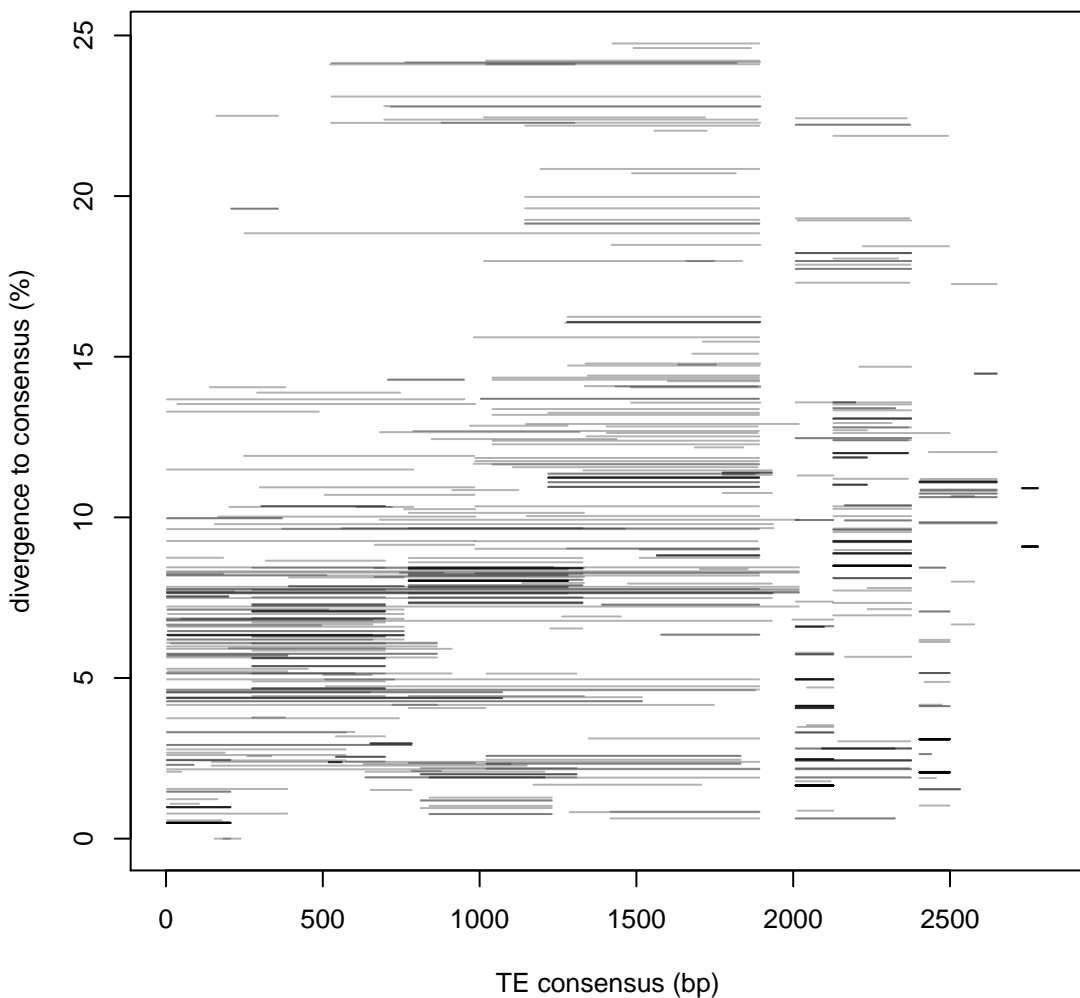
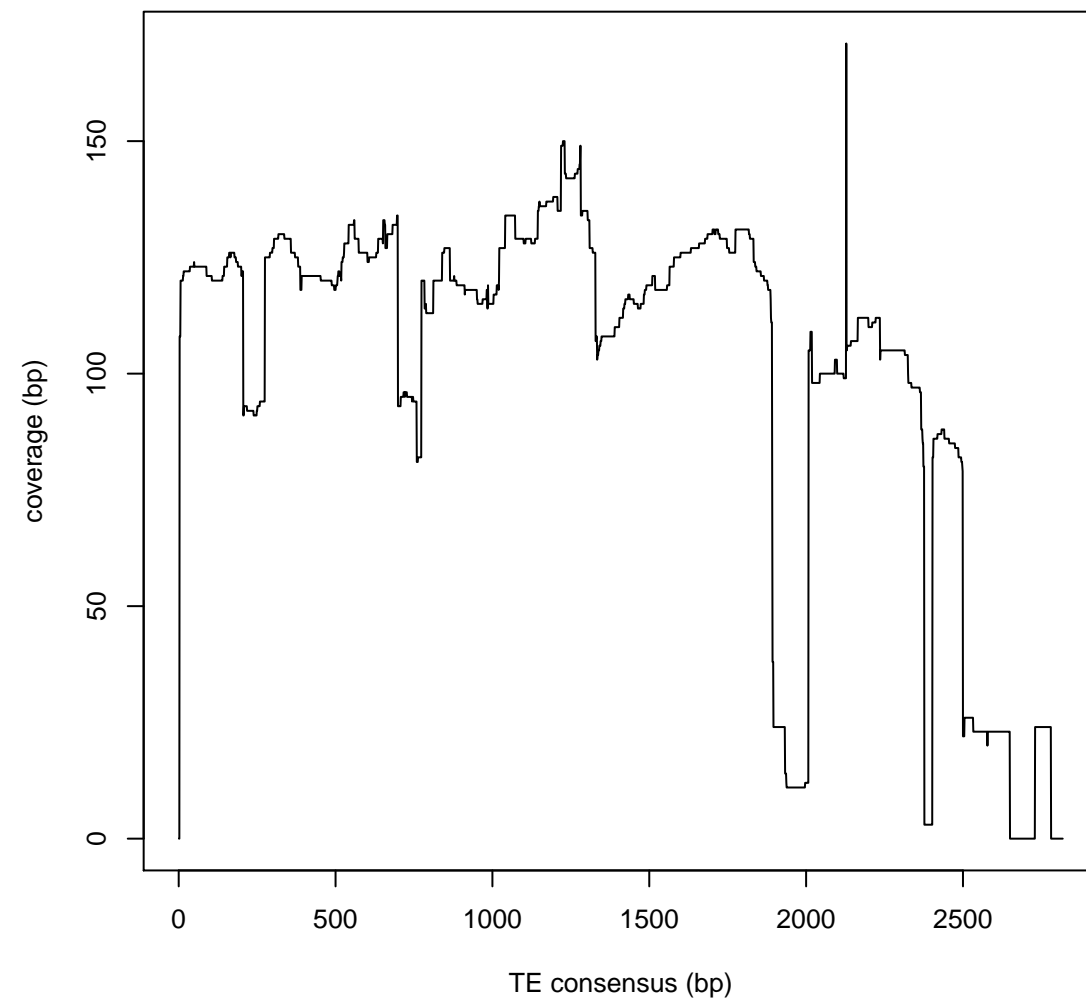


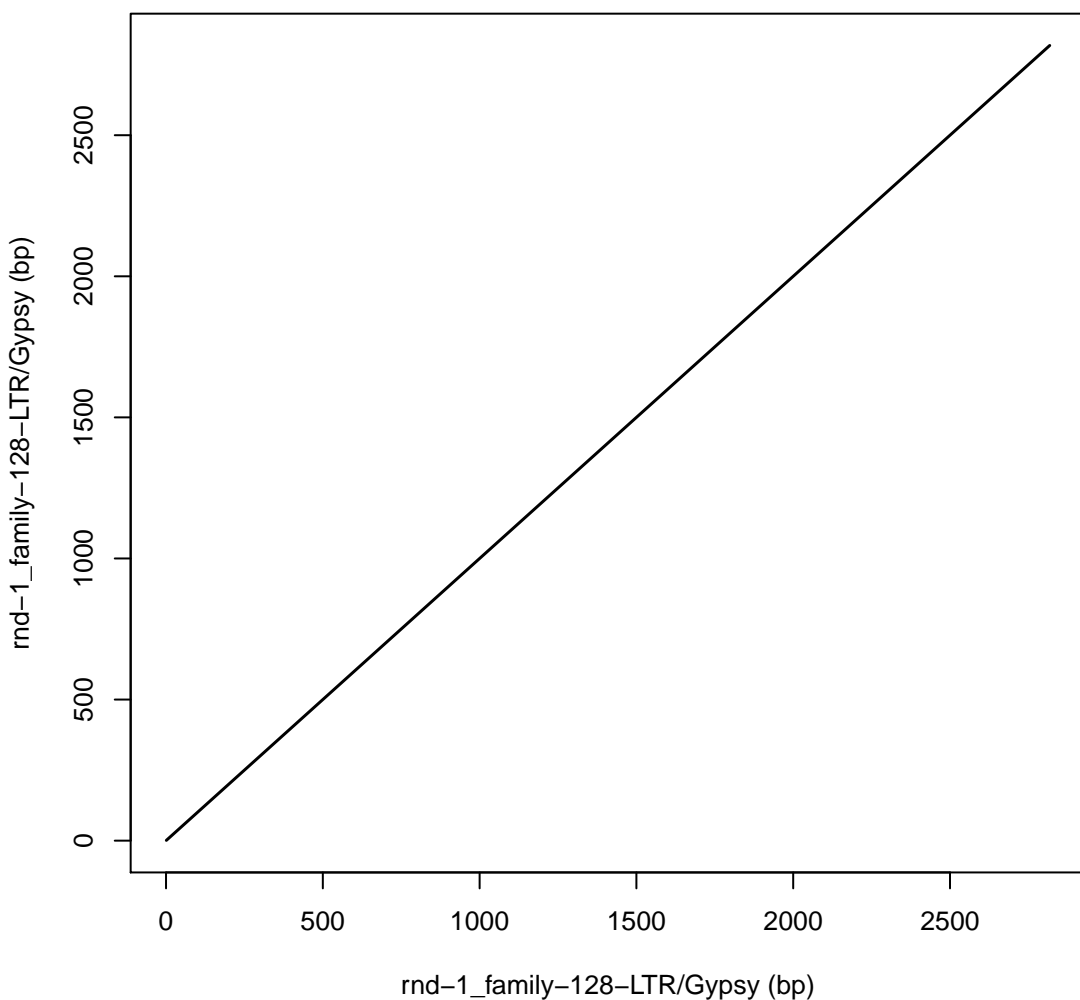
TE: rnd-1_family-128-LTR/Gypsy
consensus size: 2818bp; fragments: 685; full length: 0 (≥ 2536.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

