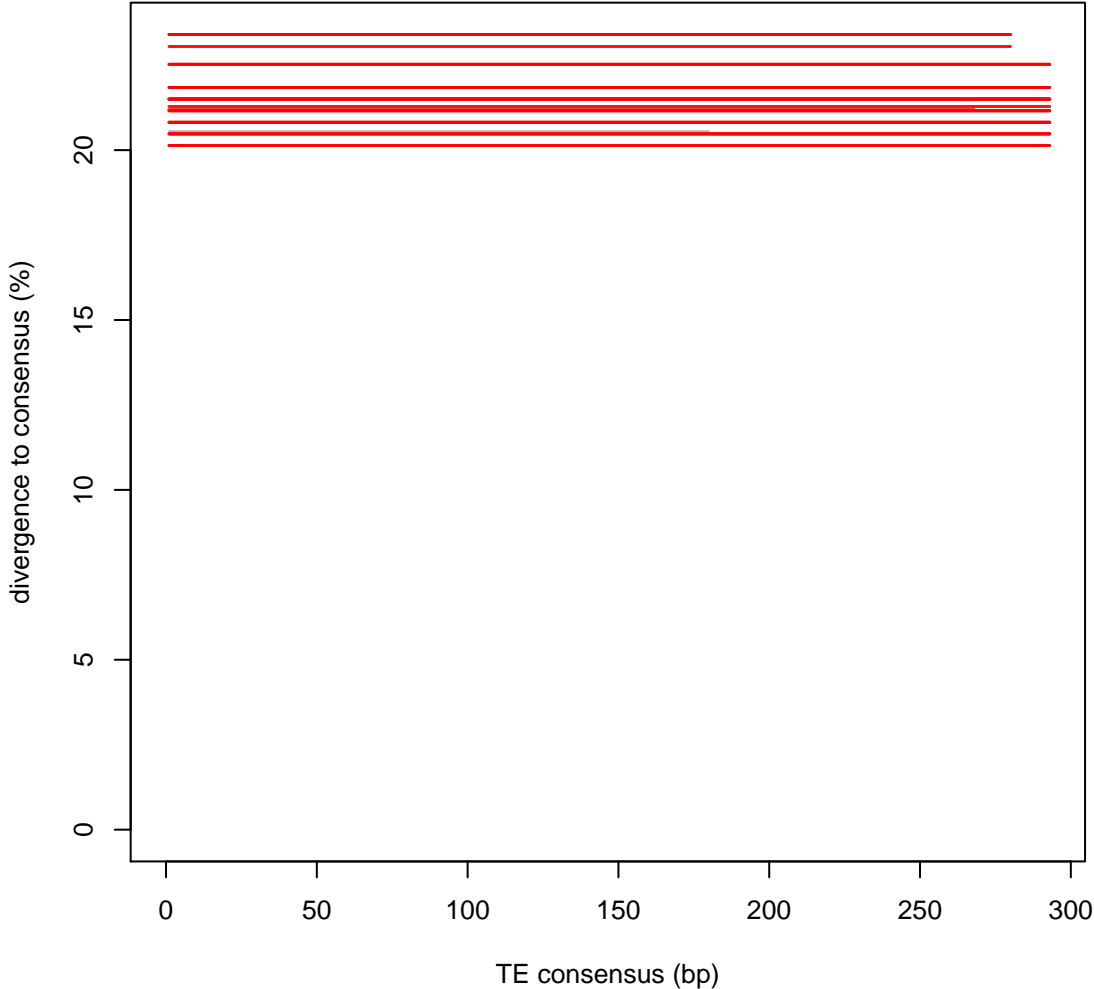
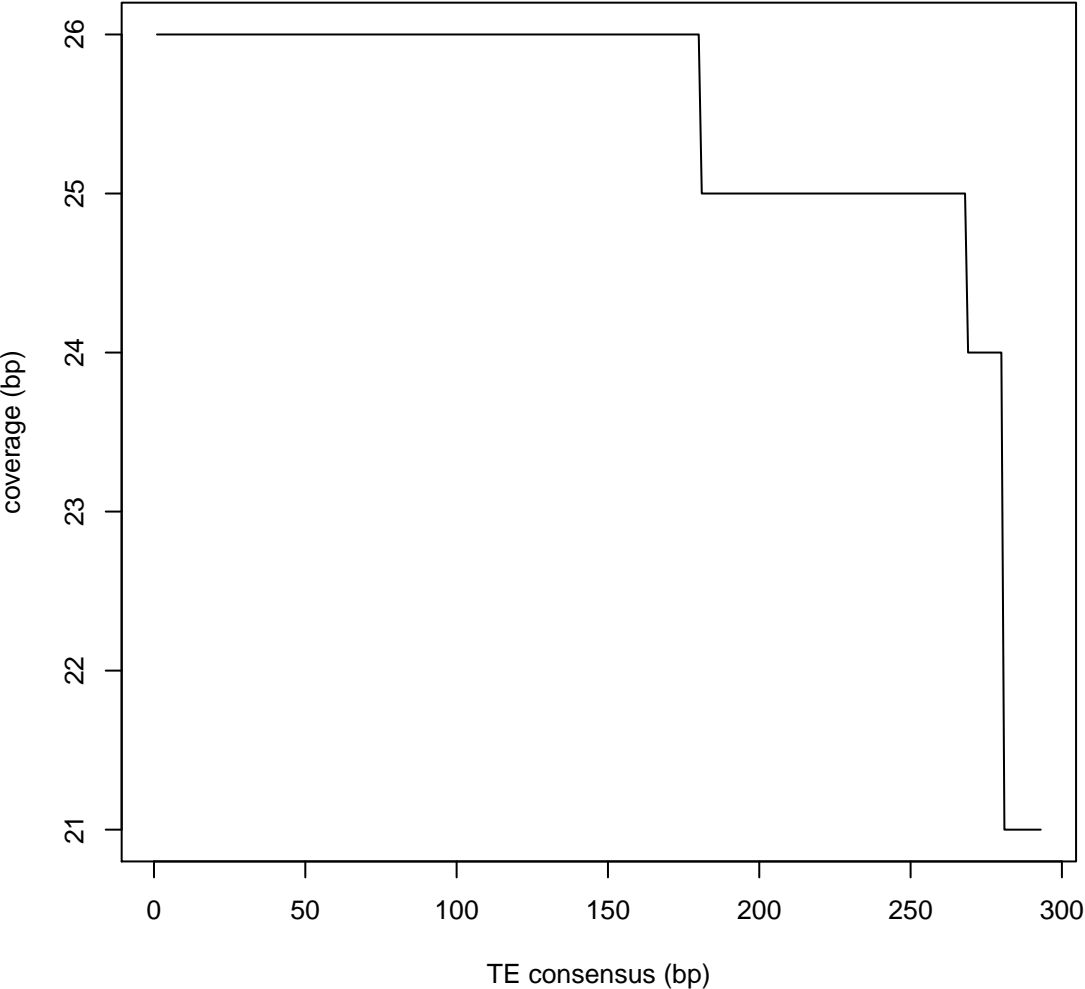


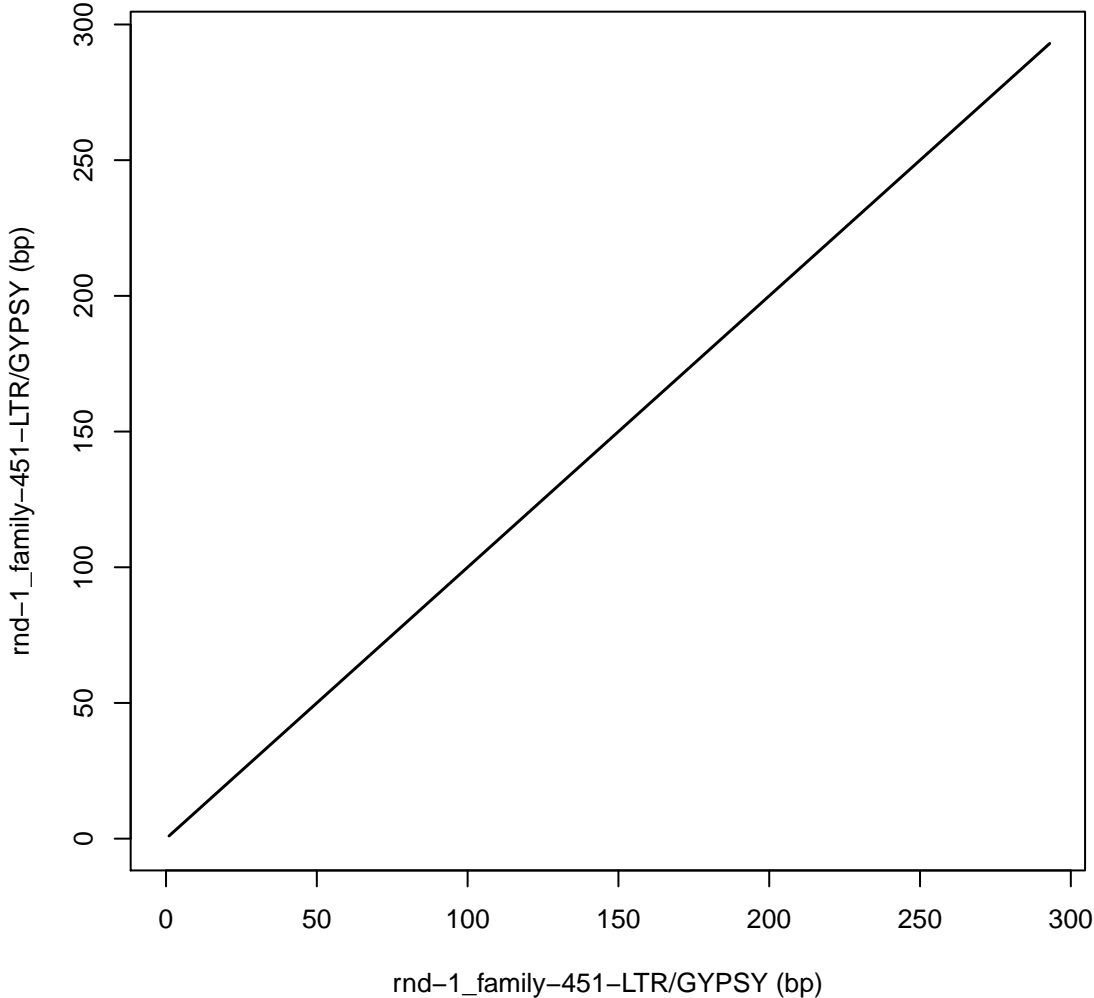
TE: rnd-1_family-451-LTR/GYPSY
consensus size: 293bp; fragments: 26; full length: 25 (≥ 263.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

