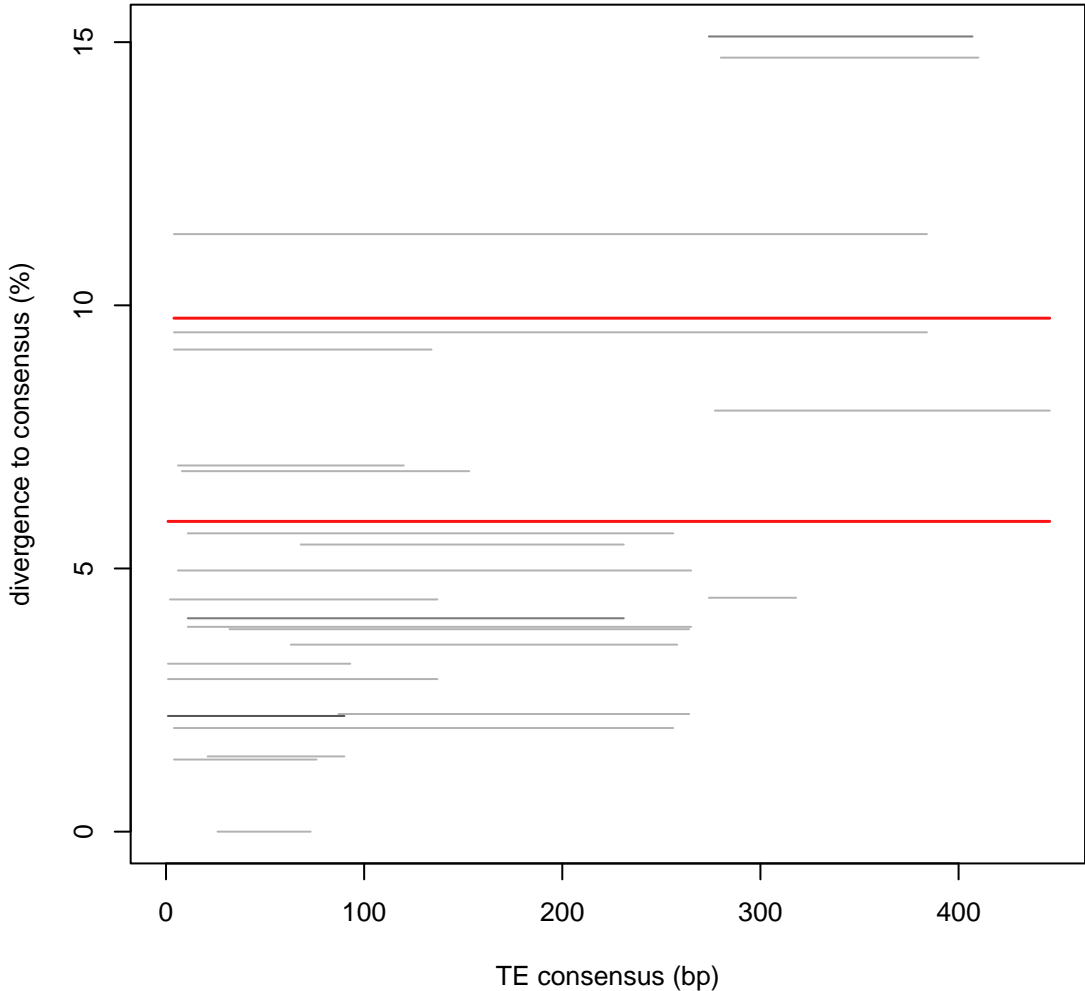
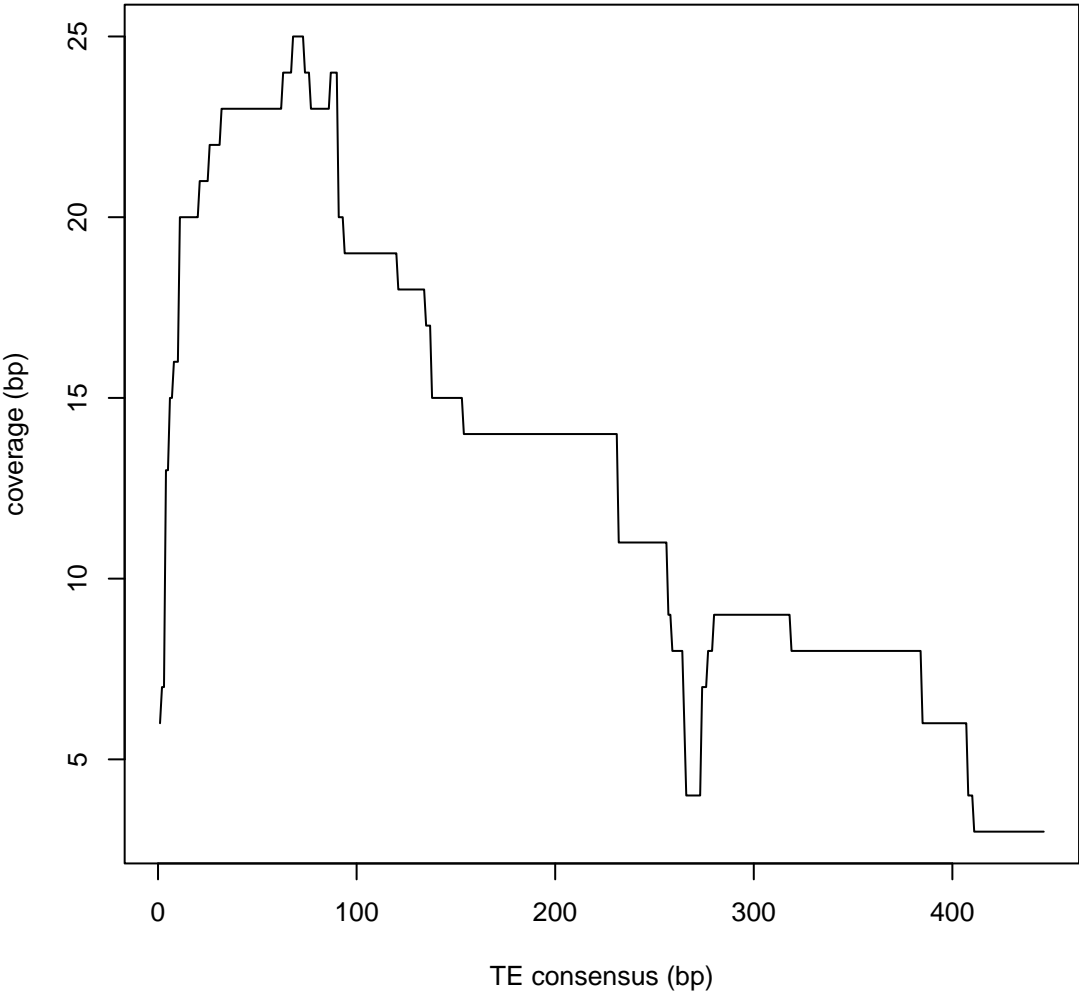


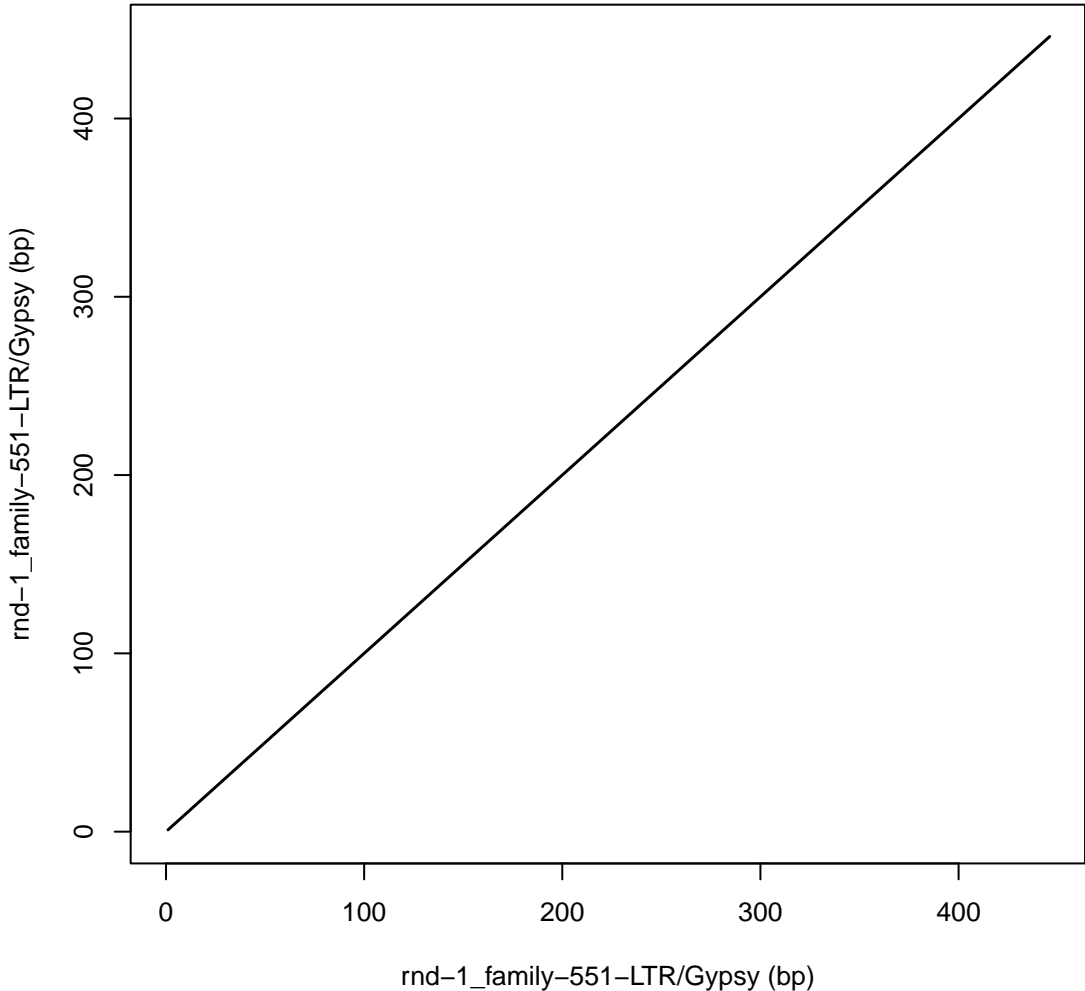
TE: rnd-1_family-551-LTR/Gypsy
consensus size: 446bp; fragments: 31; full length: 2 (≥ 401.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

