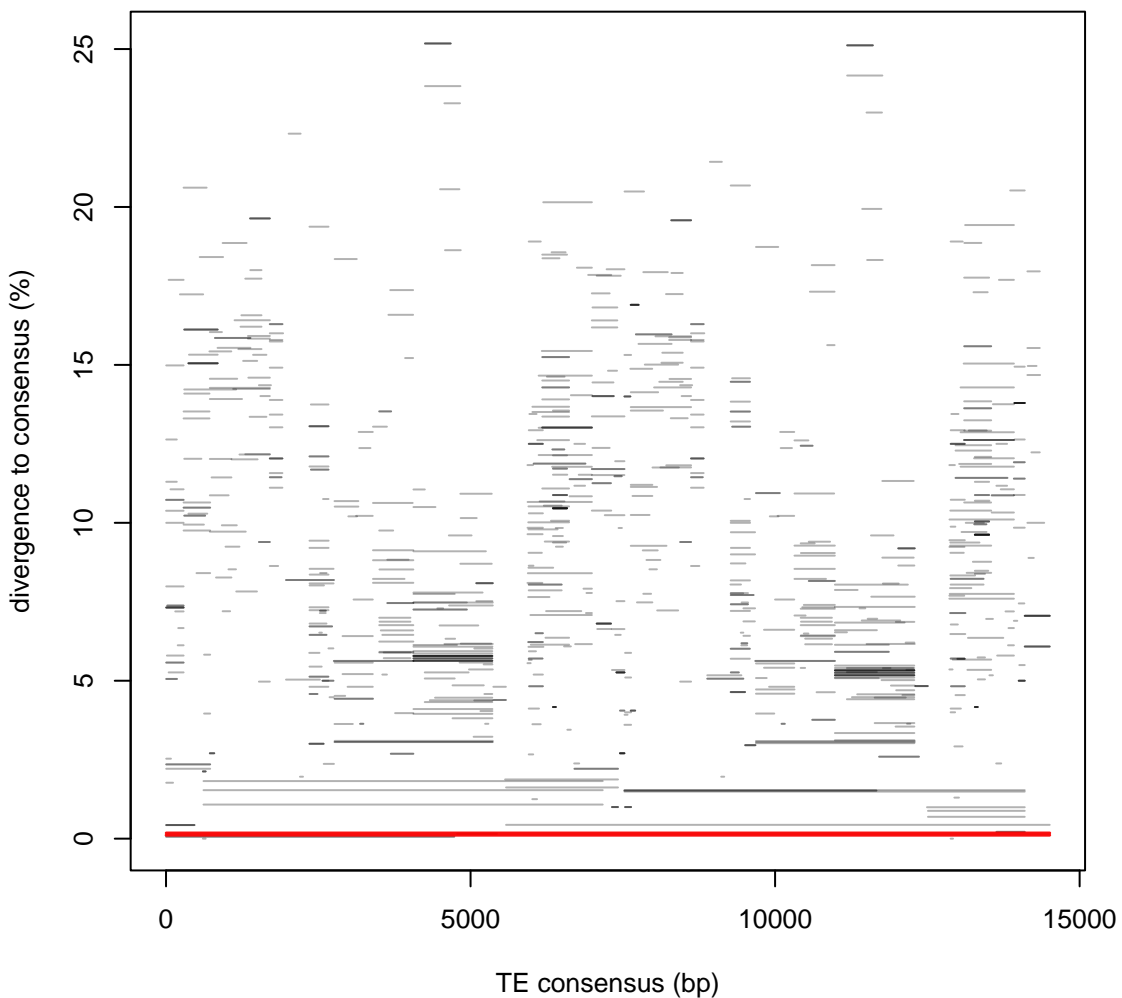
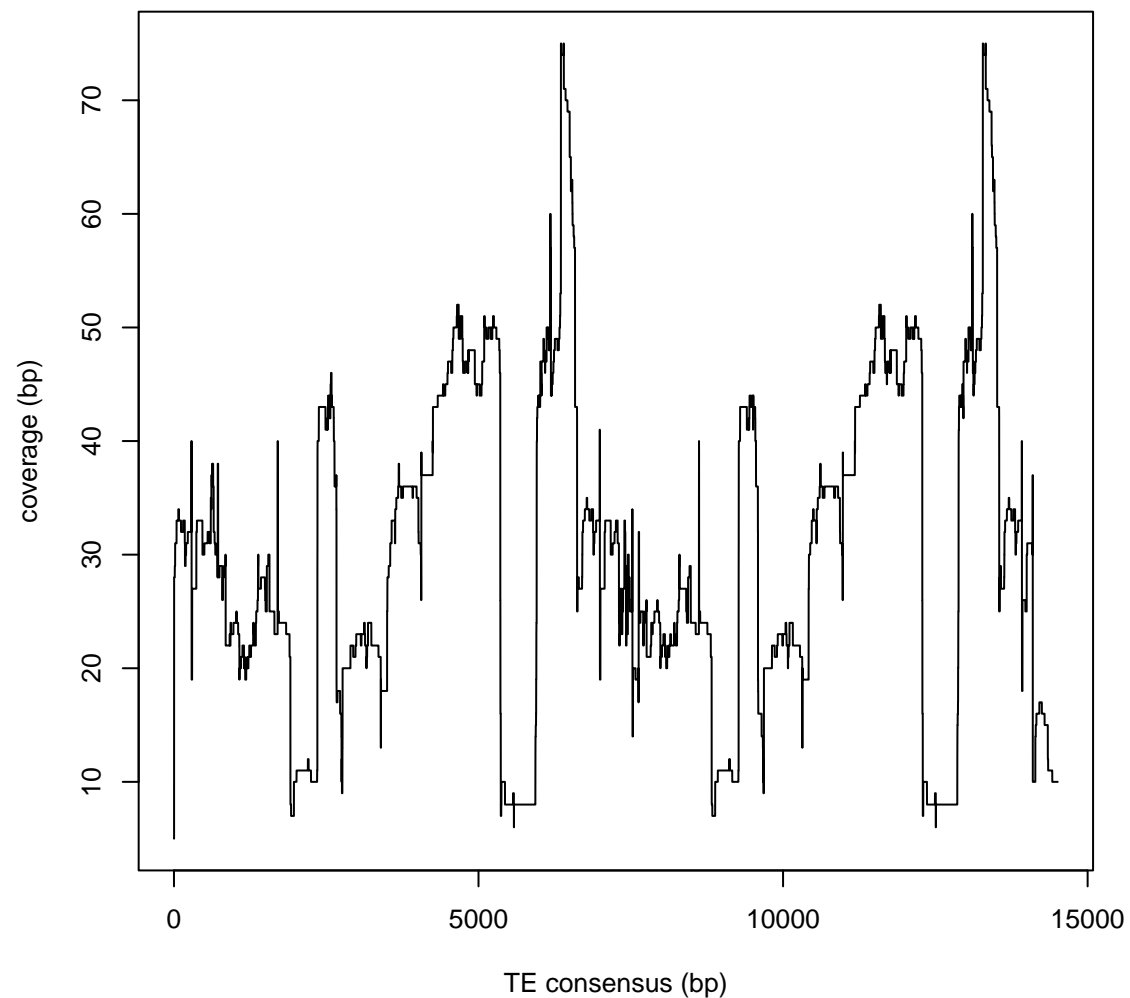


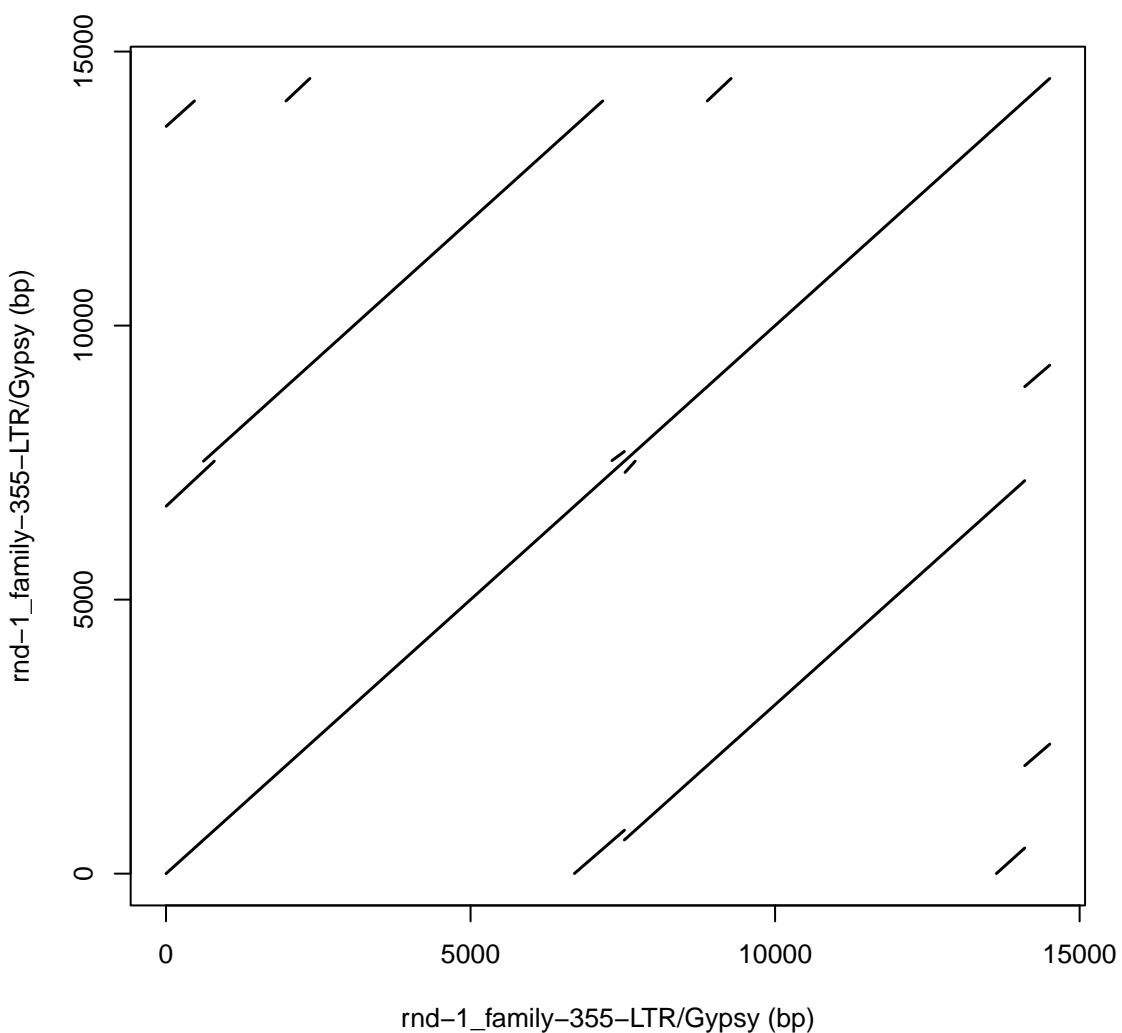
TE: rnd-1_family-355-LTR/Gypsy
consensus size: 14509bp; fragments: 889; full length: 2 (≥ 13058.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

