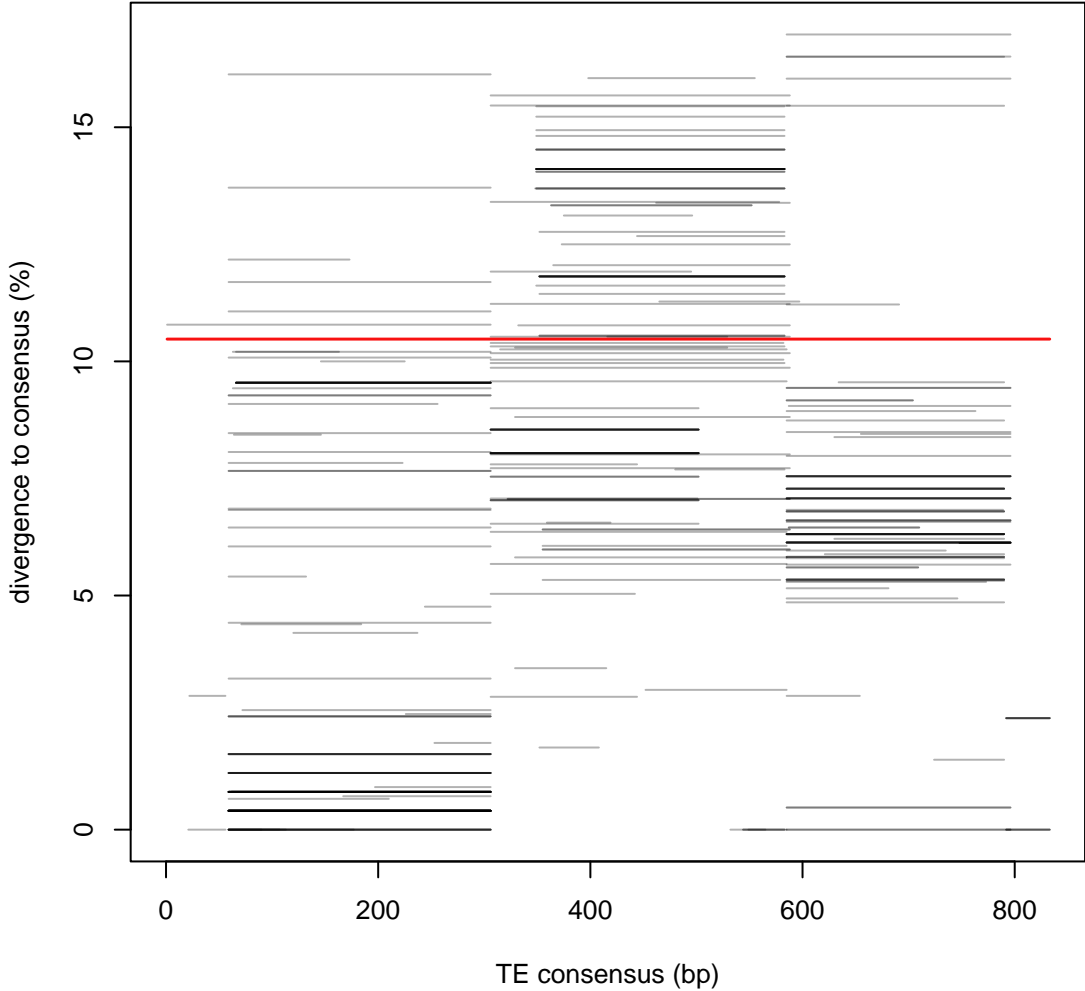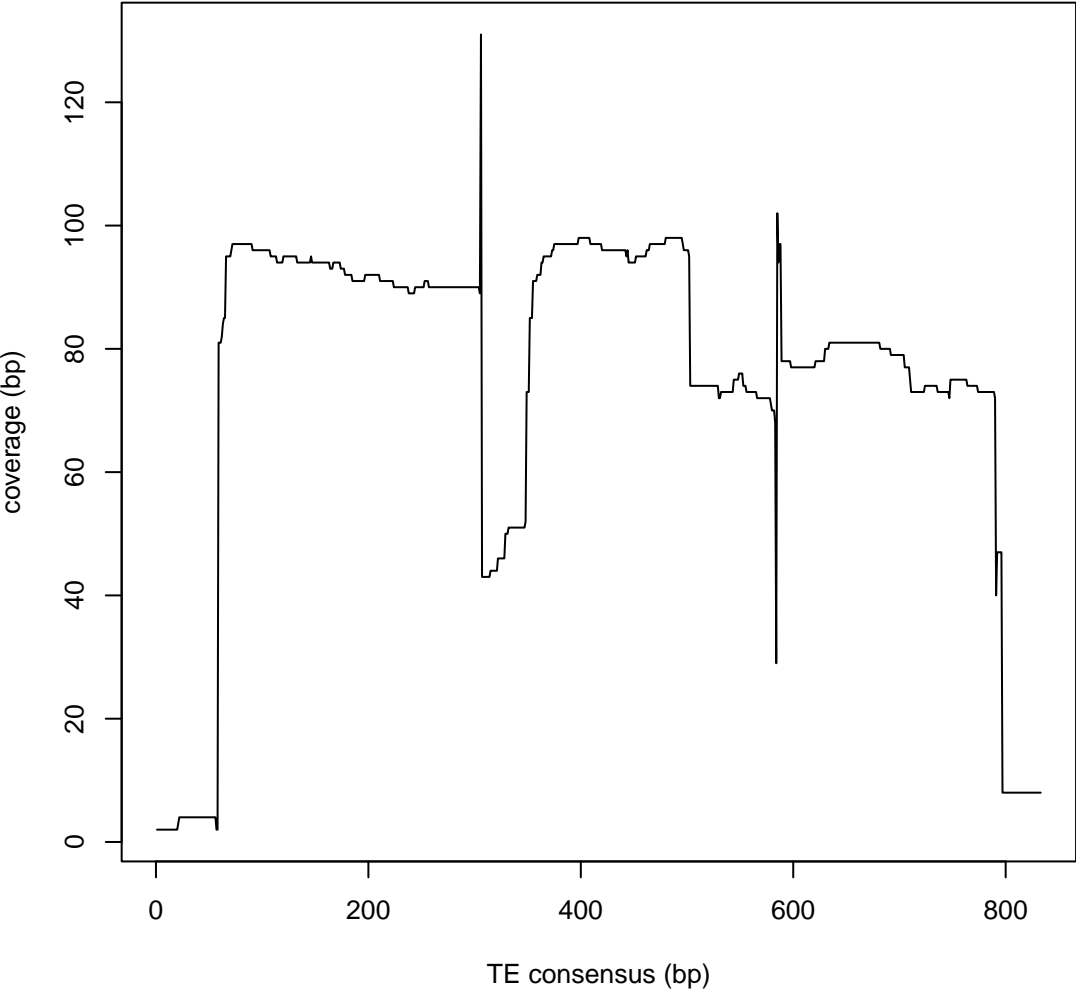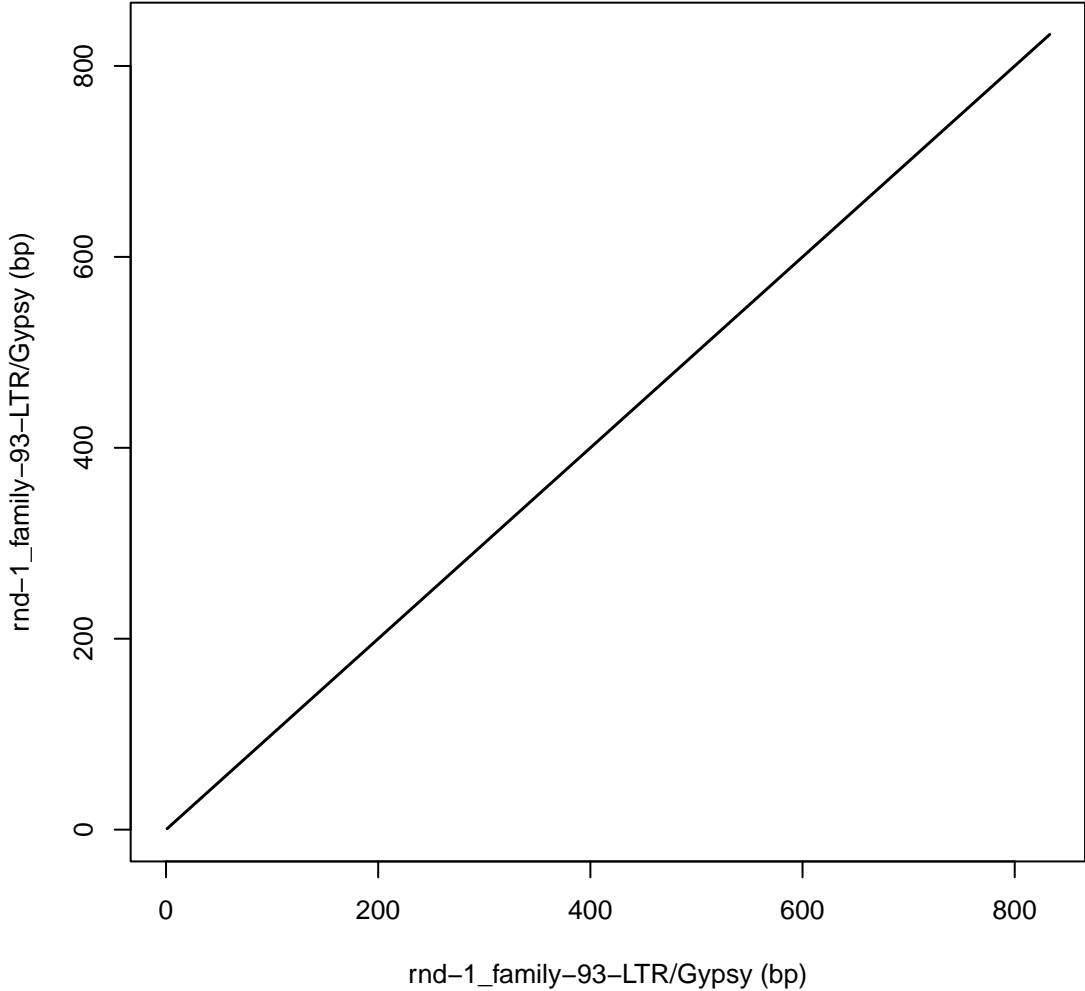**TE: rnd−1_family−93−LTR/Gypsy**
consensus size: 833bp; fragments: 305; full length: 1 (>=749.7bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected