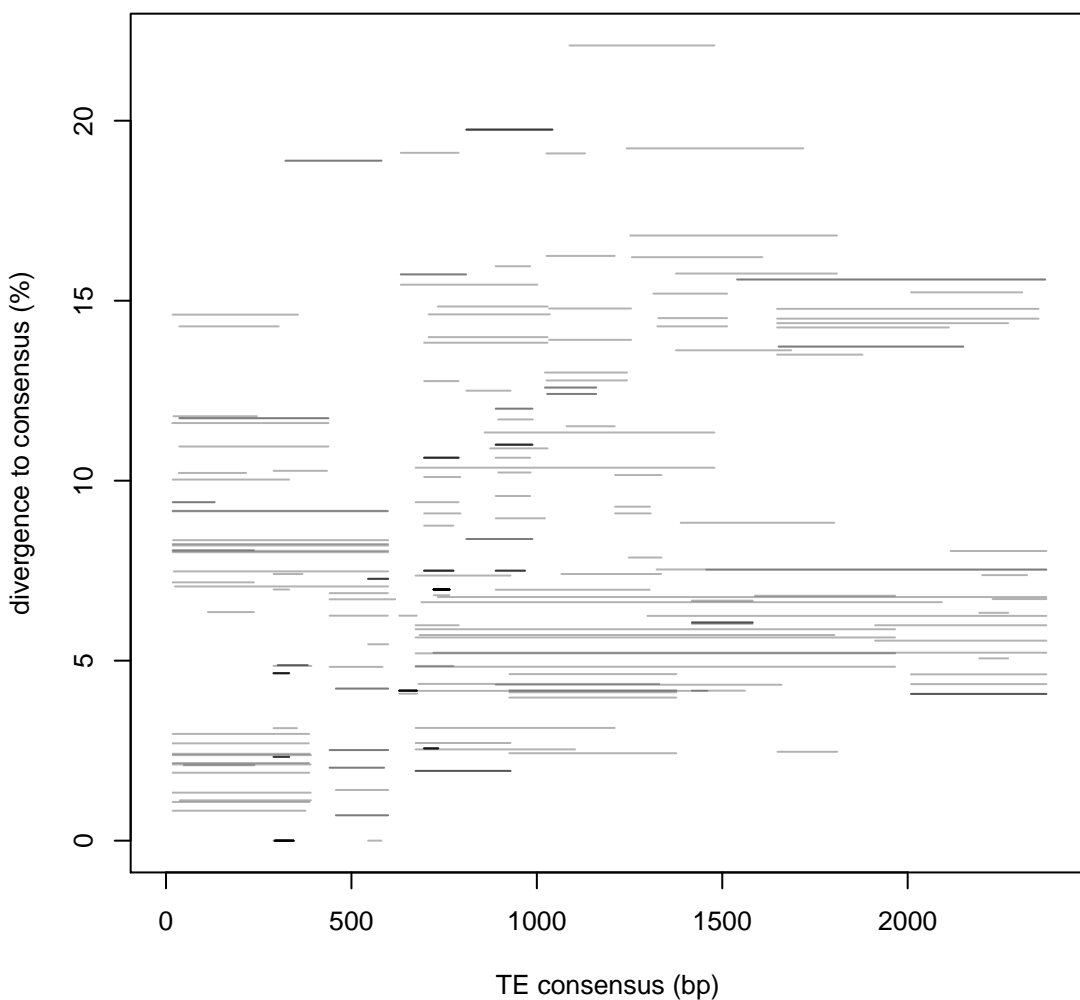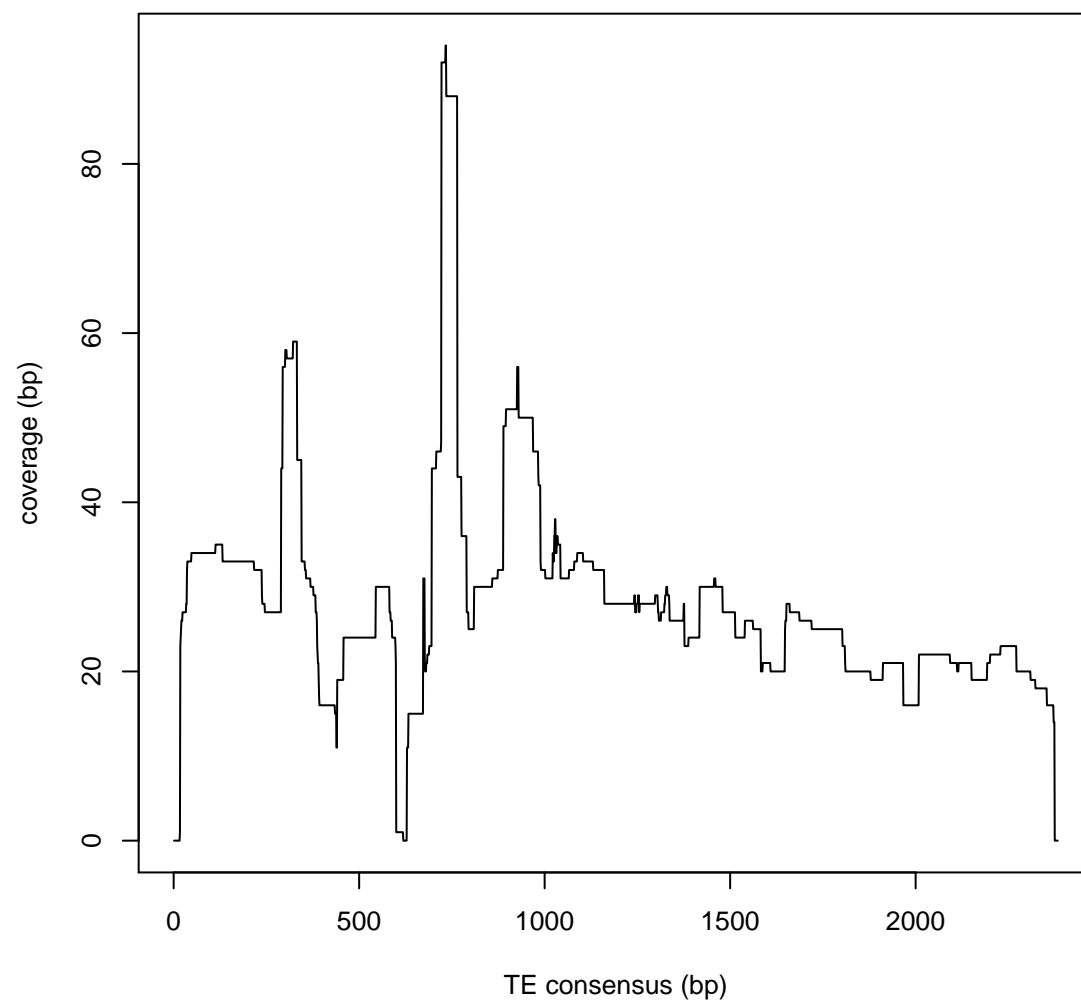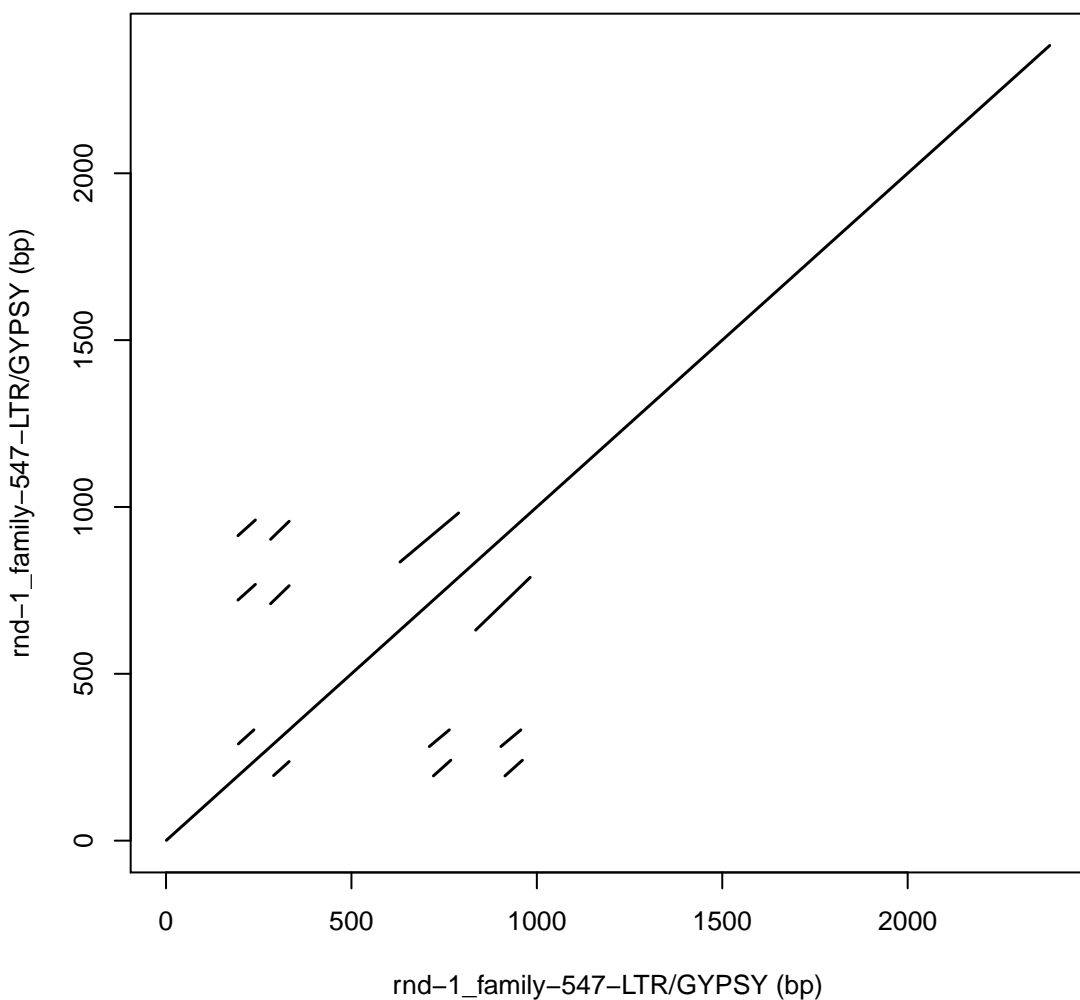TE: rnd−1_family−547−LTR/GYPSY
consensus size: 2383bp; fragments: 284; full length: 0 (>=2144.7bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected