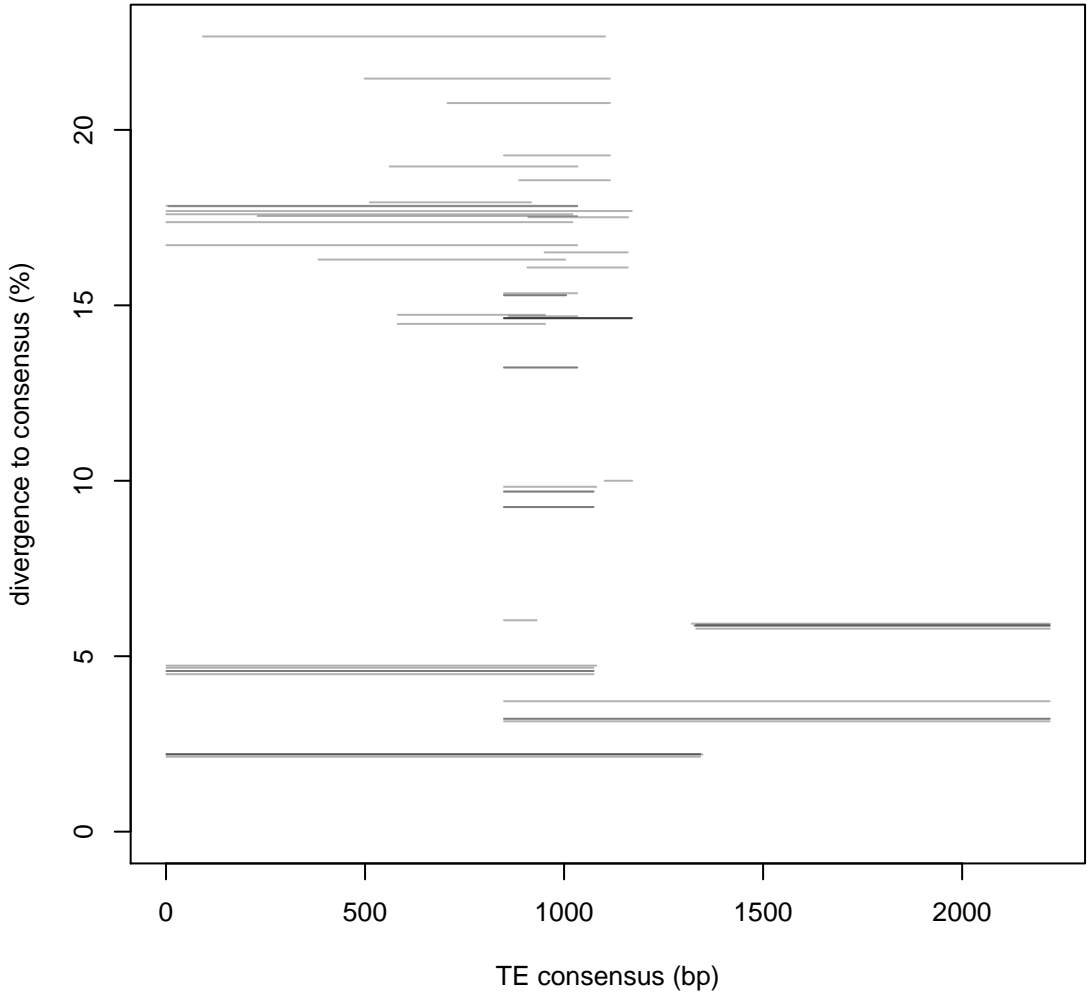
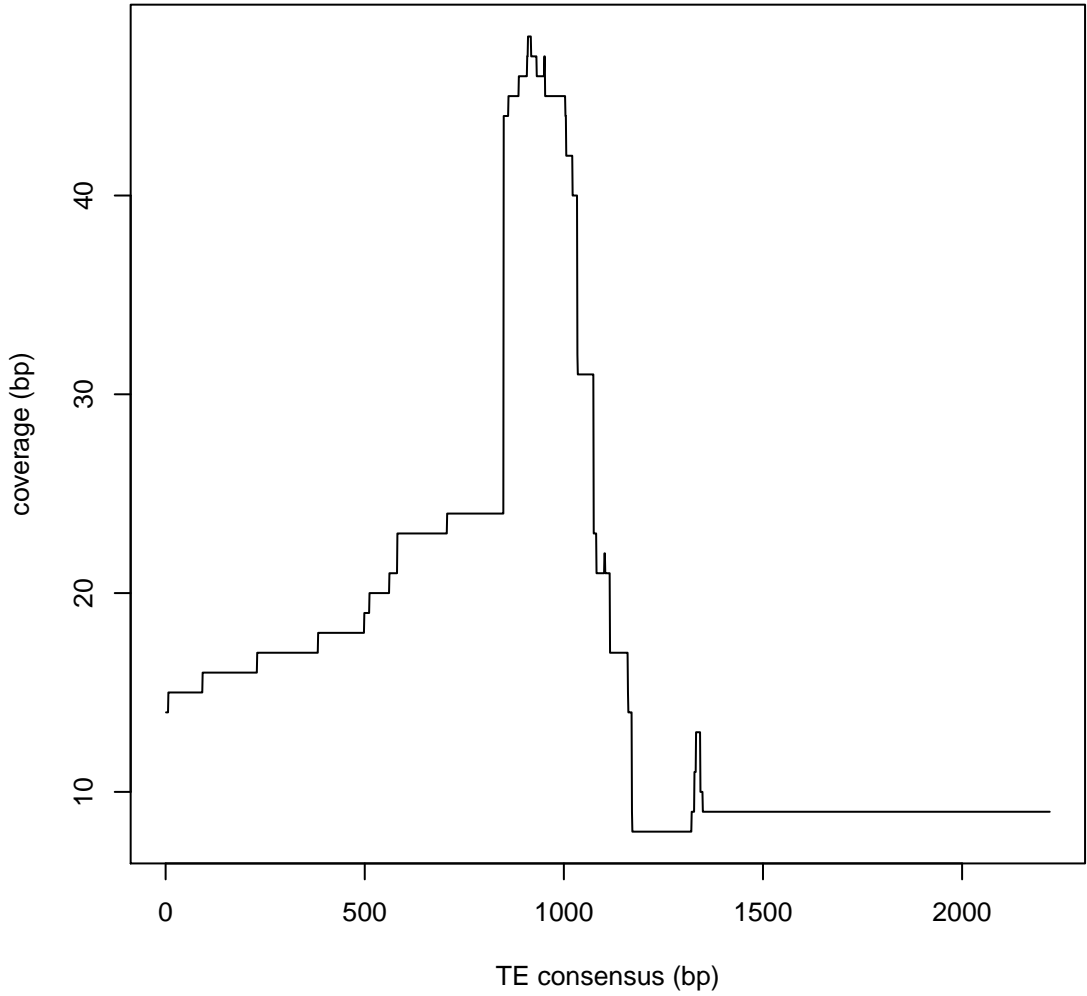


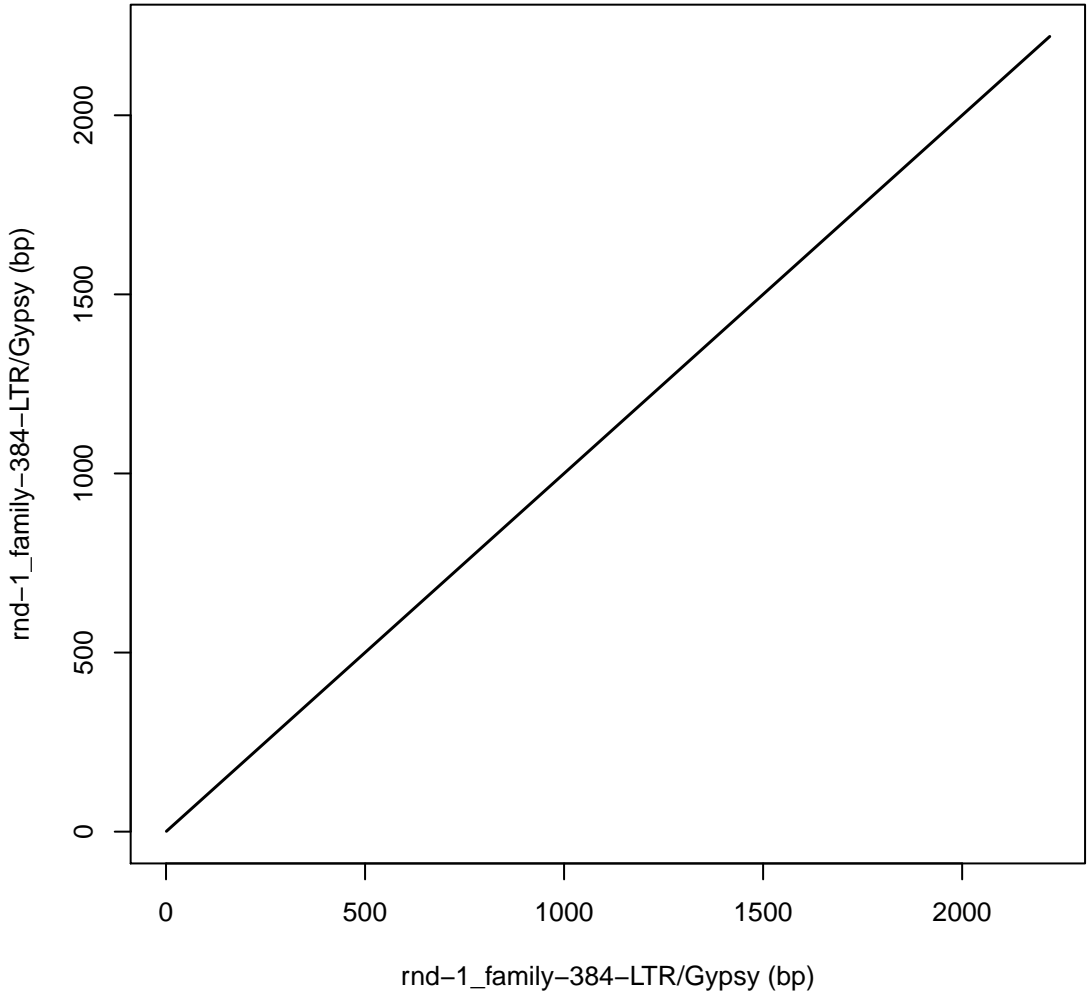
TE: rnd-1_family-384-LTR/Gypsy
consensus size: 2220bp; fragments: 55; full length: 0 (>=1998bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

