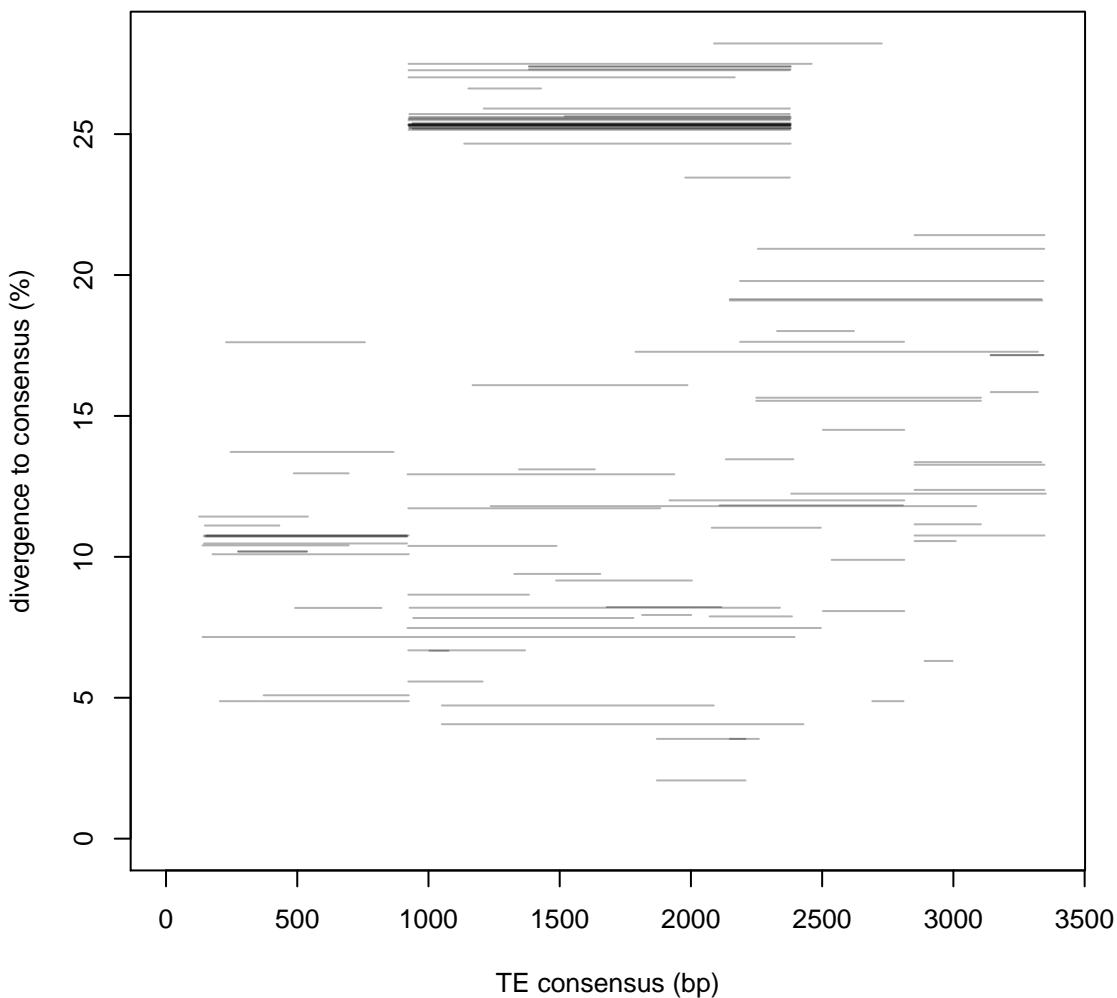
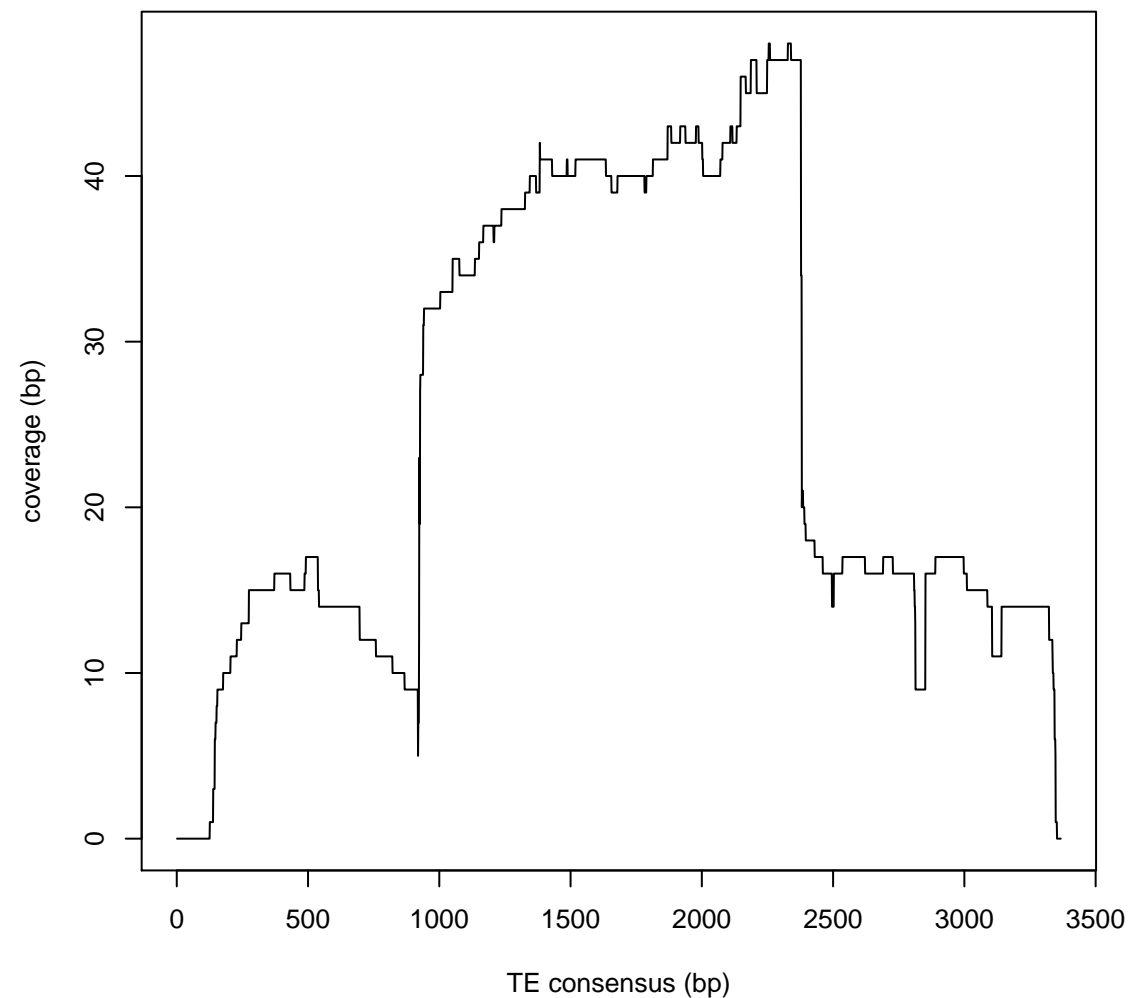


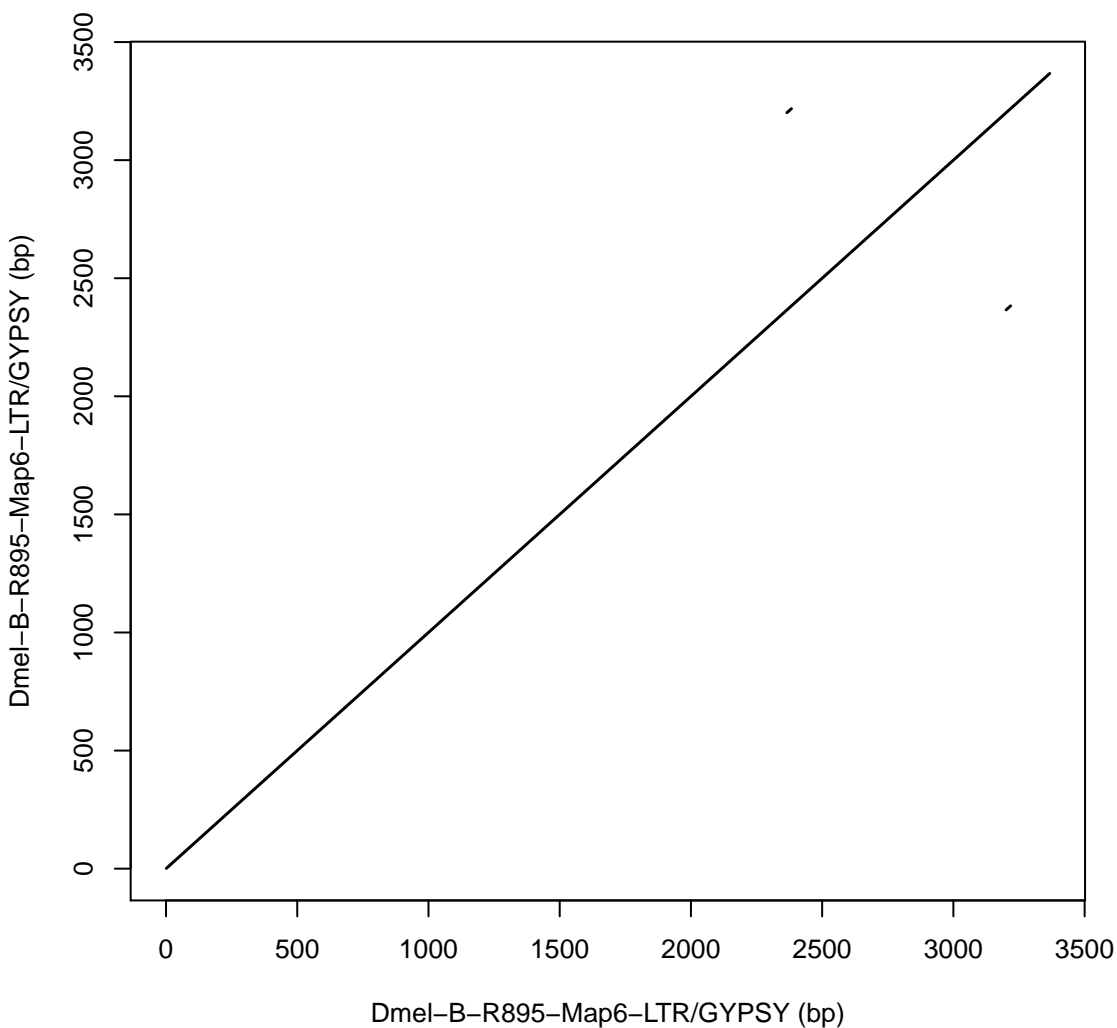
TE: Dmel-B-R895-Map6-LTR/GYPSY
consensus size: 3367bp; fragments: 101; full length: 0 (≥ 3030.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

