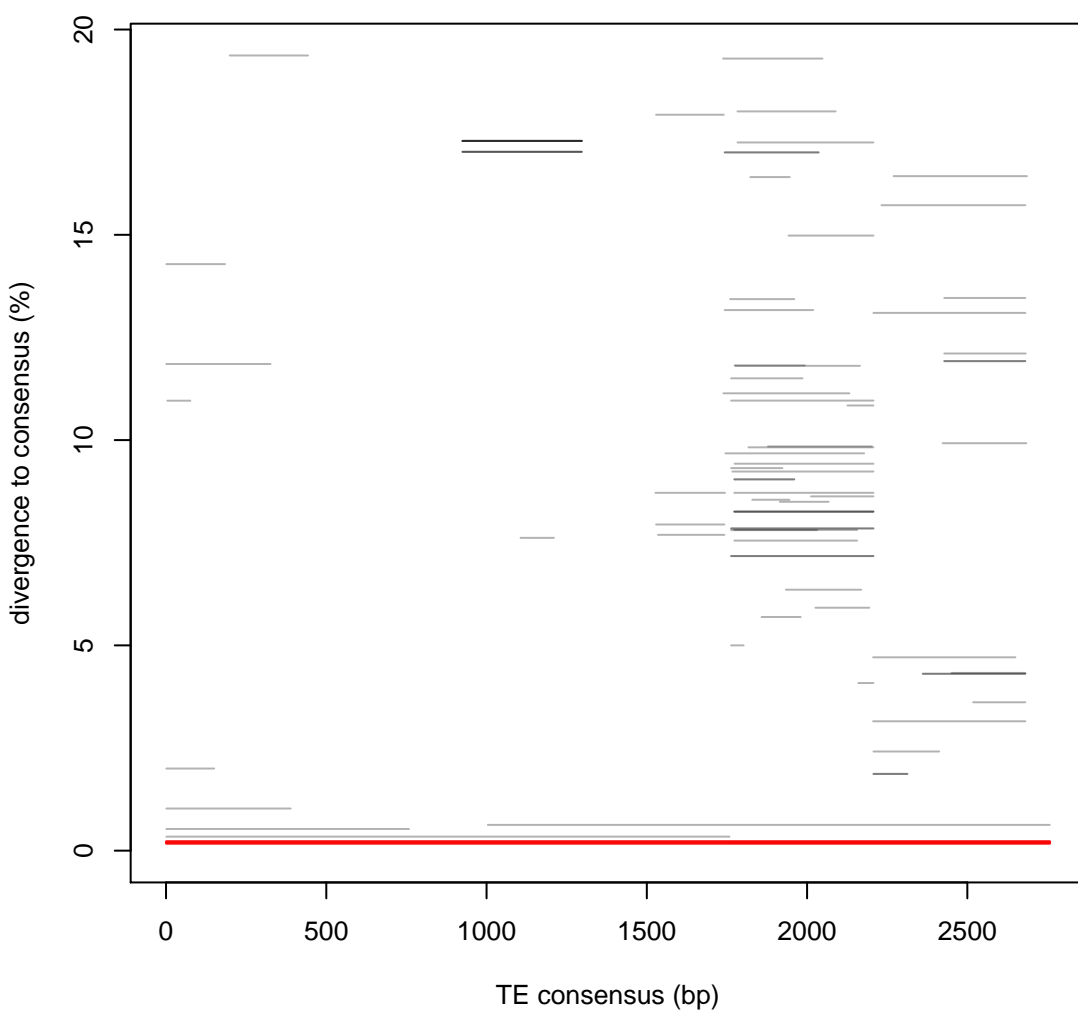
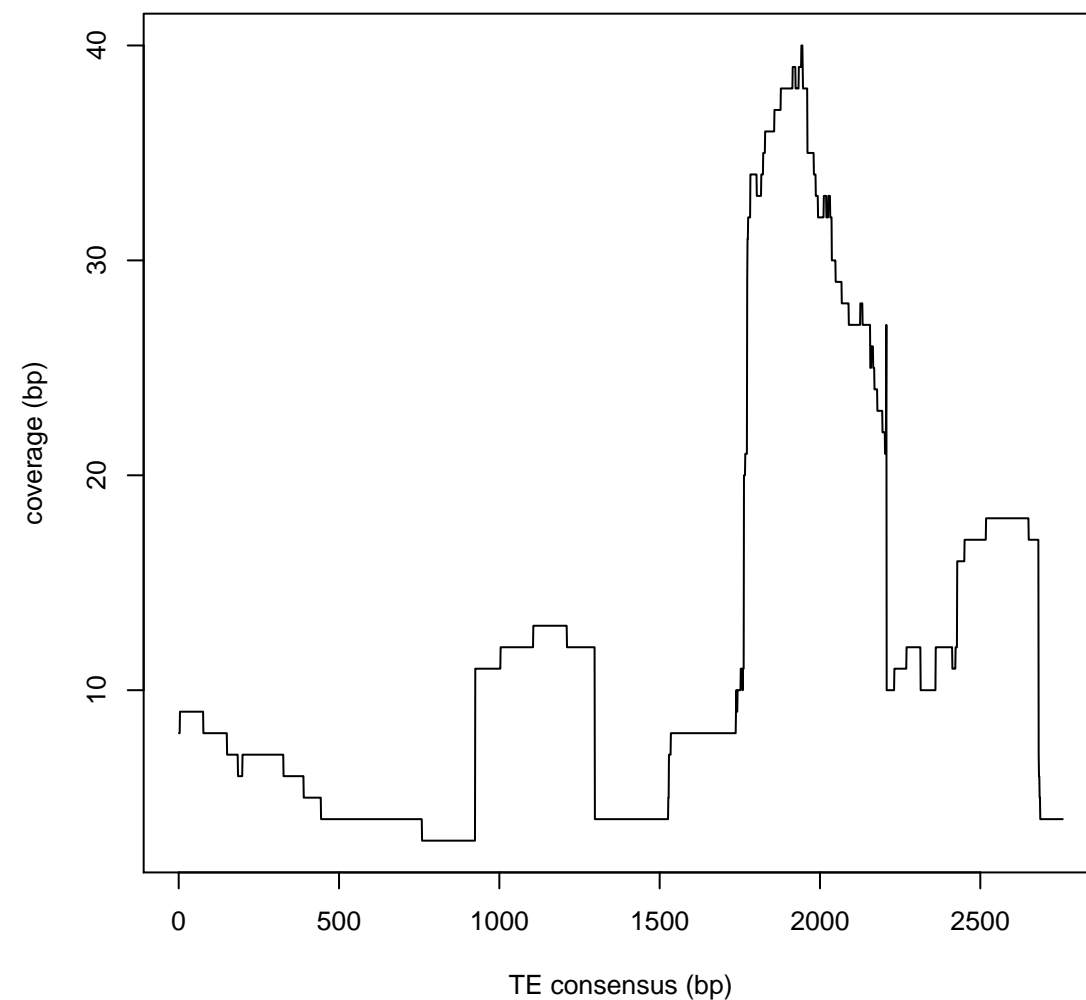


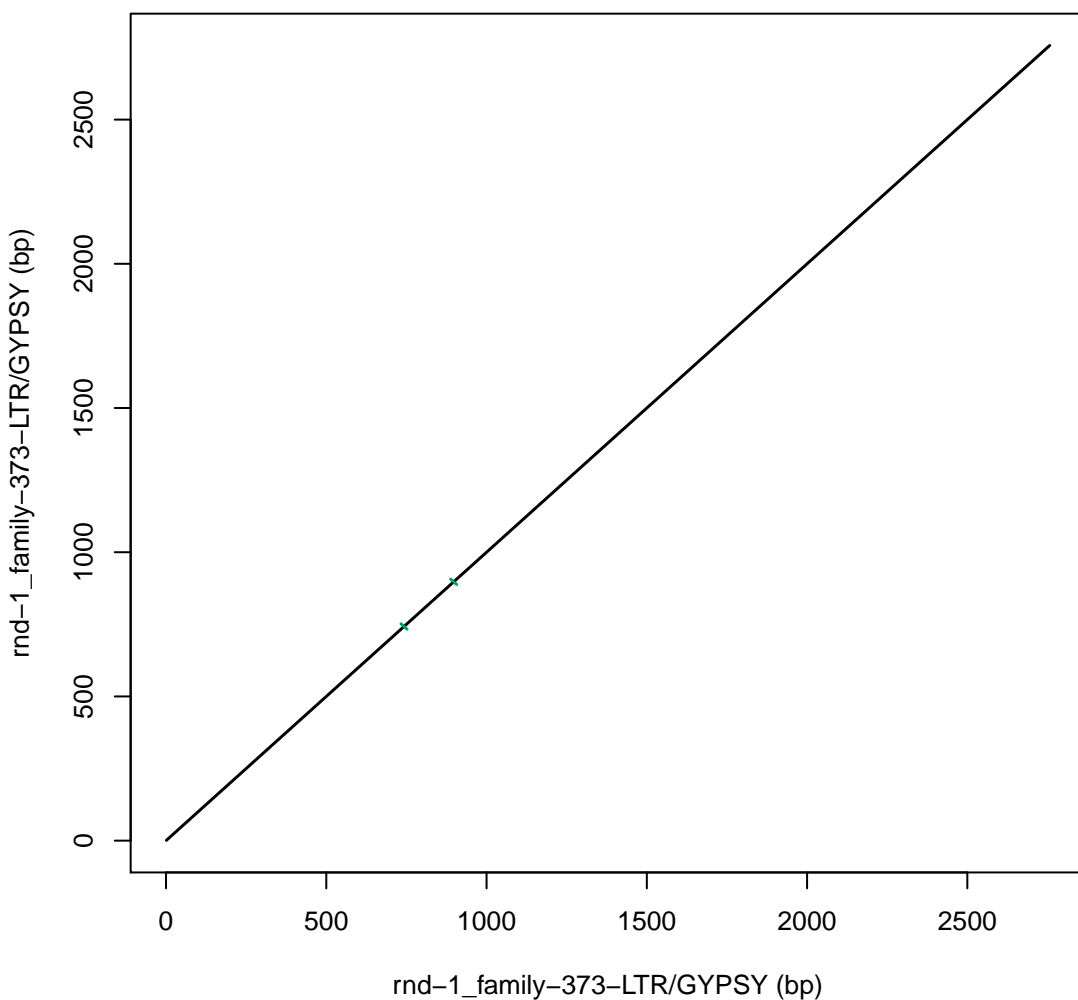
TE: rnd-1_family-373-LTR/GYPSY
consensus size: 2757bp; fragments: 84; full length: 2 (≥ 2481.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

