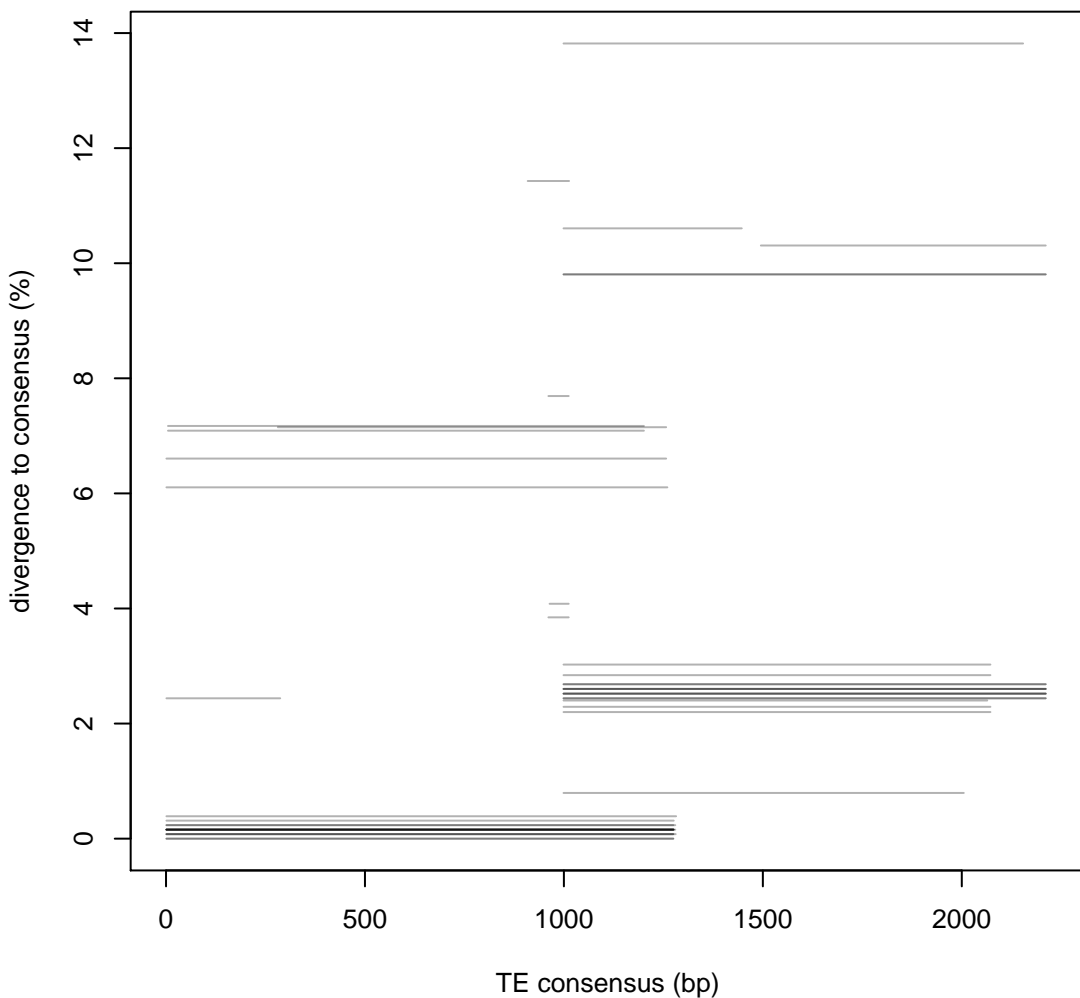
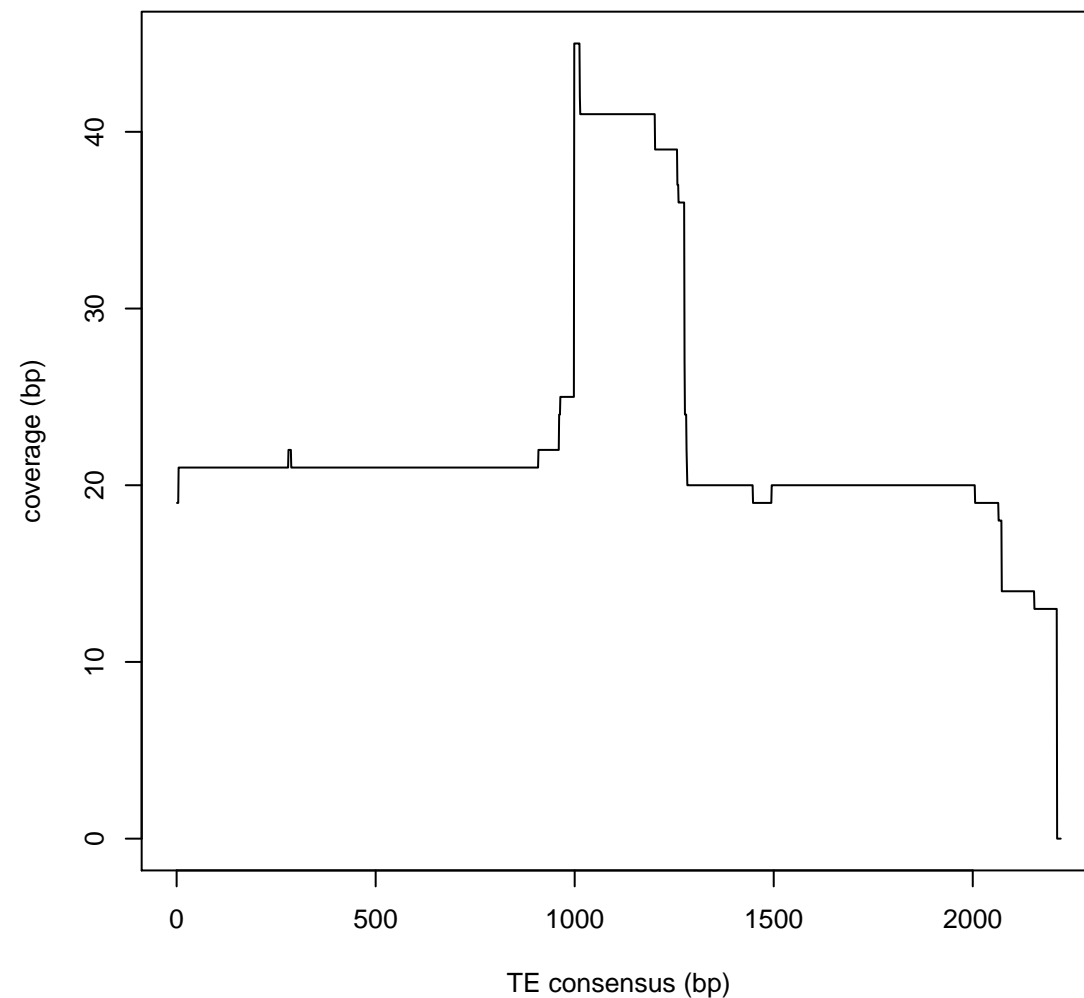


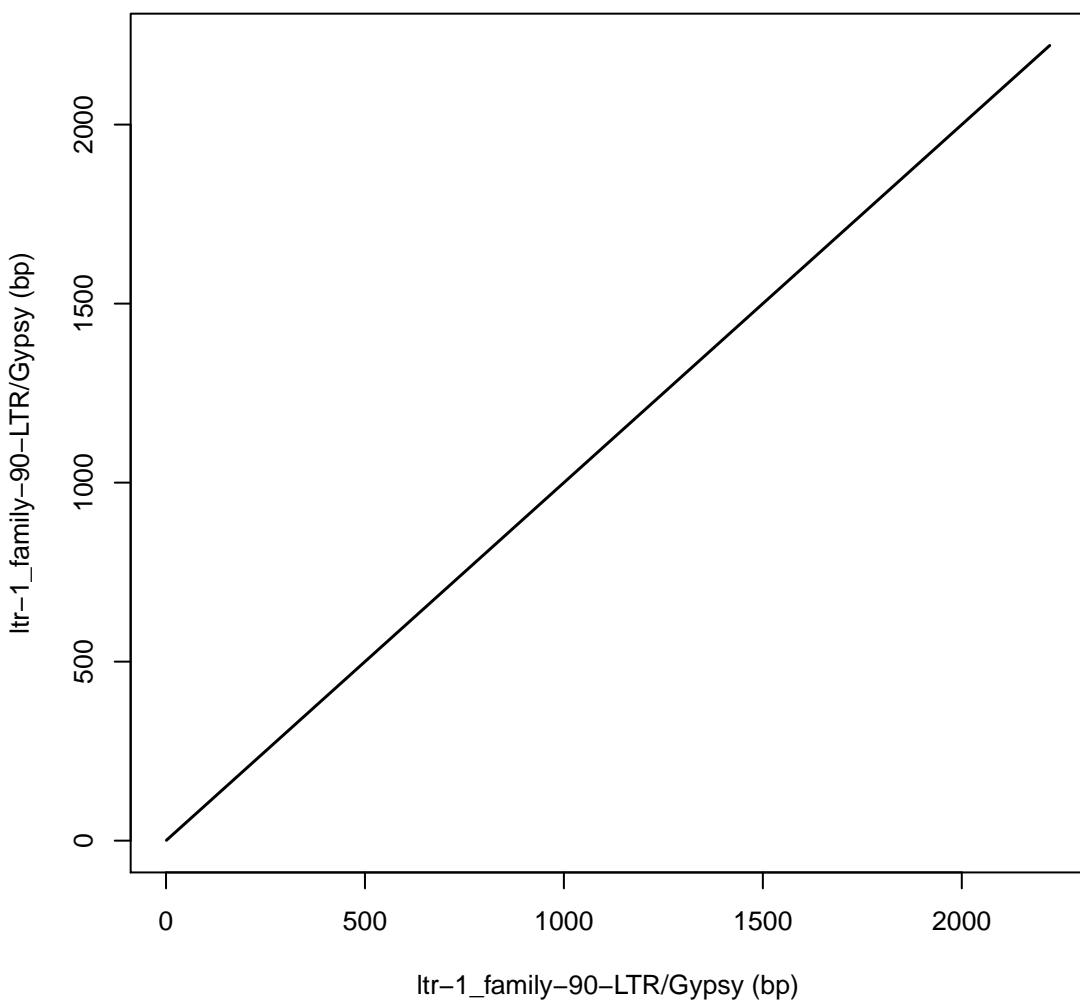
TE: ltr-1_family-90-LTR/Gypsy
consensus size: 2221bp; fragments: 47; full length: 0 (≥ 1998.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

