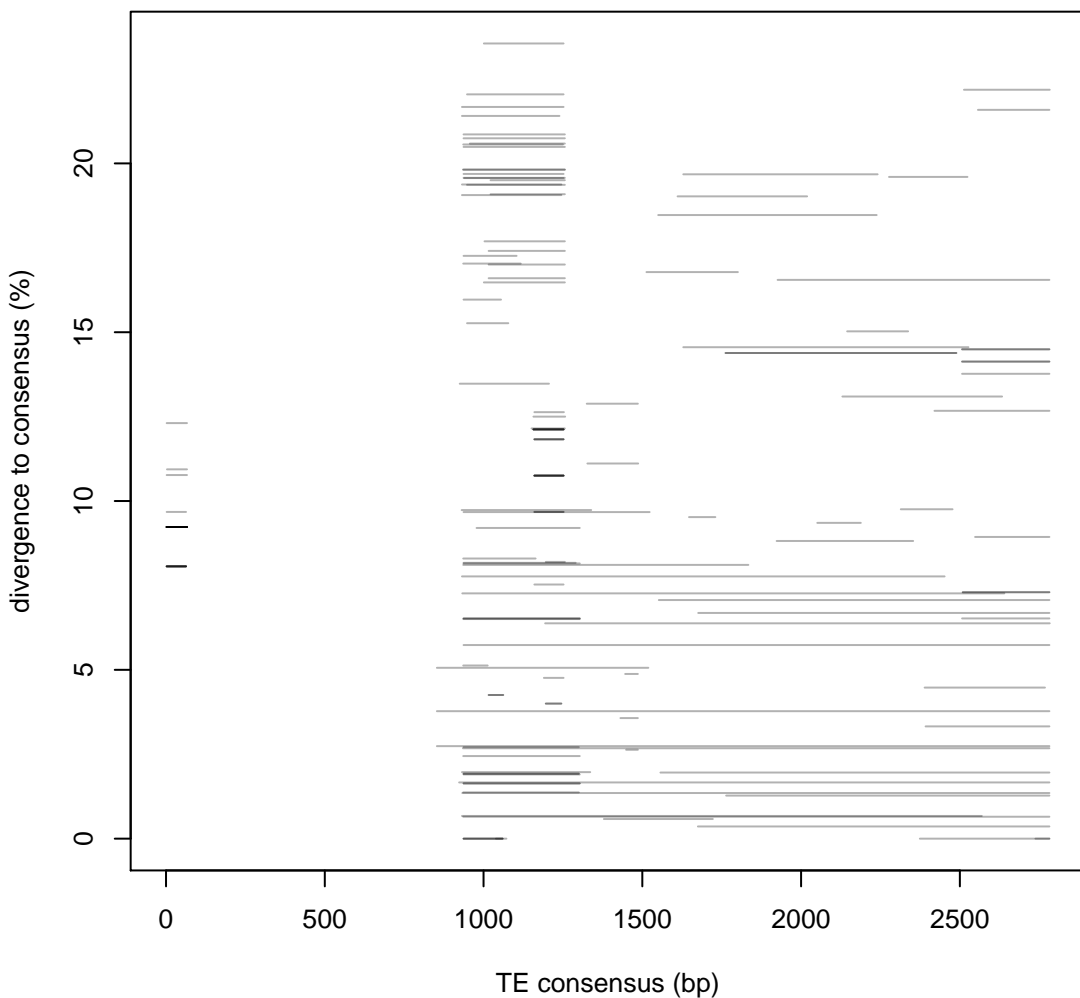
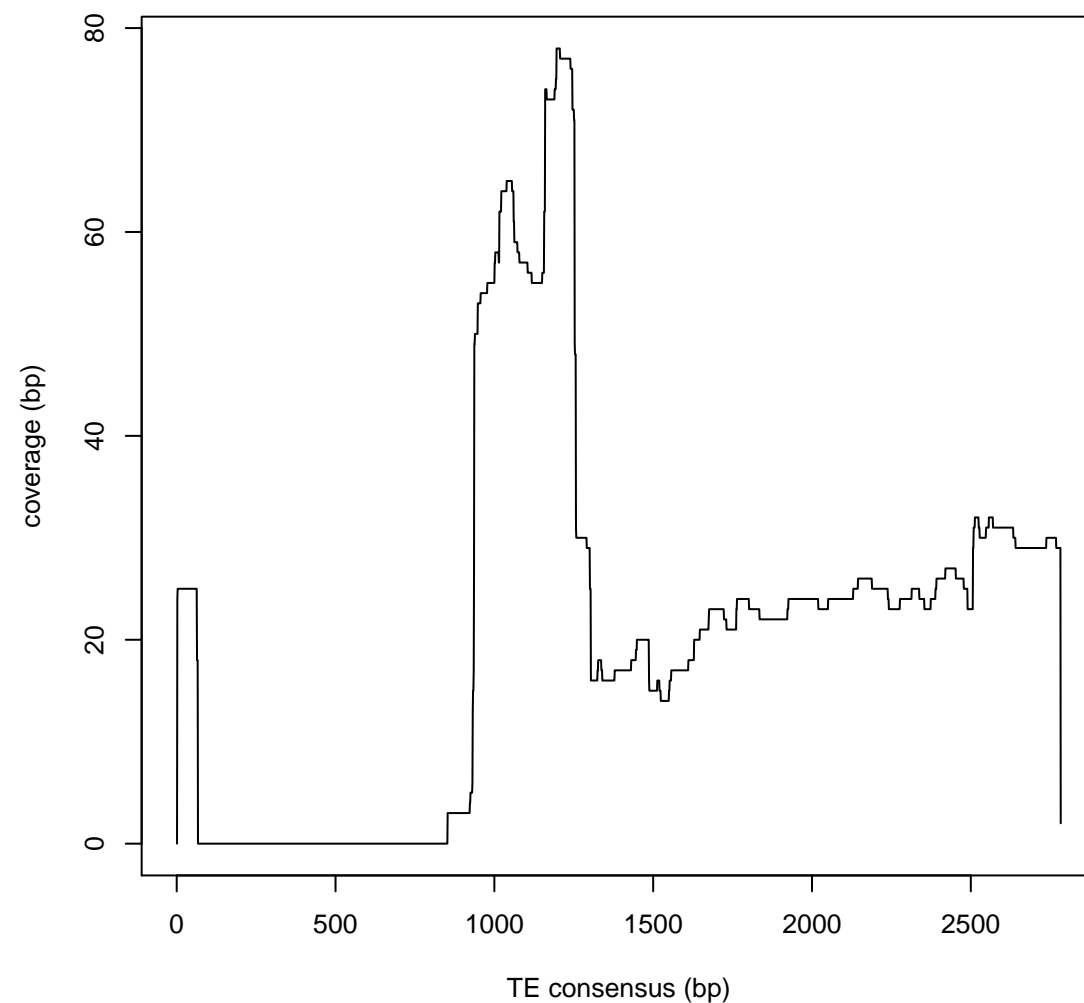


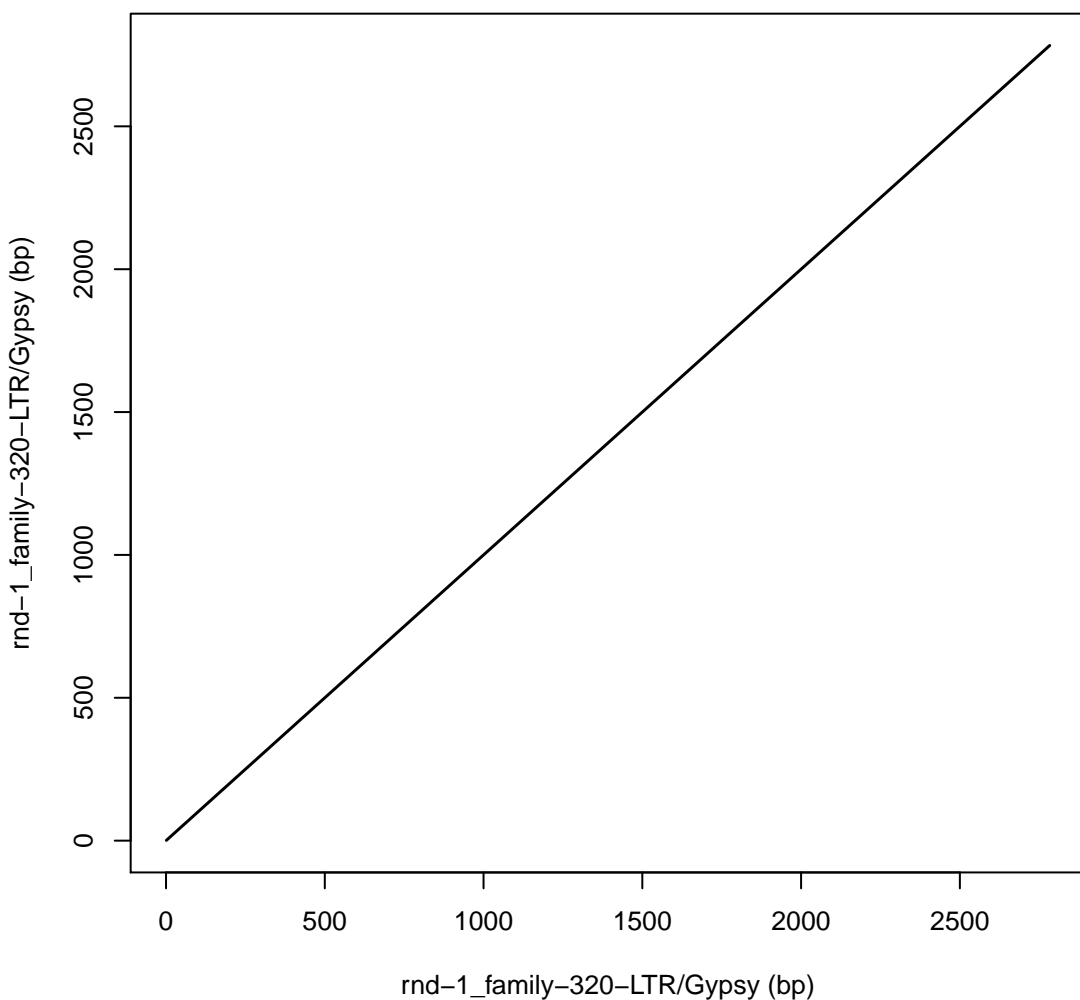
TE: rnd-1_family-320-LTR/Gypsy
consensus size: 2783bp; fragments: 157; full length: 0 (>=2504.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

