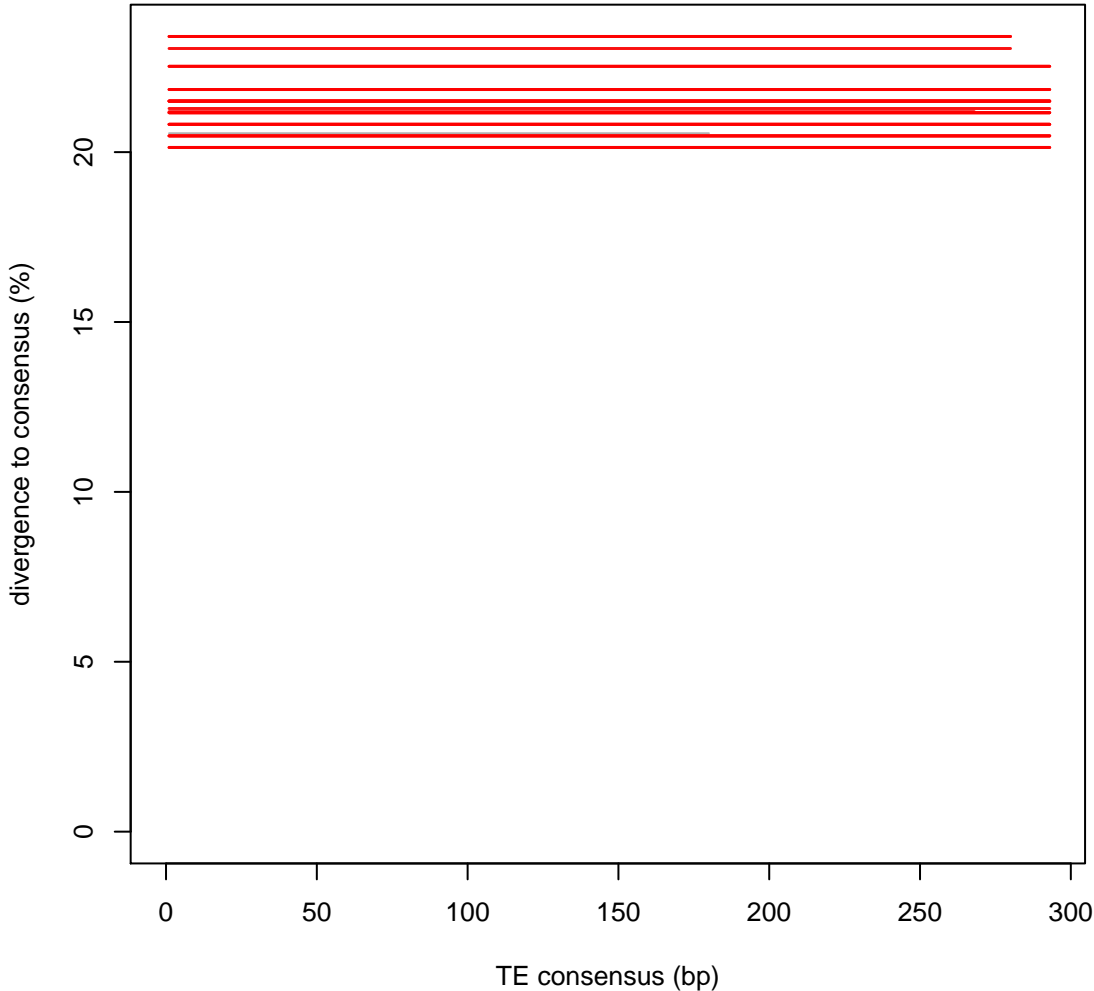
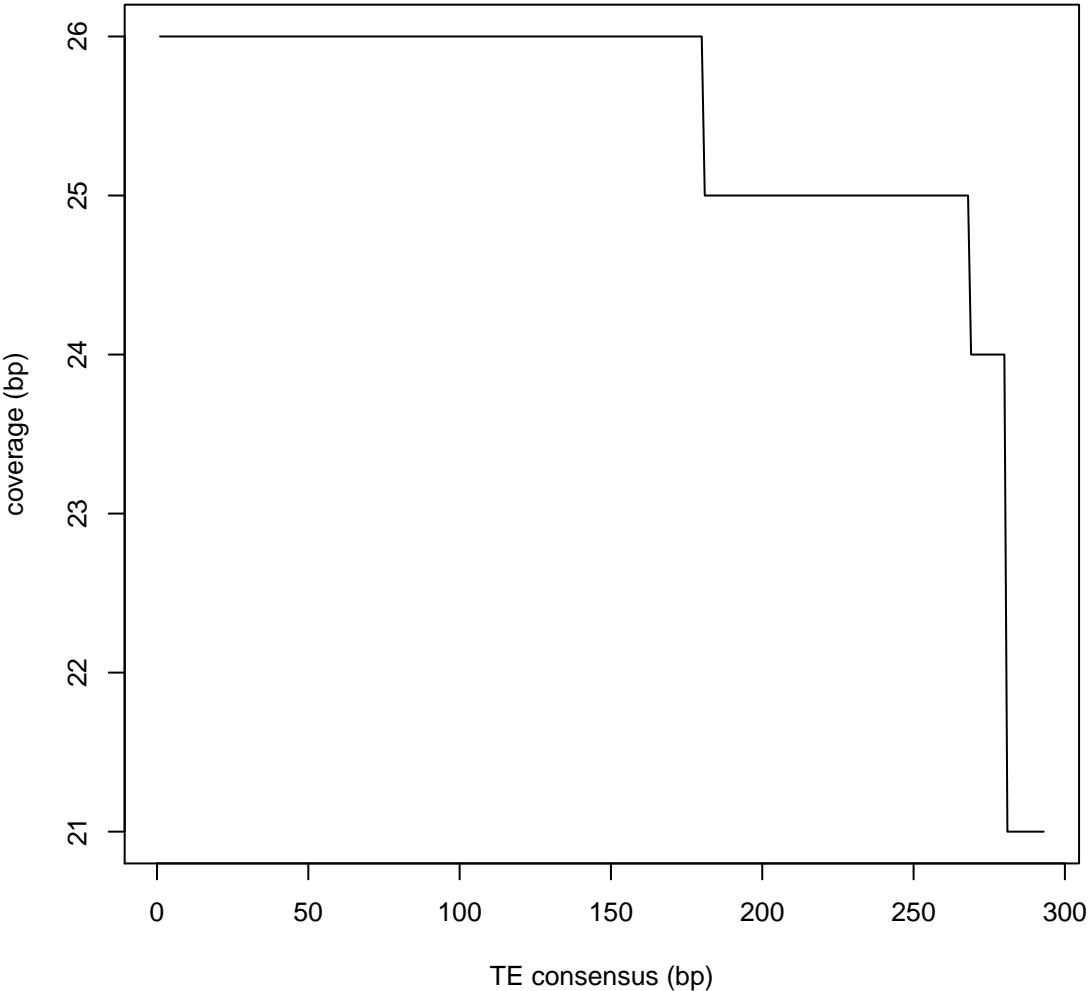


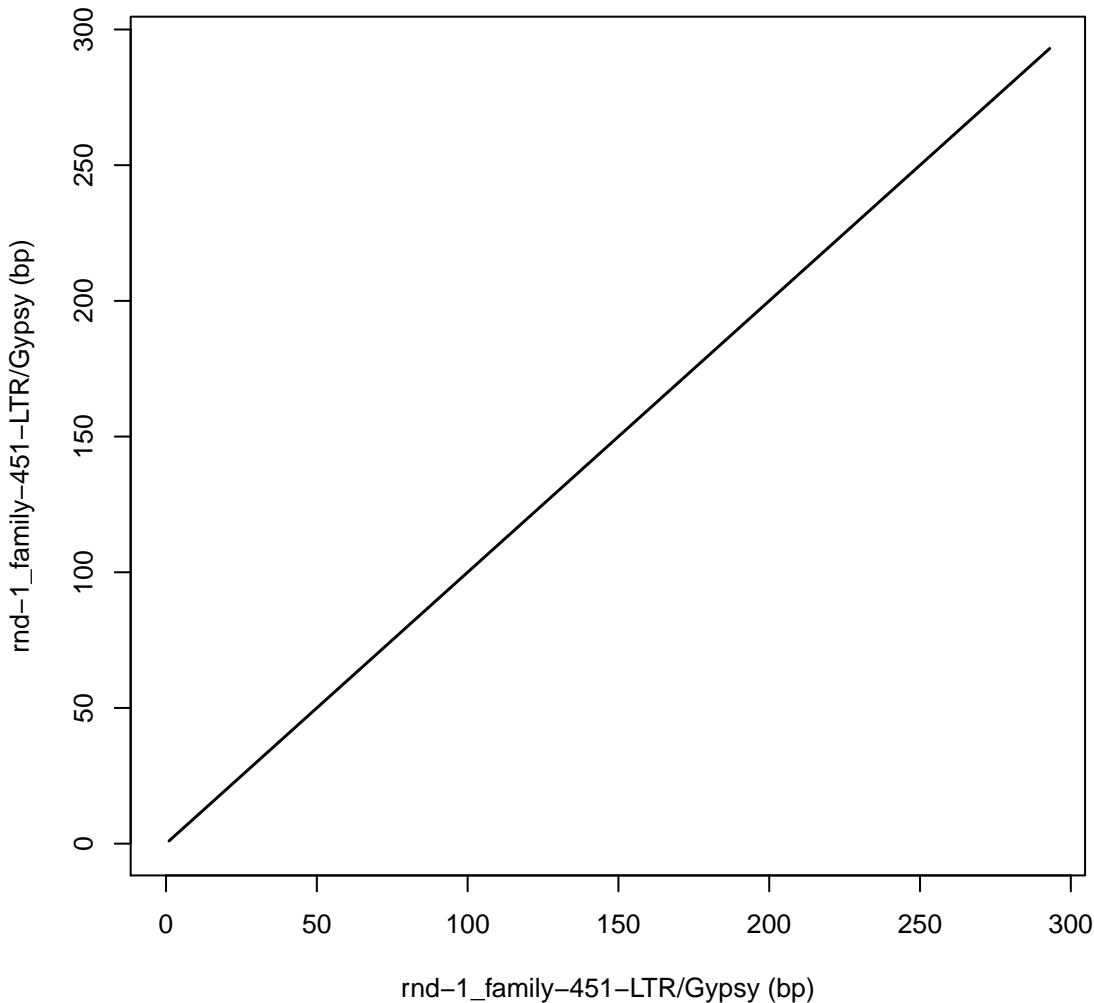
TE: rnd-1_family-451-LTR/Gypsy
consensus size: 293bp; fragments: 26; full length: 25 (>=263.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

