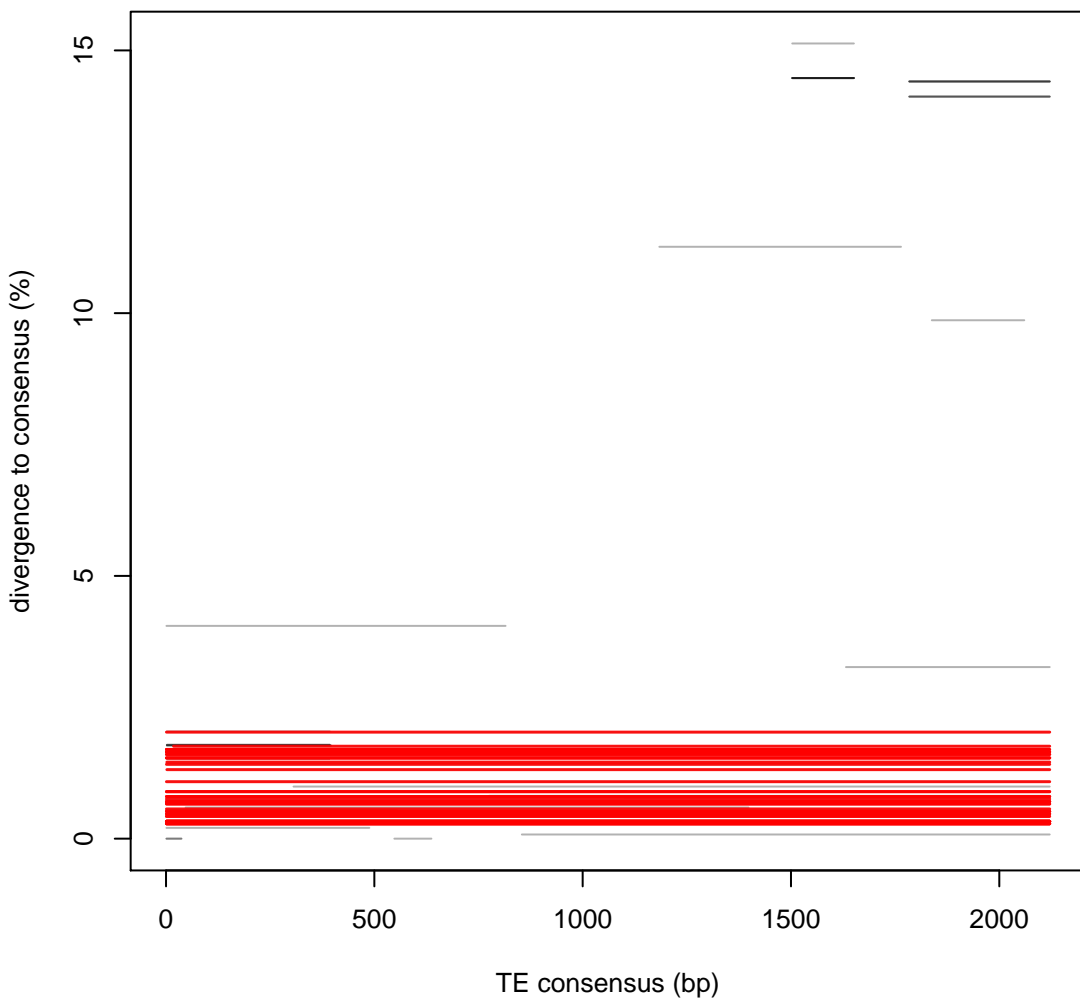
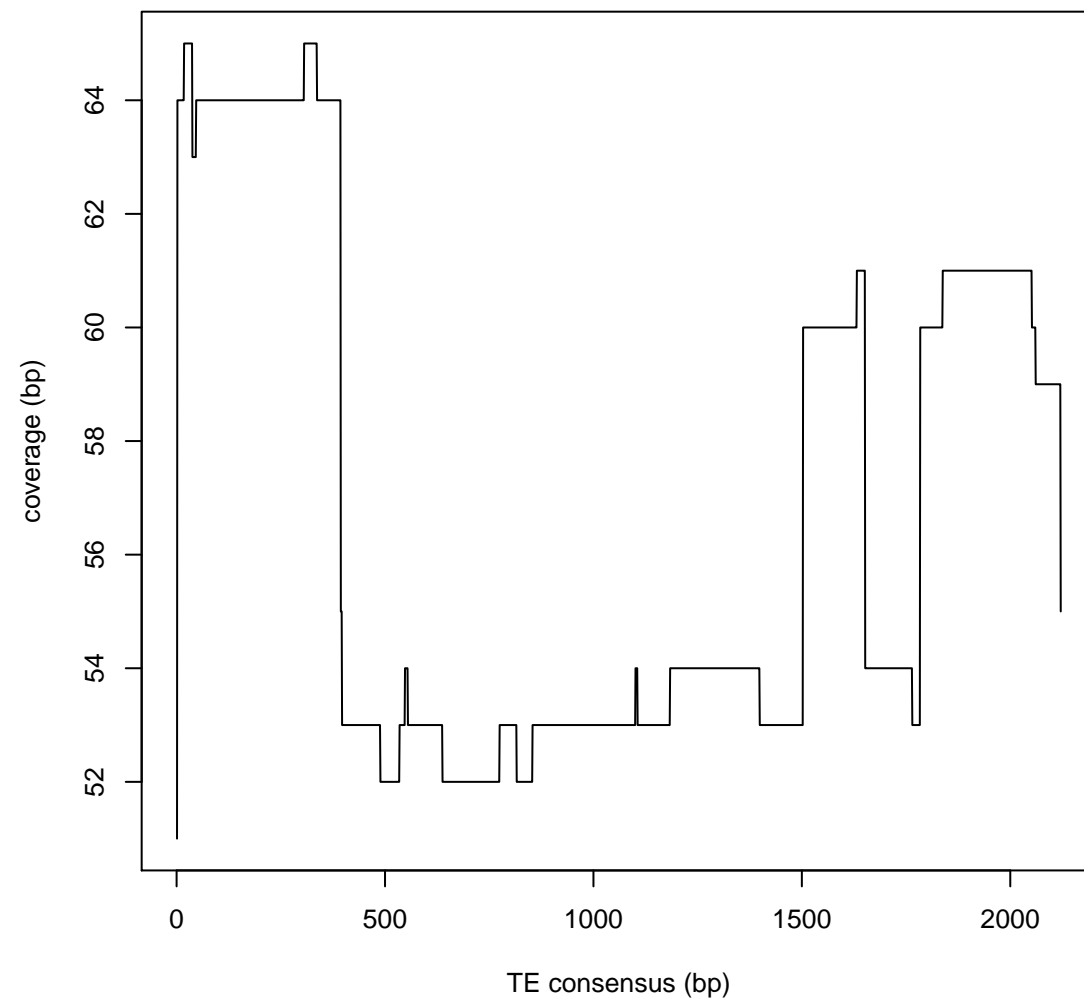


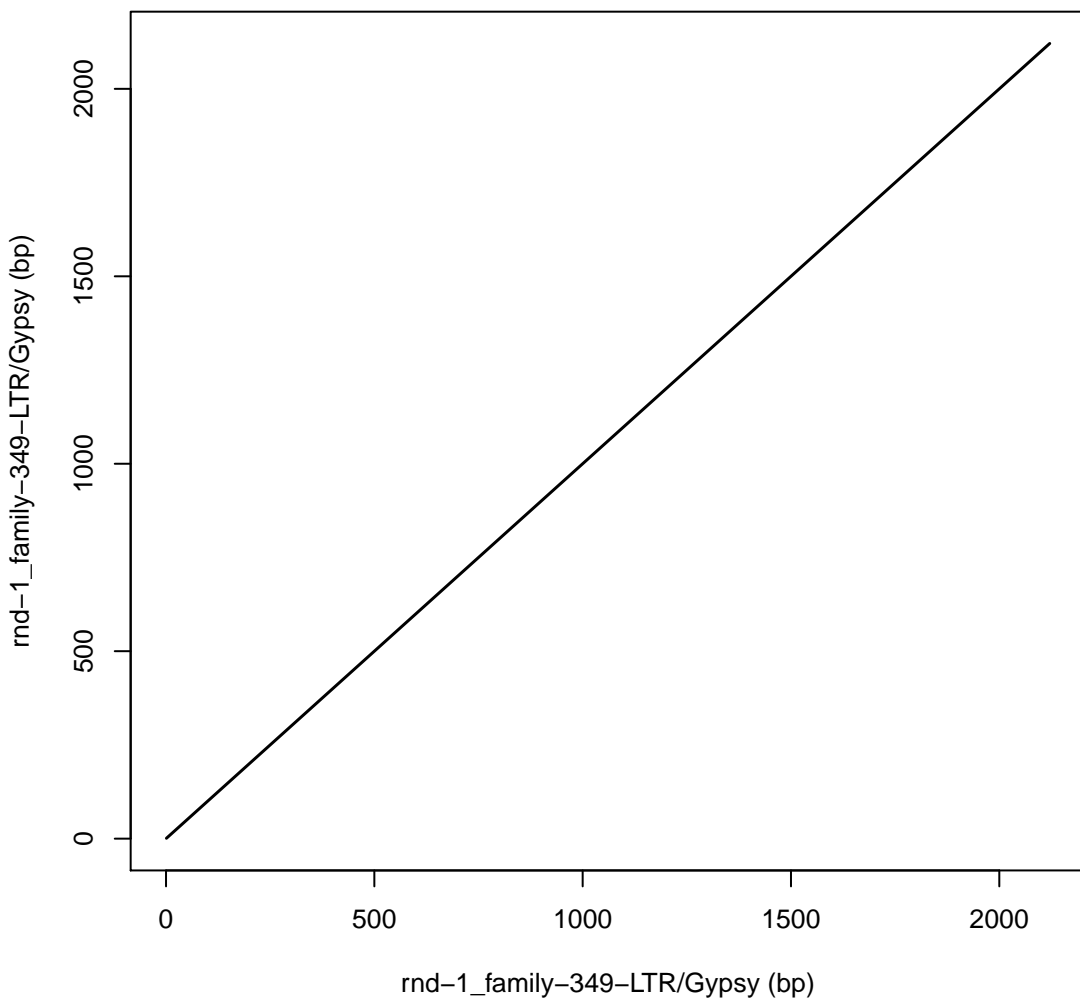
TE: rnd-1_family-349-LTR/Gypsy
consensus size: 2121bp; fragments: 89; full length: 47 (≥ 1908.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

