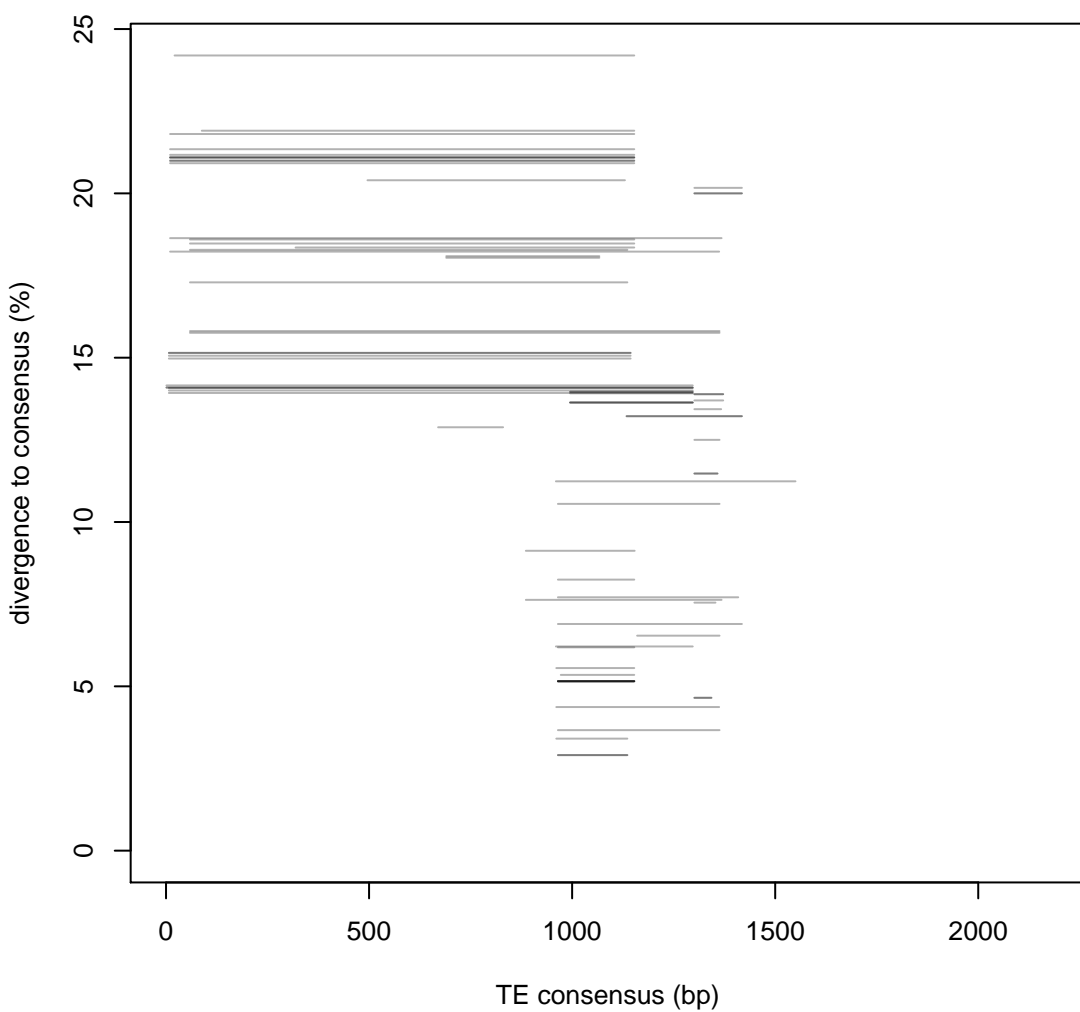
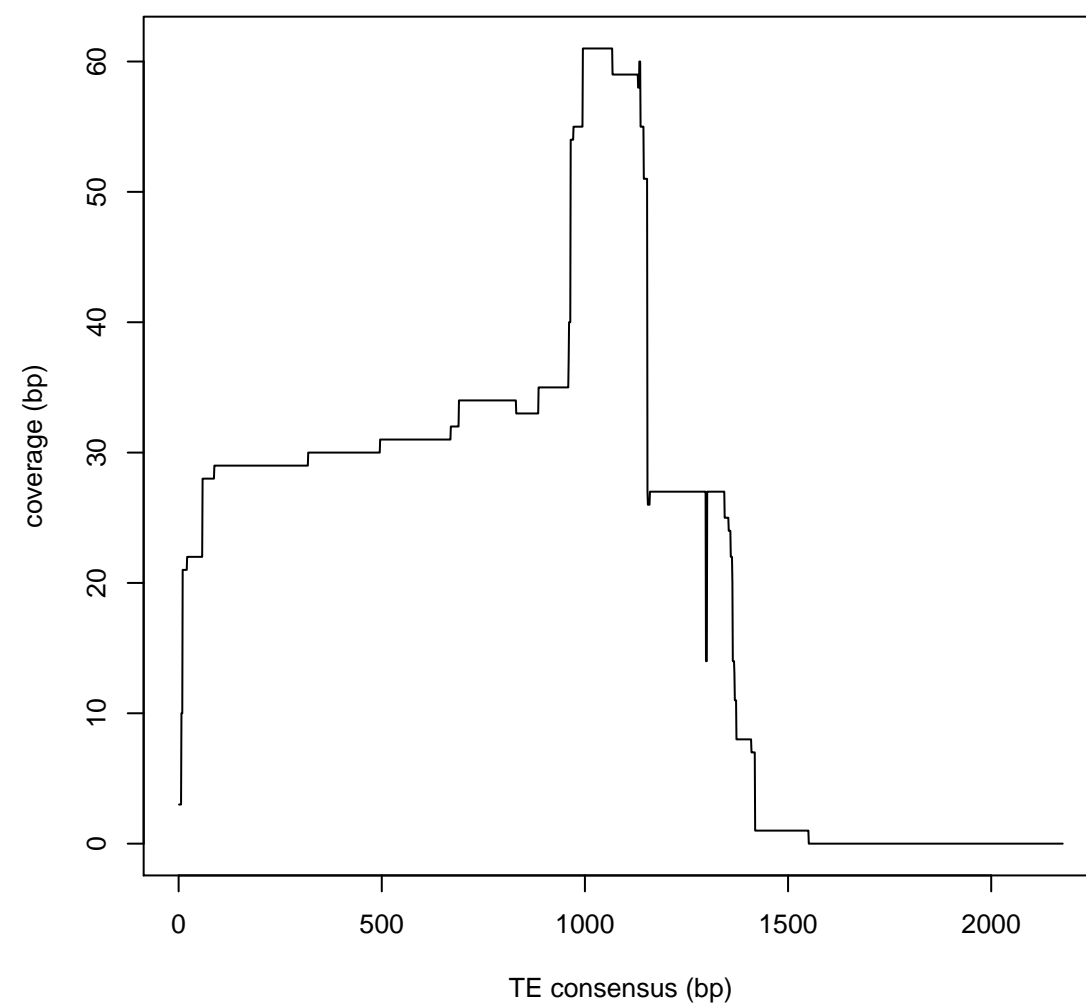


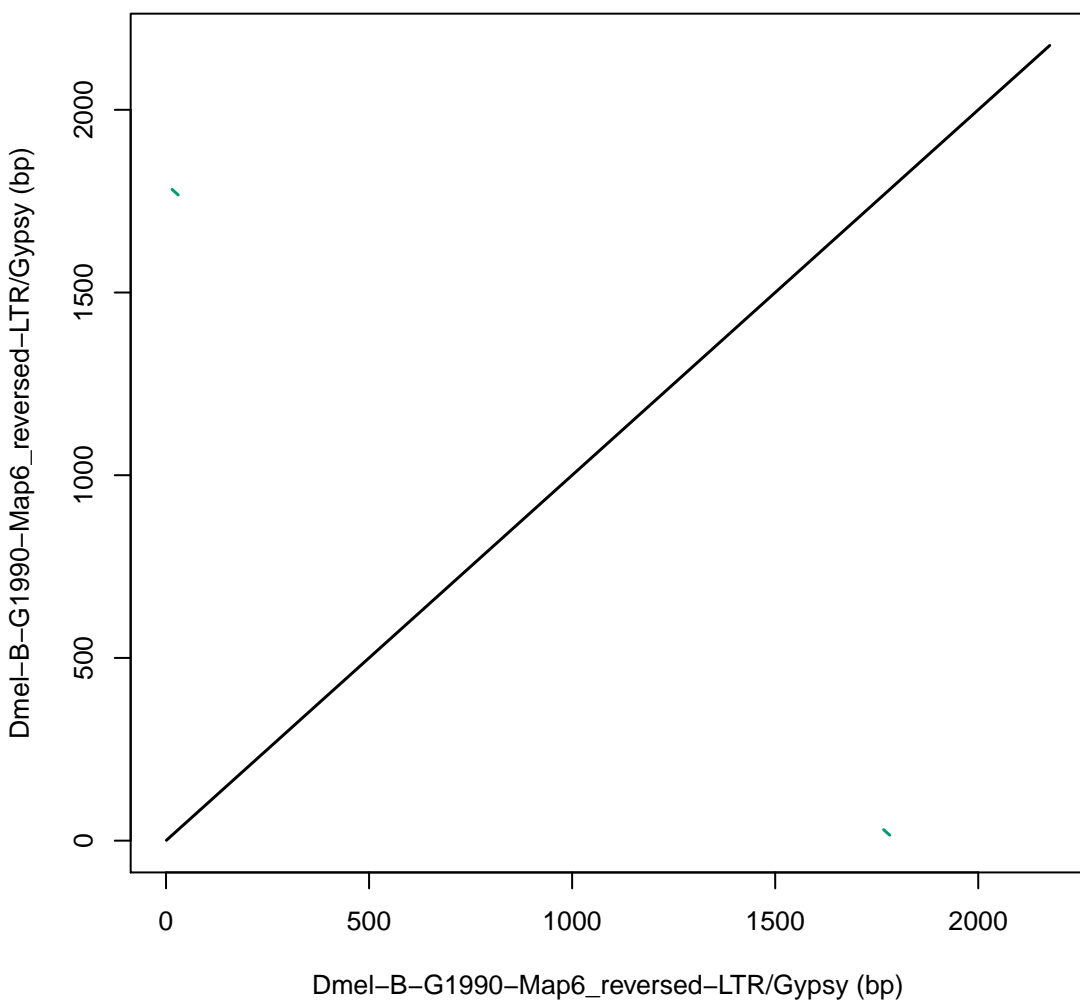
TE: Dmel-B-G1990-Map6_reversed-LTR/Gypsy
consensus size: 2176bp; fragments: 78; full length: 0 (≥ 1958.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

