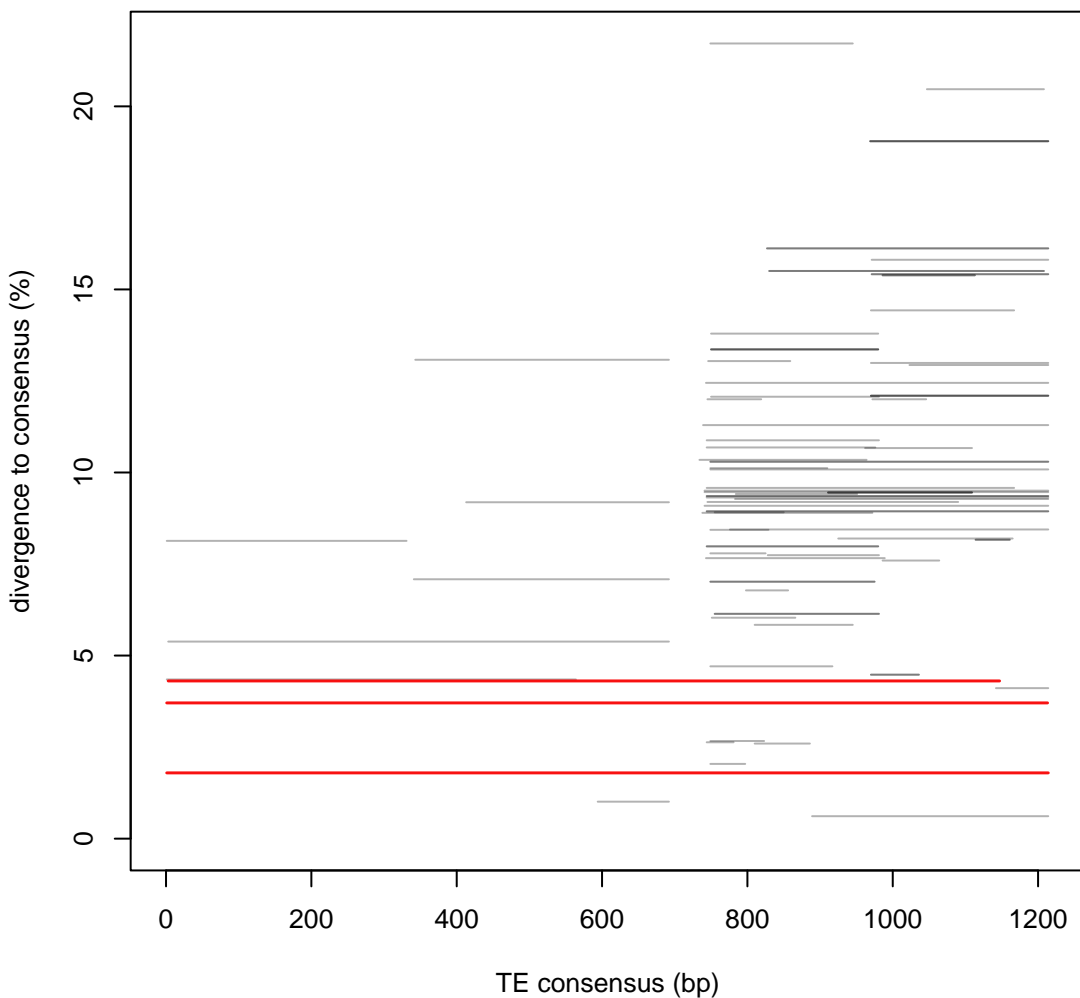
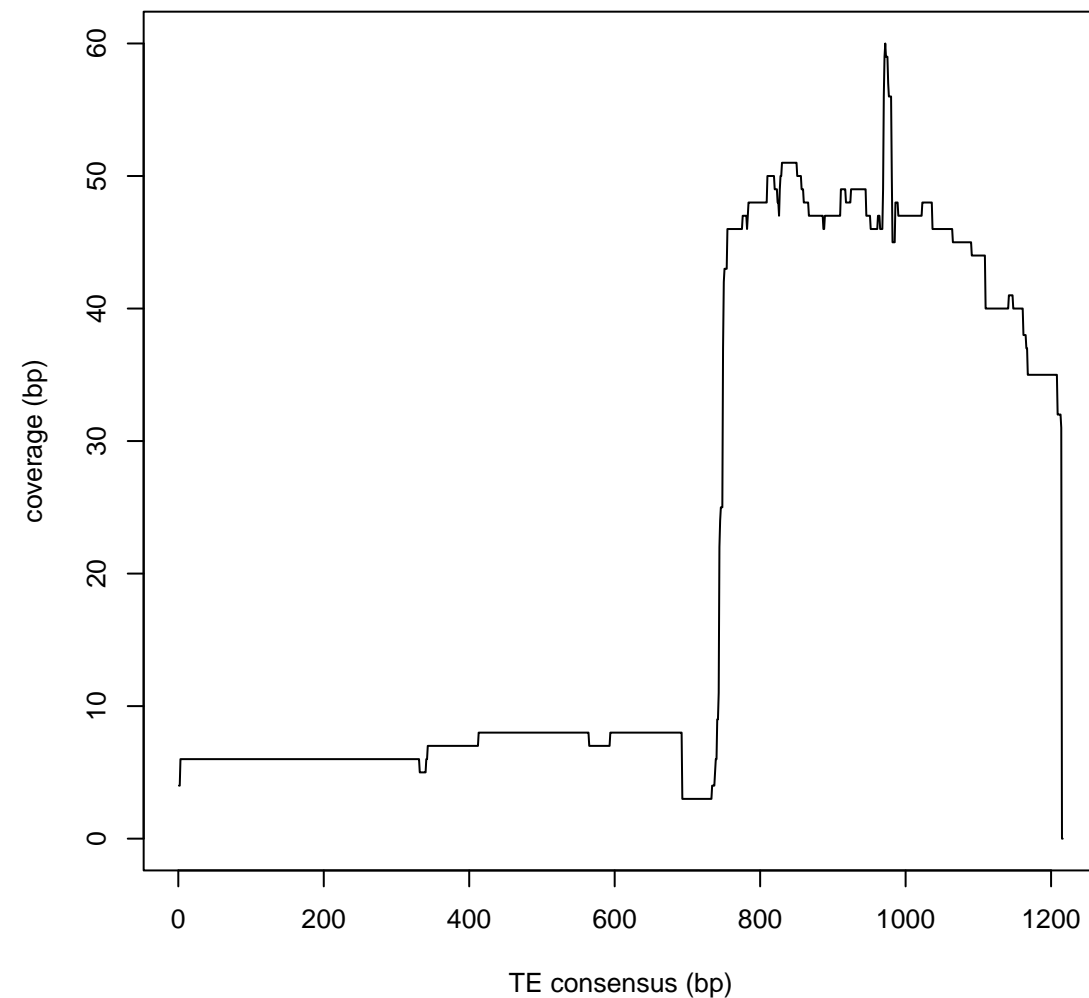


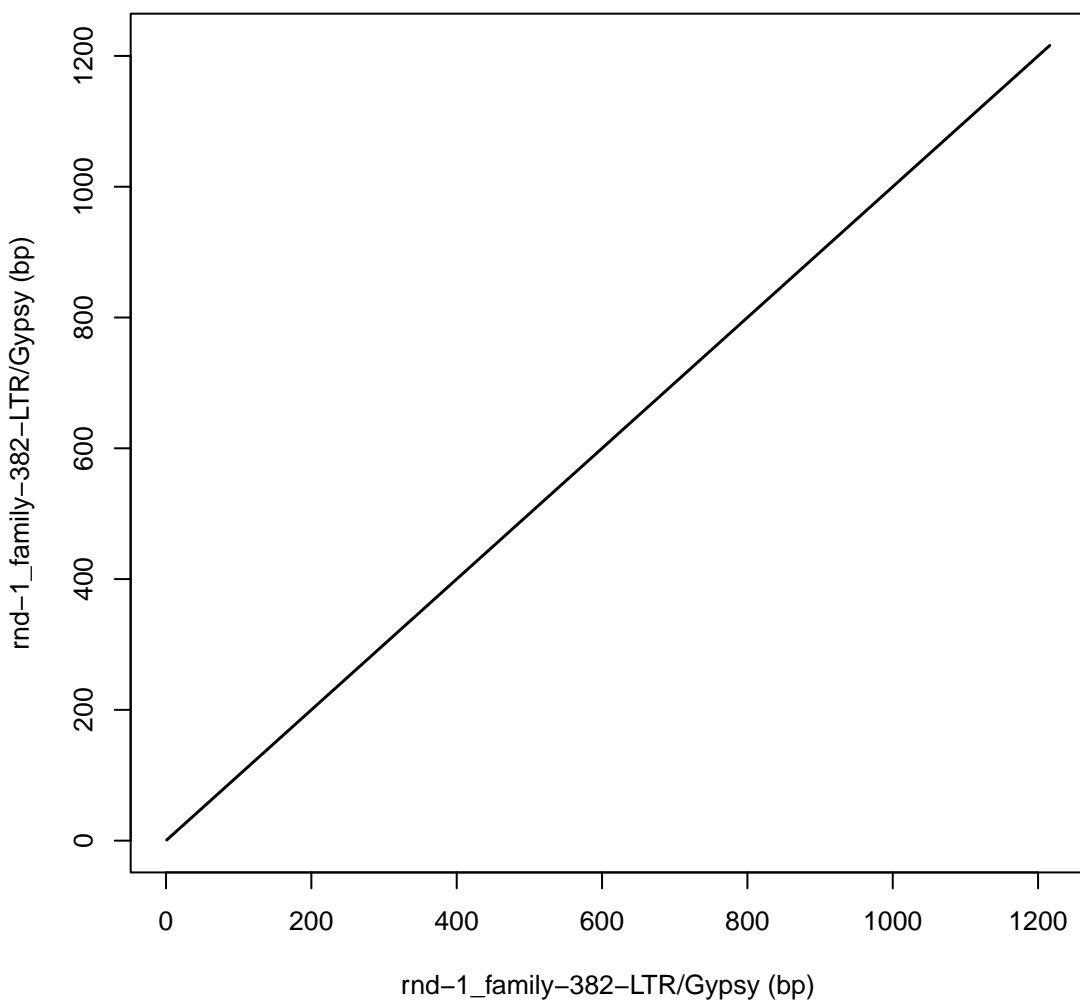
TE: rnd-1_family-382-LTR/Gypsy
consensus size: 1216bp; fragments: 92; full length: 3 (≥ 1094.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

