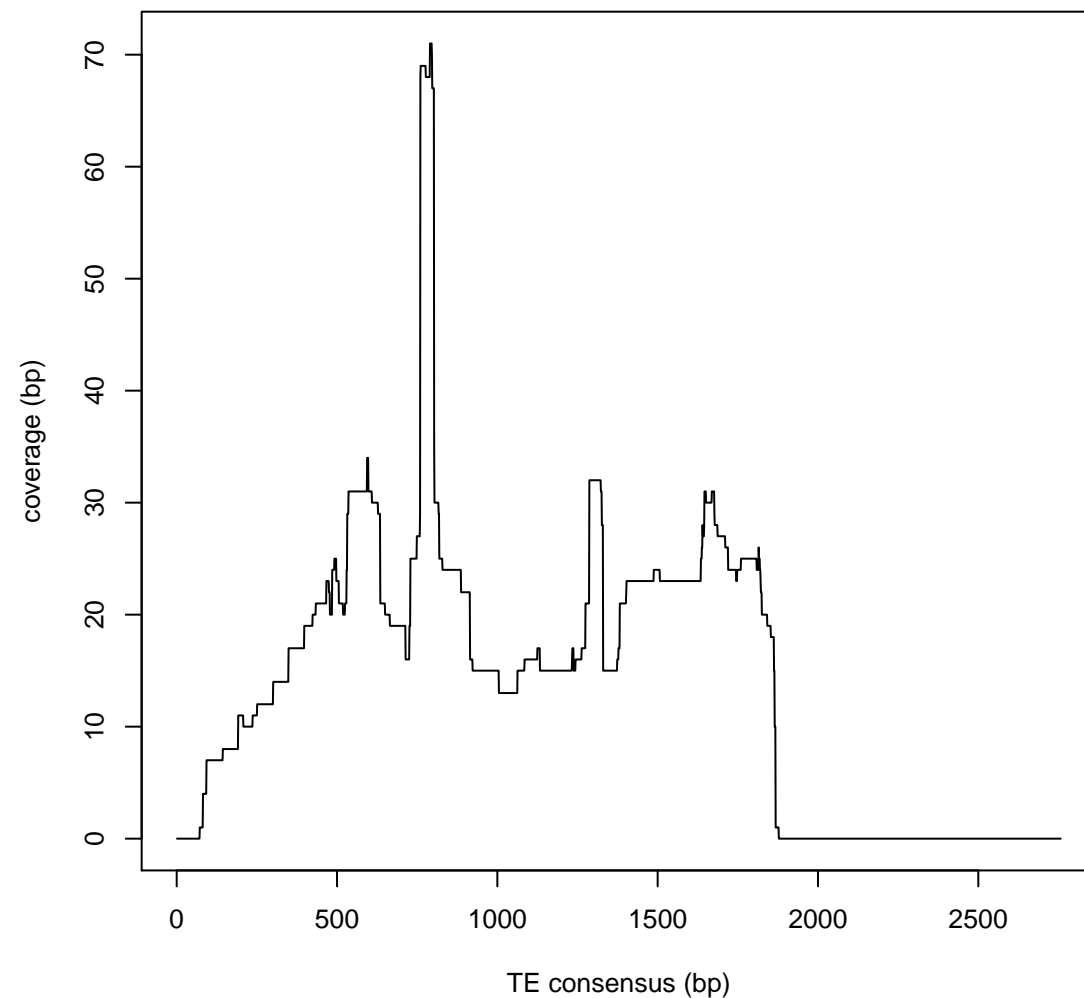


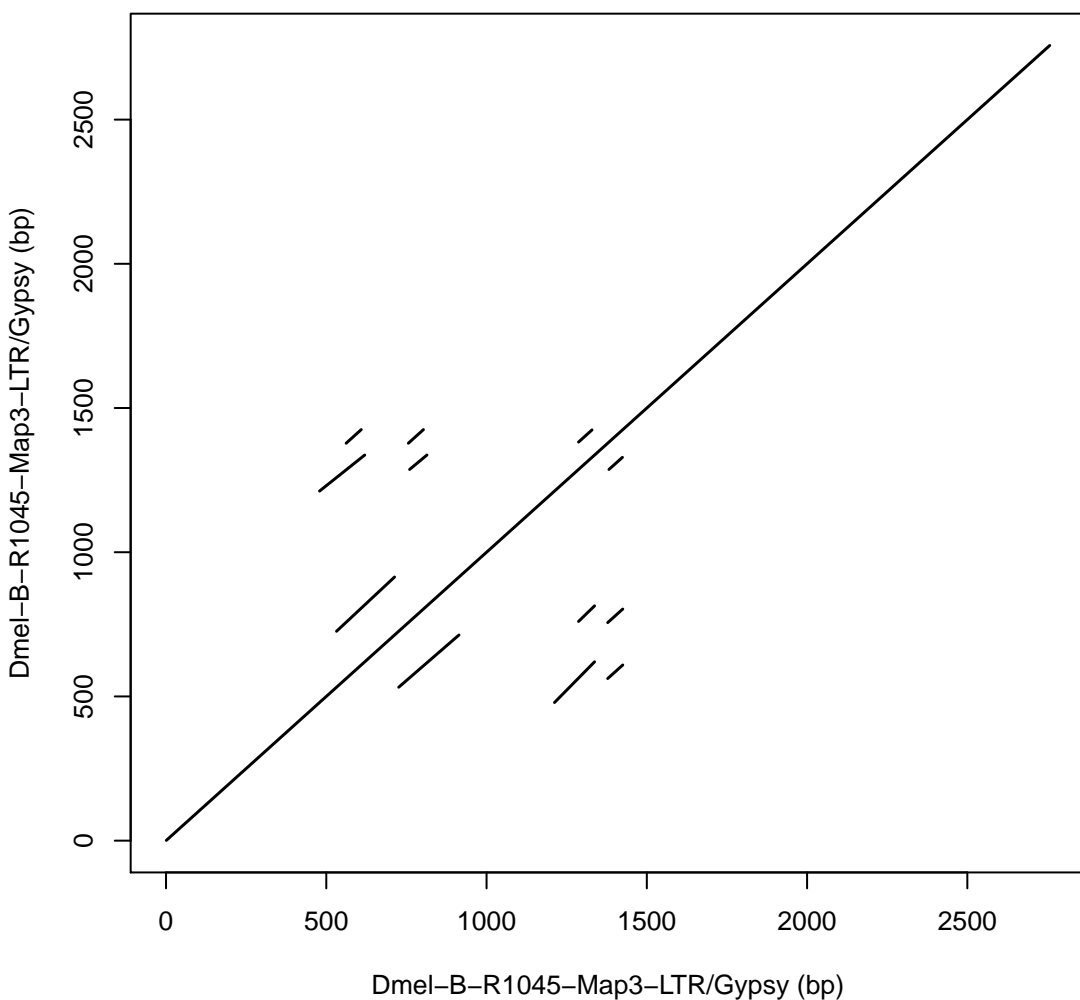
TE: Dmel-B-R1045-Map3-LTR/Gypsy
consensus size: 2757bp; fragments: 148; full length: 0 (≥ 2481.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

