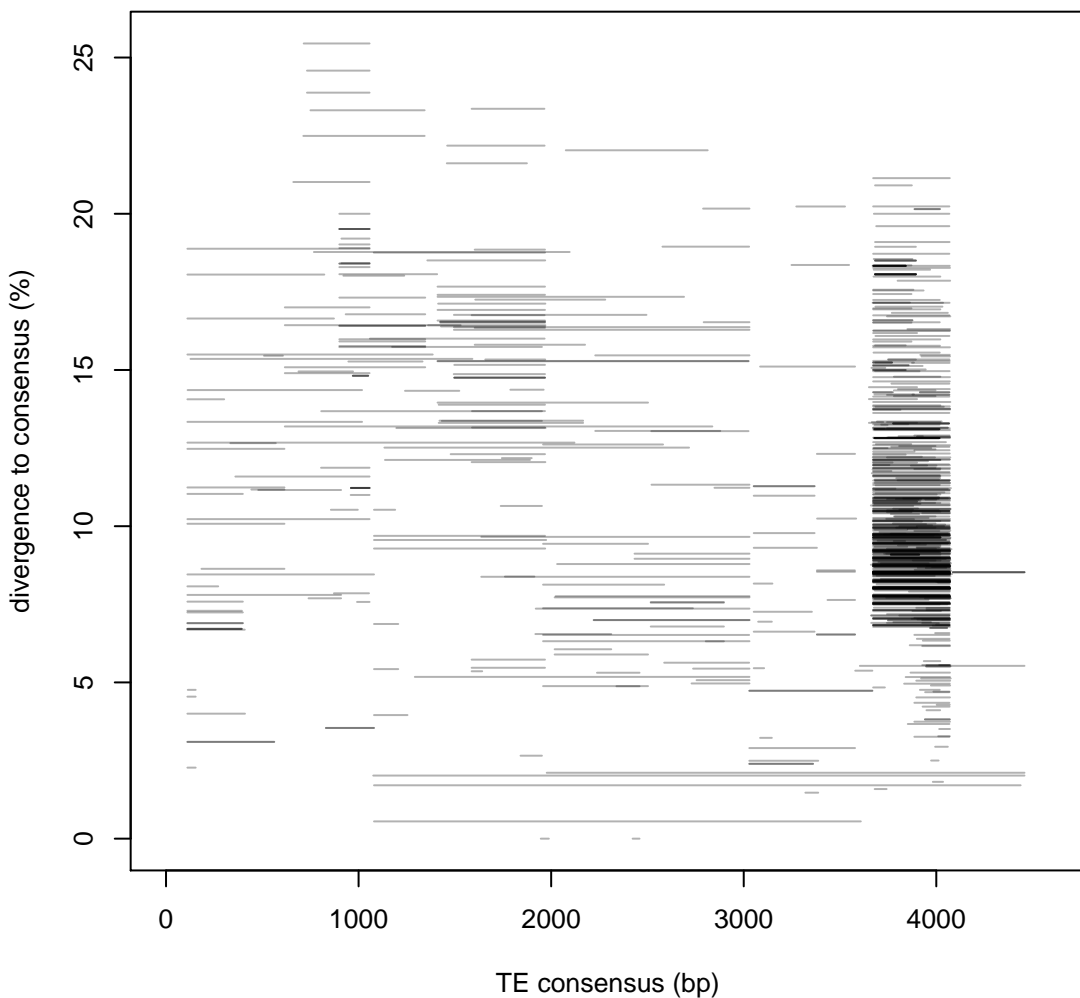
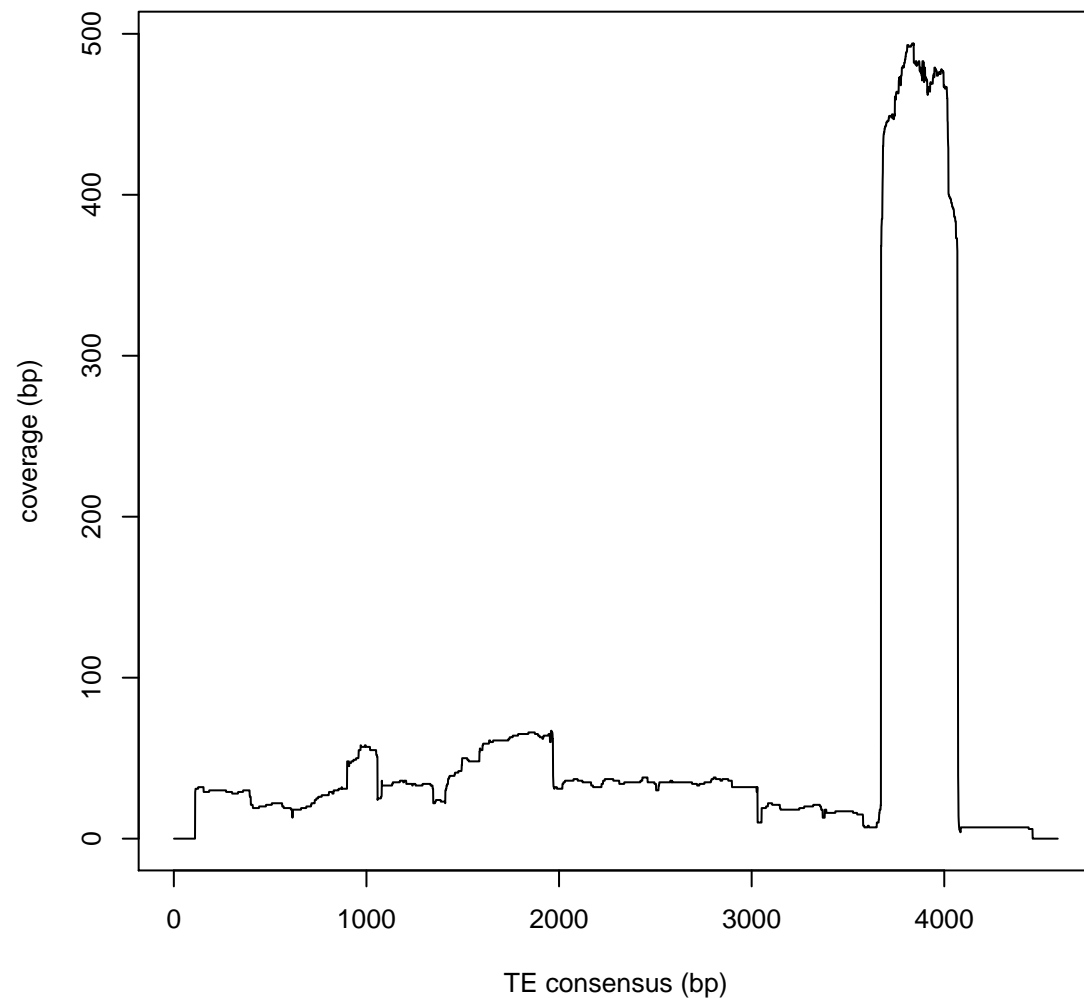


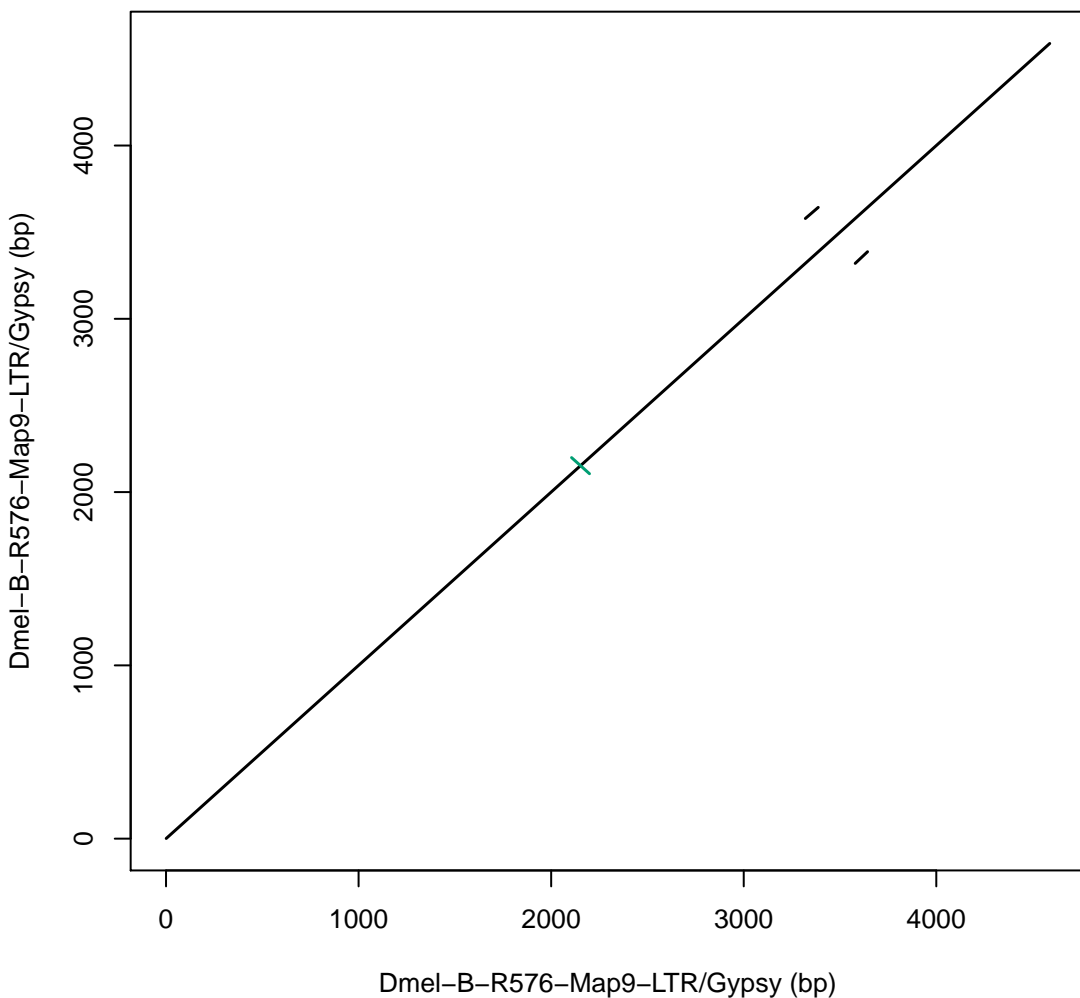
TE: Dmel-B-R576-Map9-LTR/Gypsy
consensus size: 4589bp; fragments: 819; full length: 0 (≥ 4130.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

