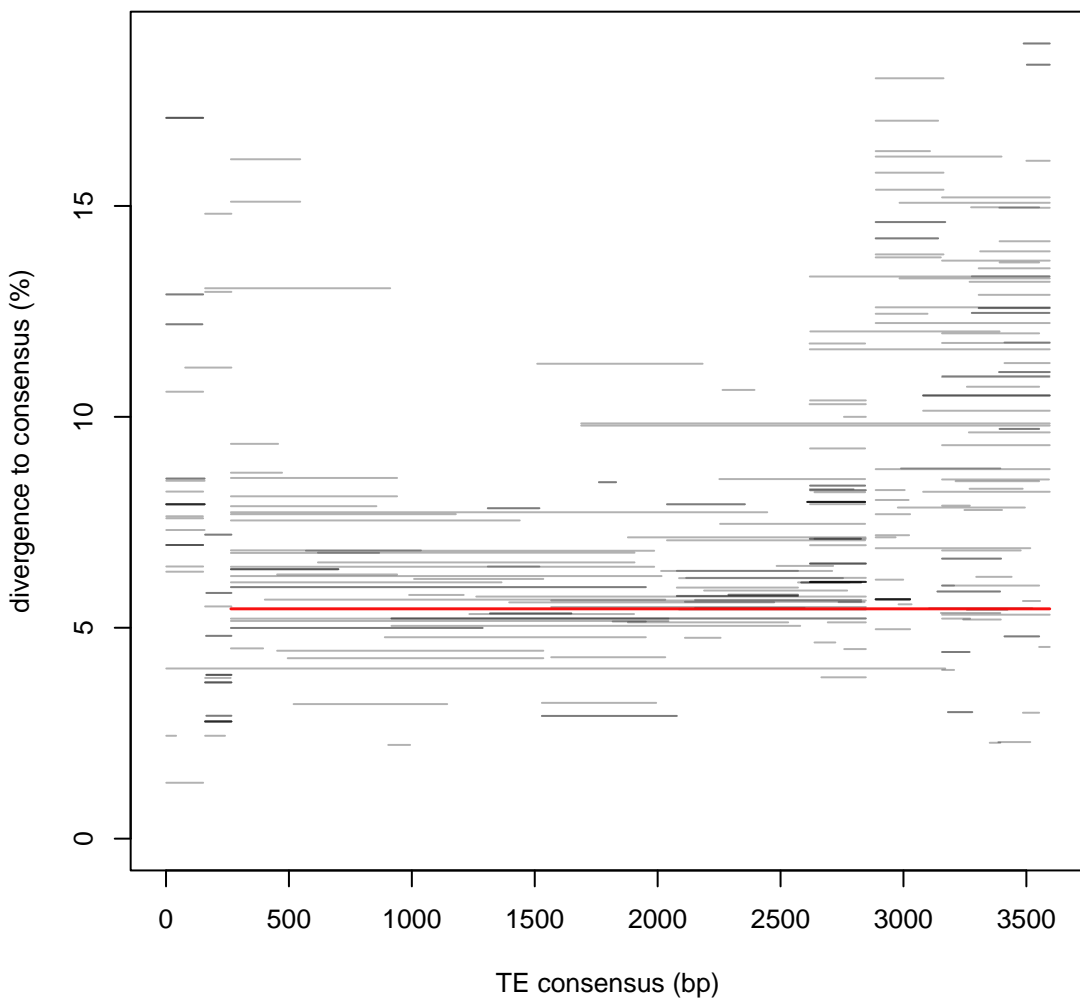
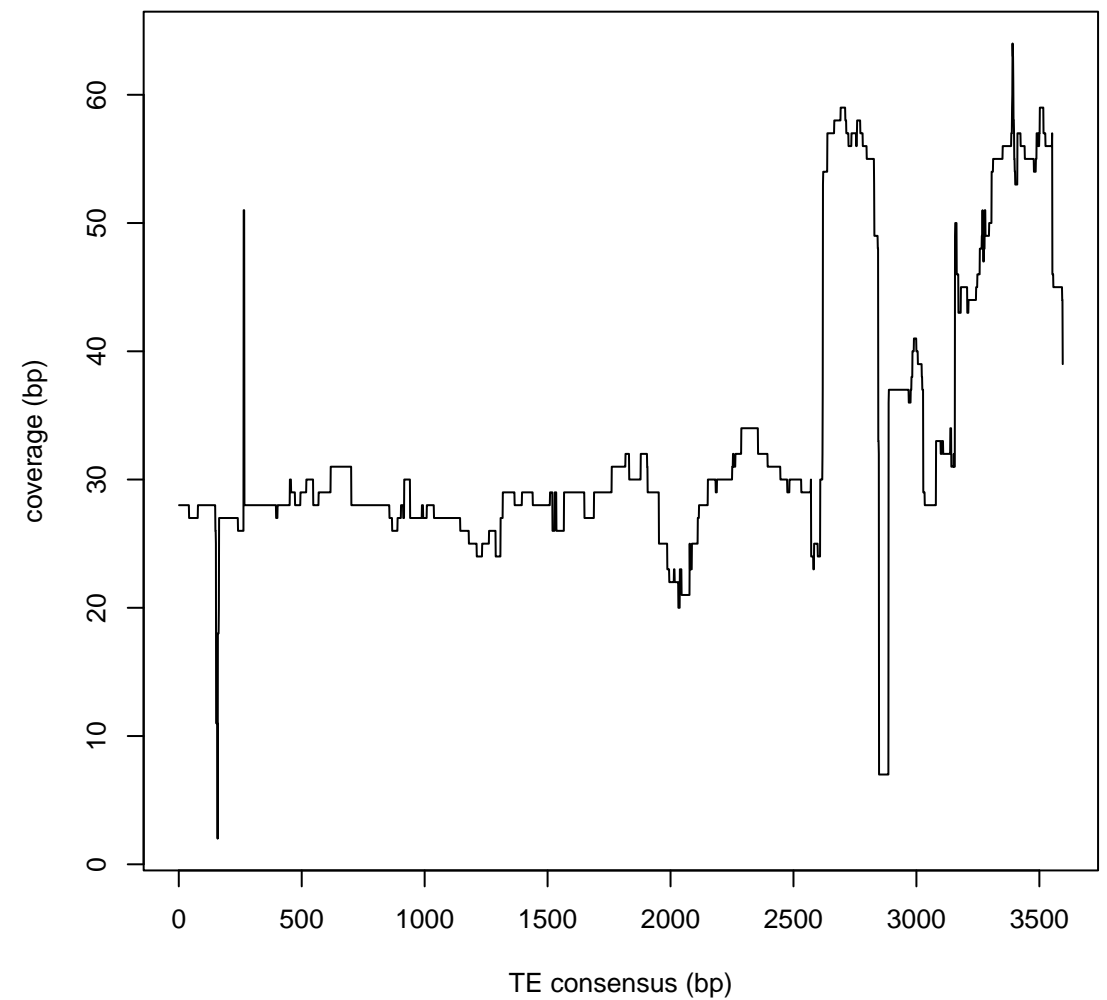


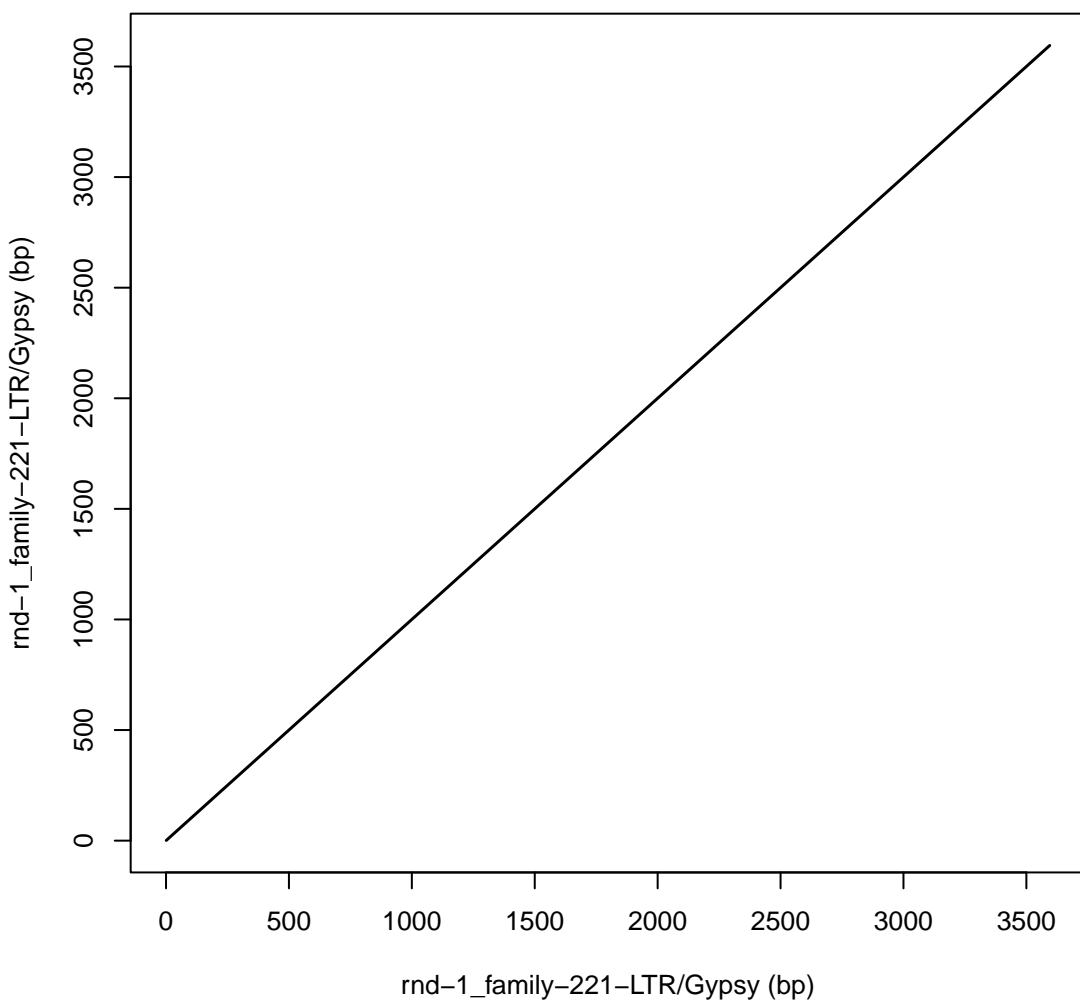
TE: rnd-1_family-221-LTR/Gypsy
consensus size: 3595bp; fragments: 284; full length: 1 (≥ 3235.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

