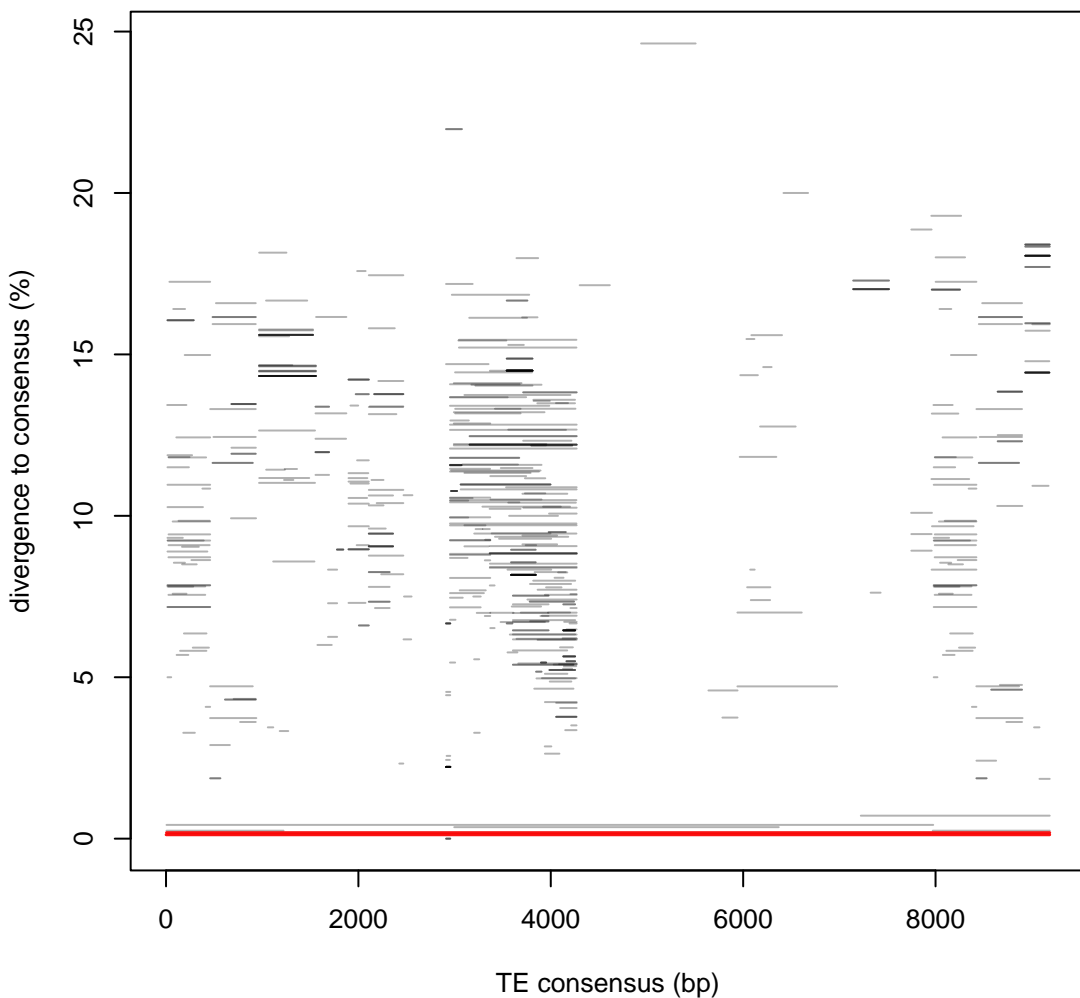
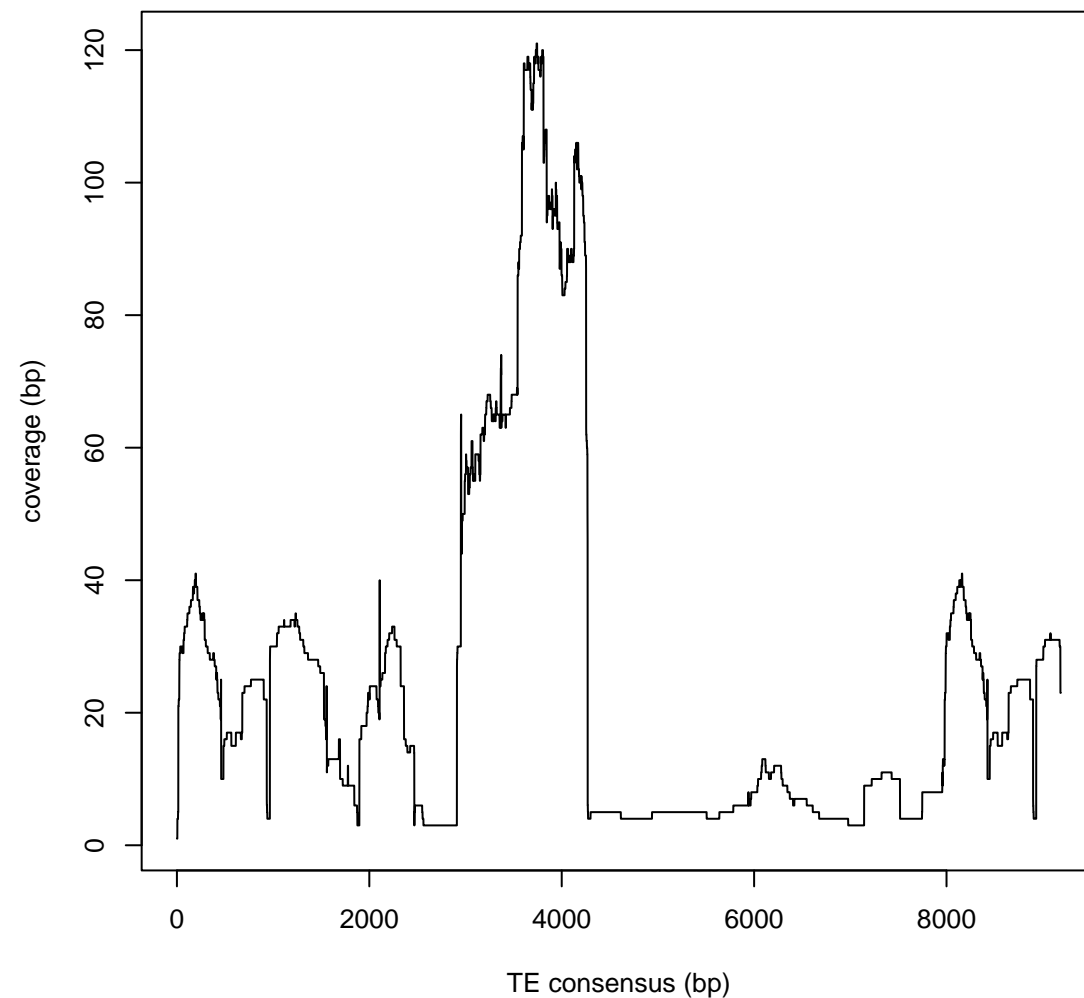


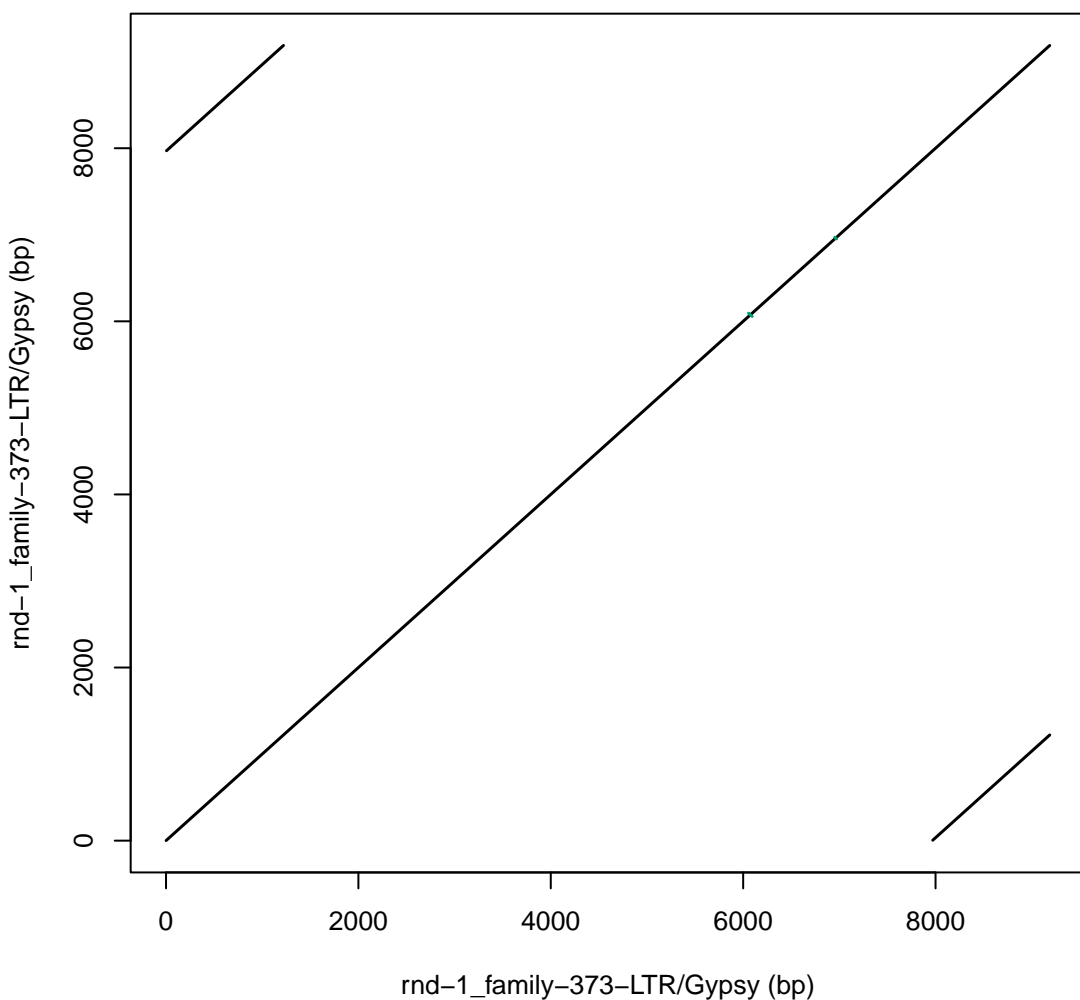
TE: rnd-1_family-373-LTR/Gypsy
consensus size: 9187bp; fragments: 578; full length: 2 (≥ 8268.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

