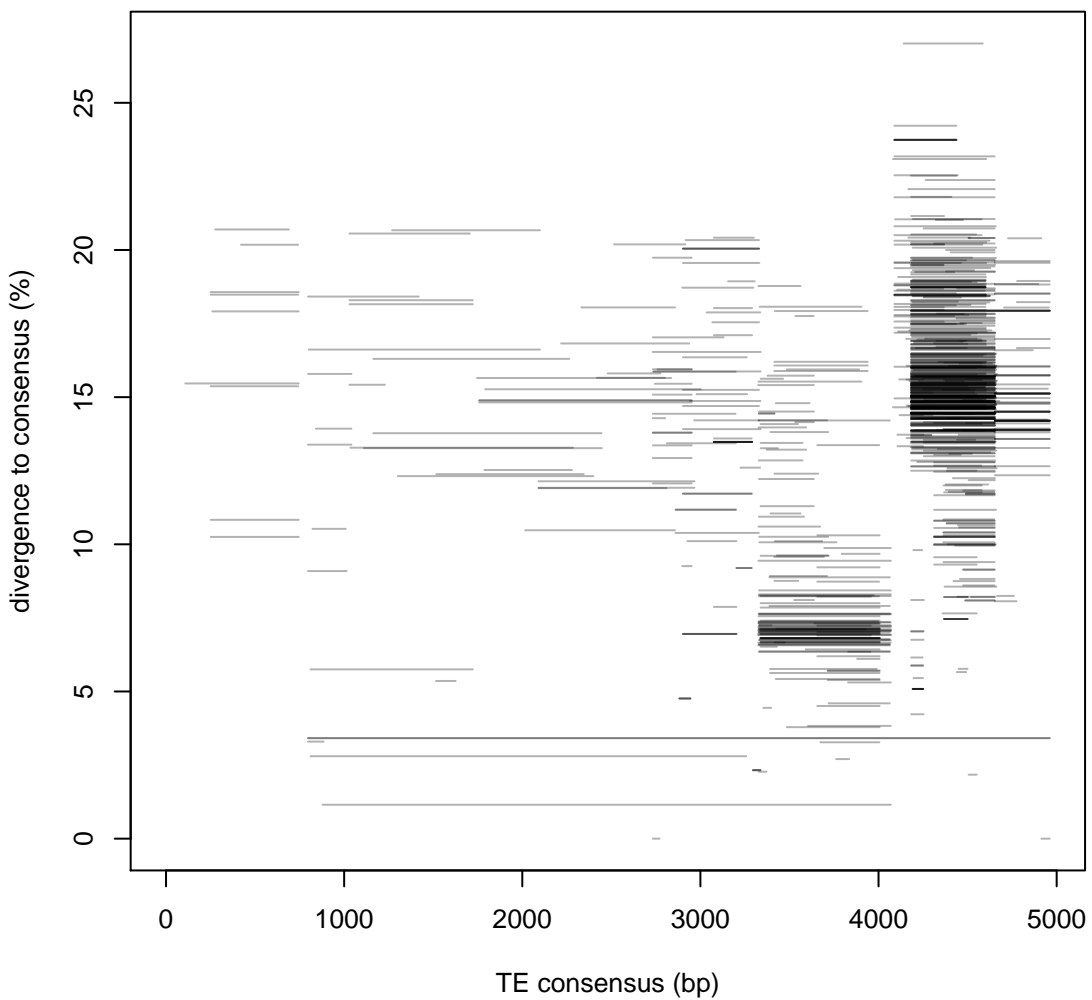
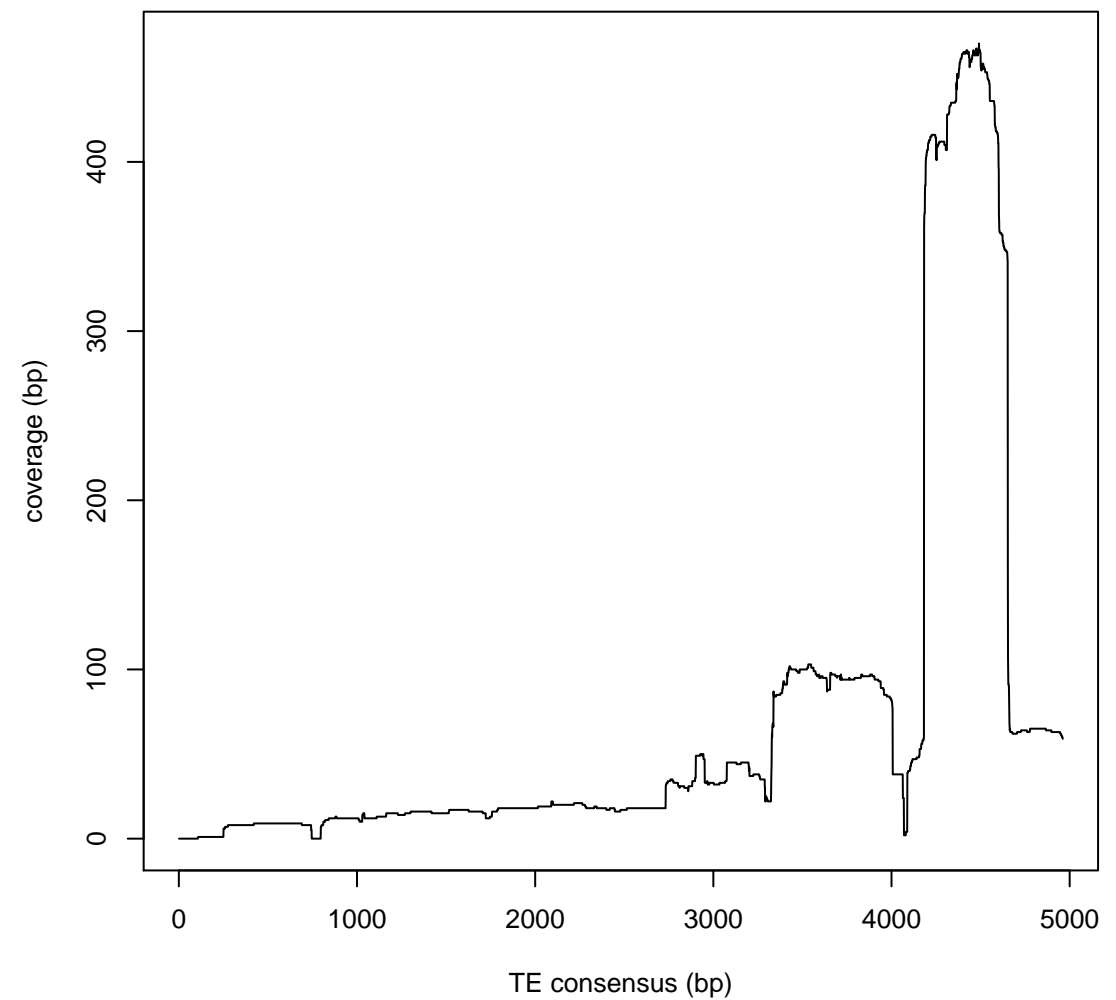


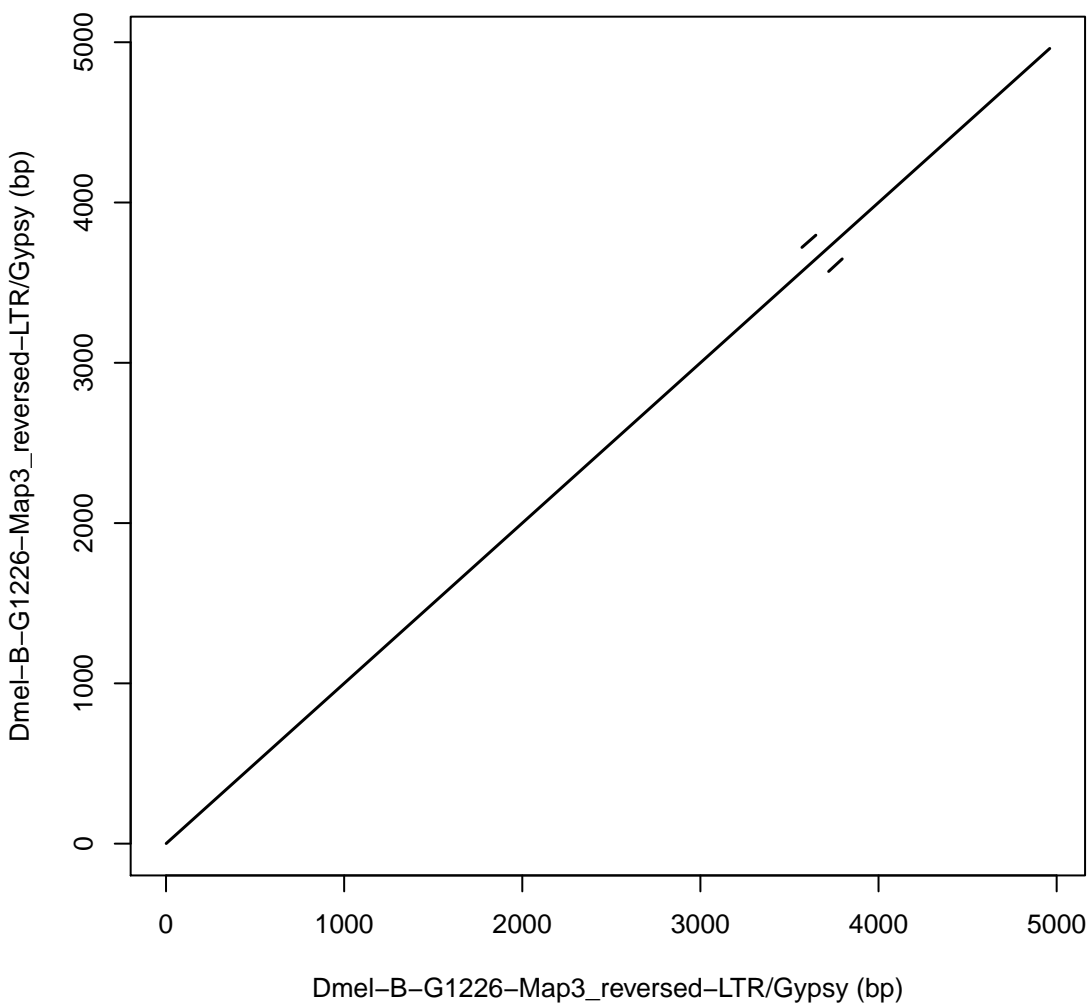
TE: Dmel-B-G1226-Map3_reversed-LTR/Gypsy
consensus size: 4961bp; fragments: 838; full length: 0 (>=4464.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

