TE: rnd−5_family−845−LTR/GYPSY
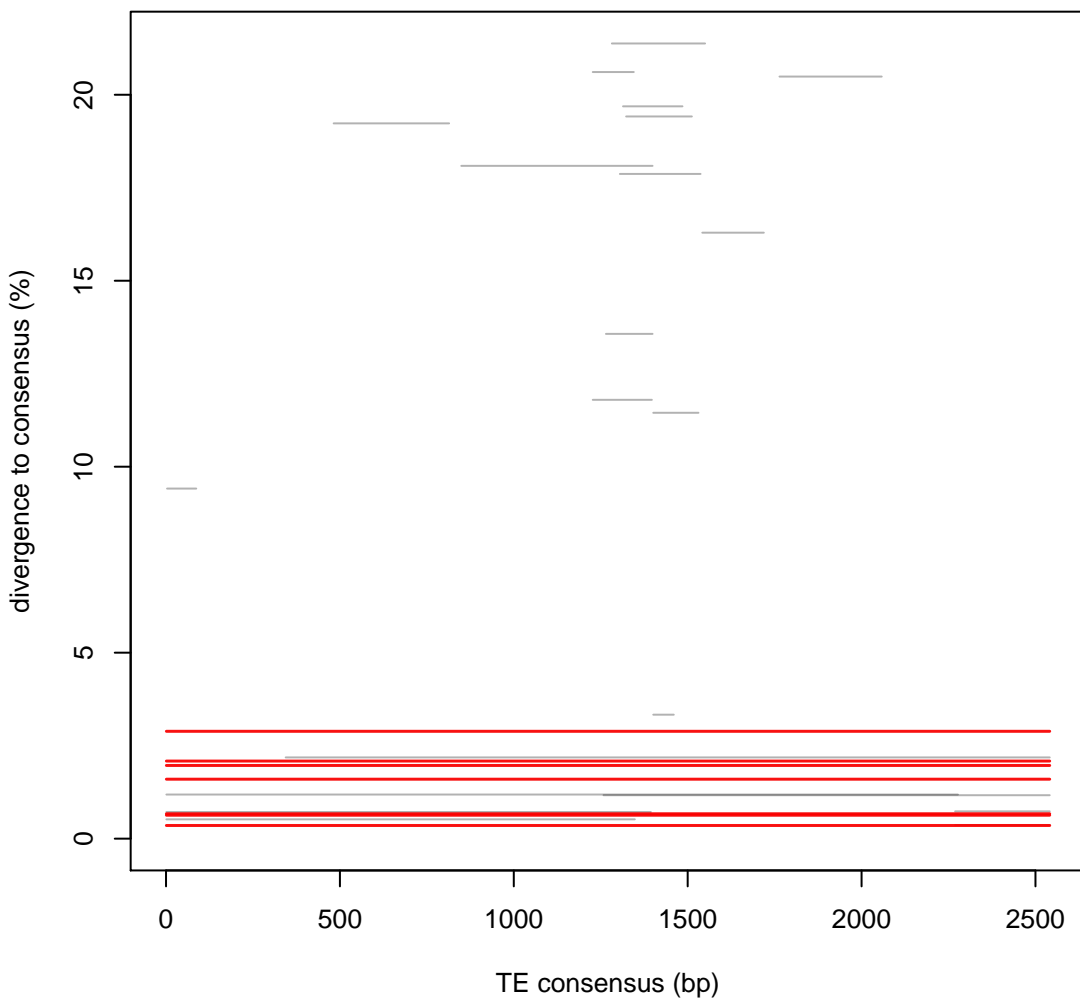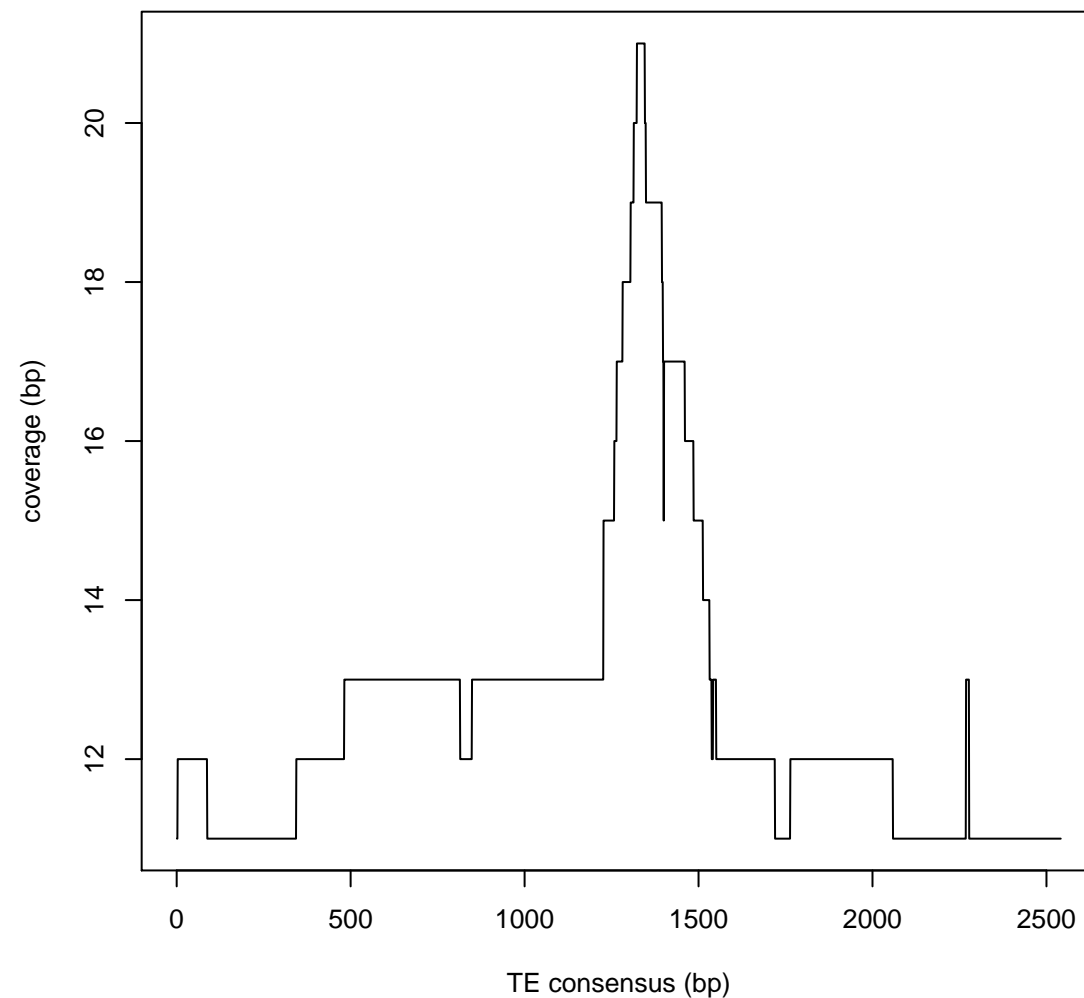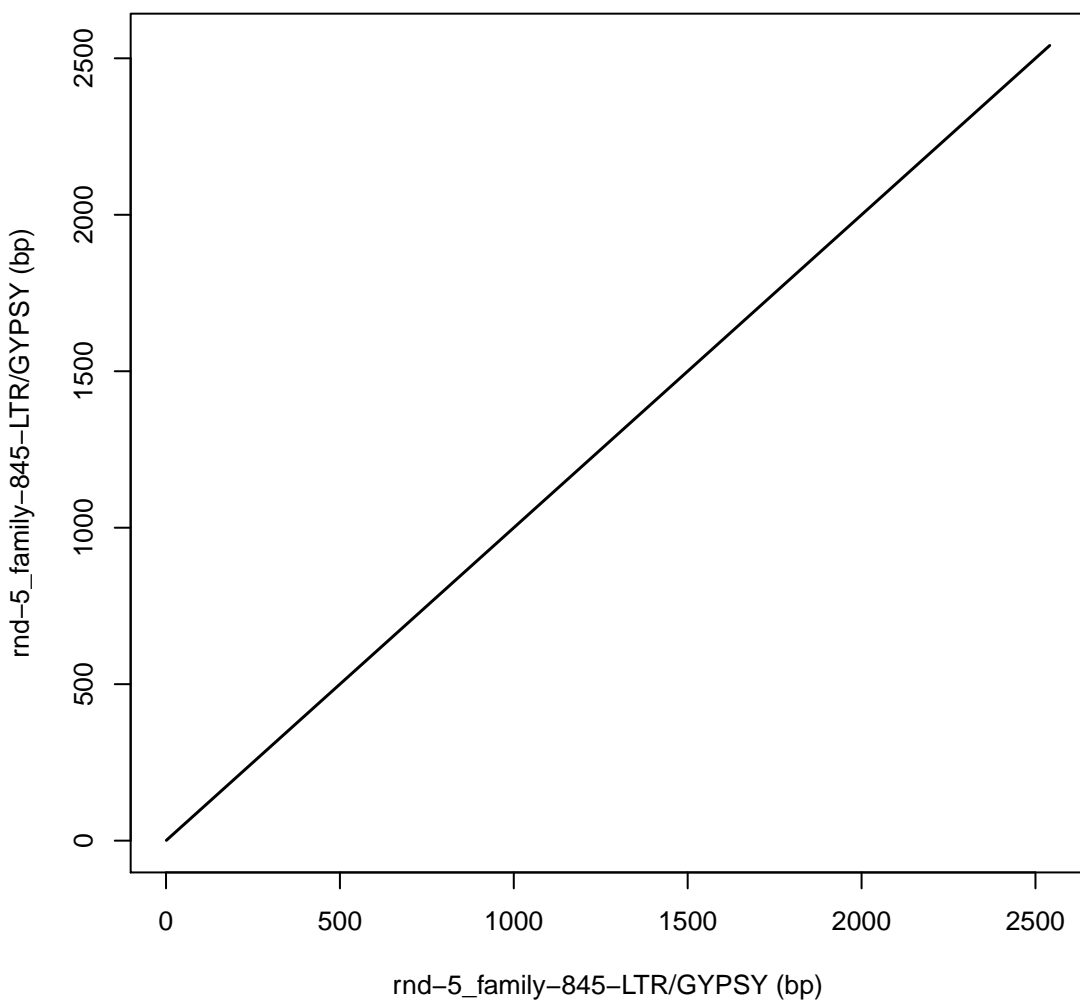consensus size: 2541bp; fragments: 29; full length: 7 (>=2286.9bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected