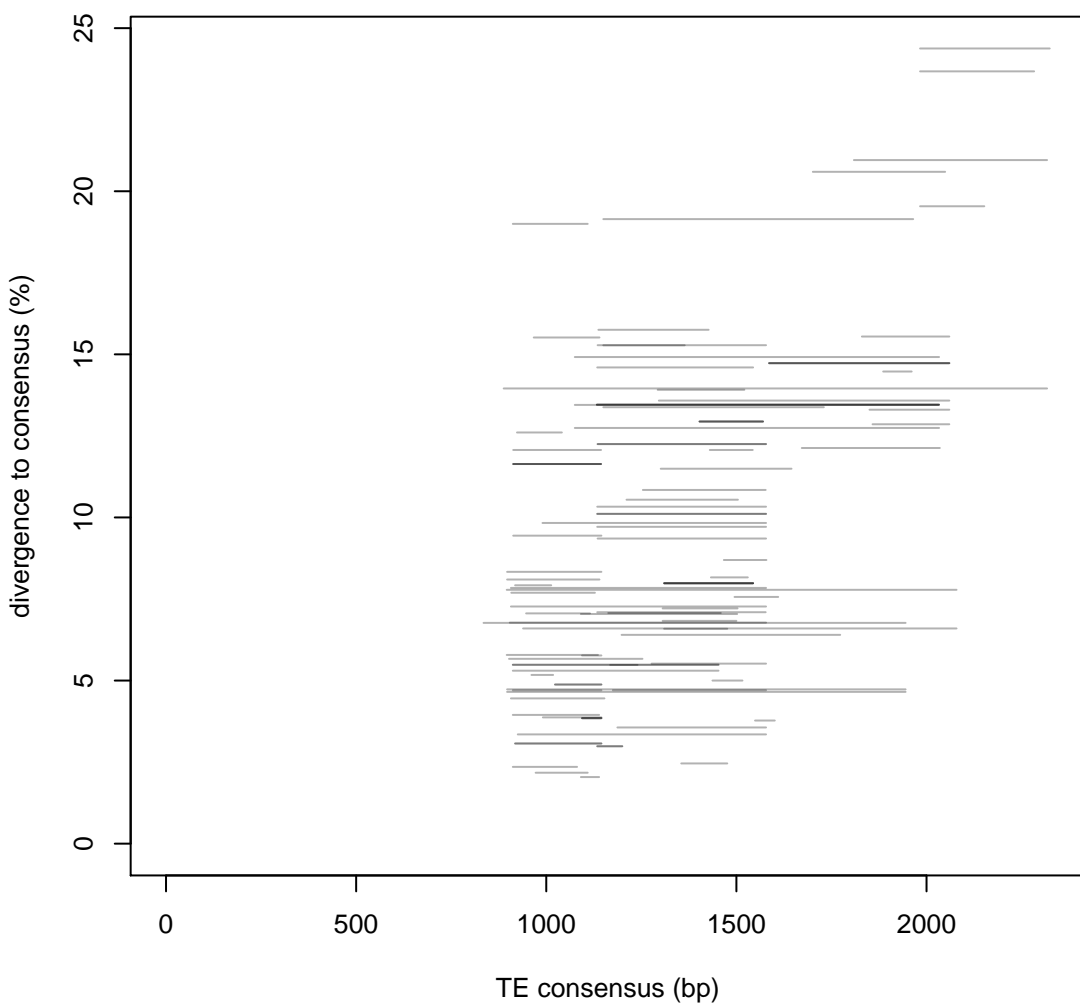
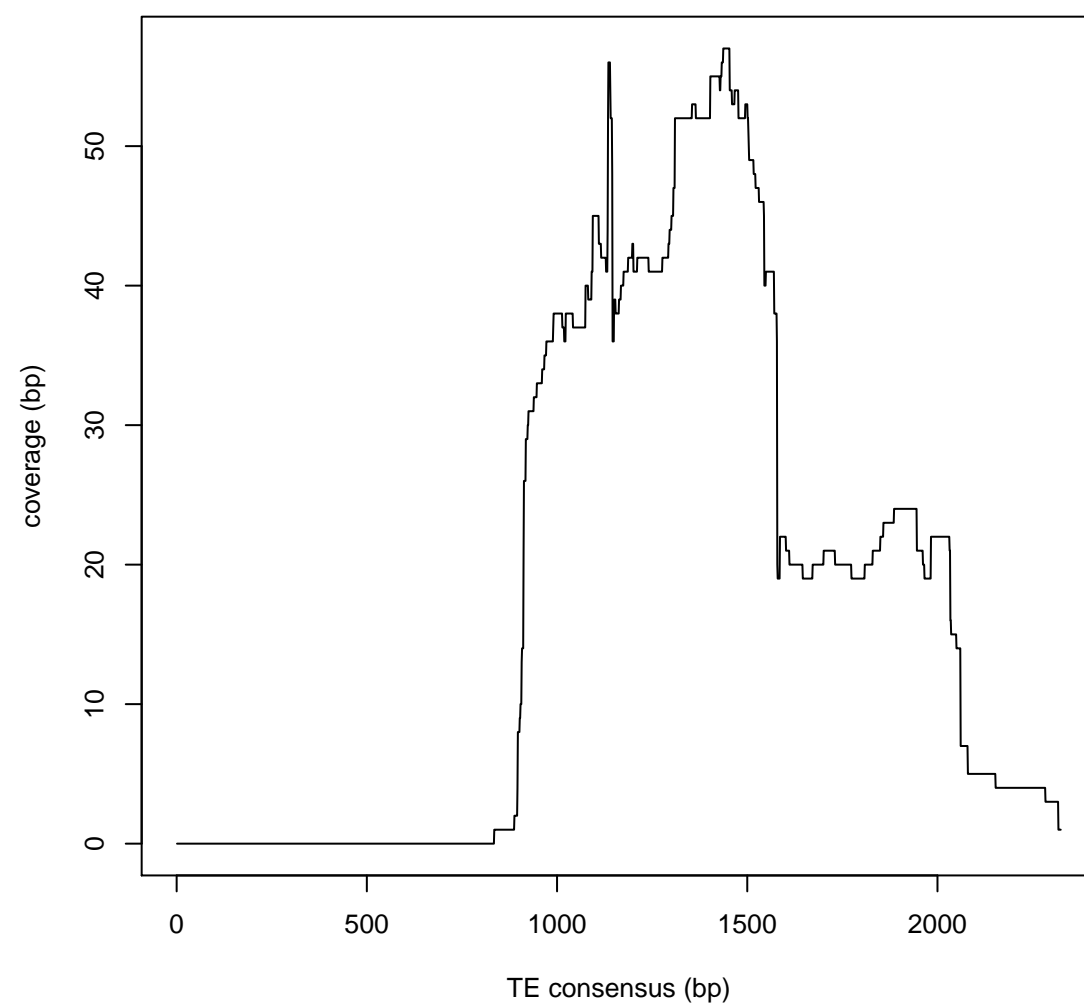


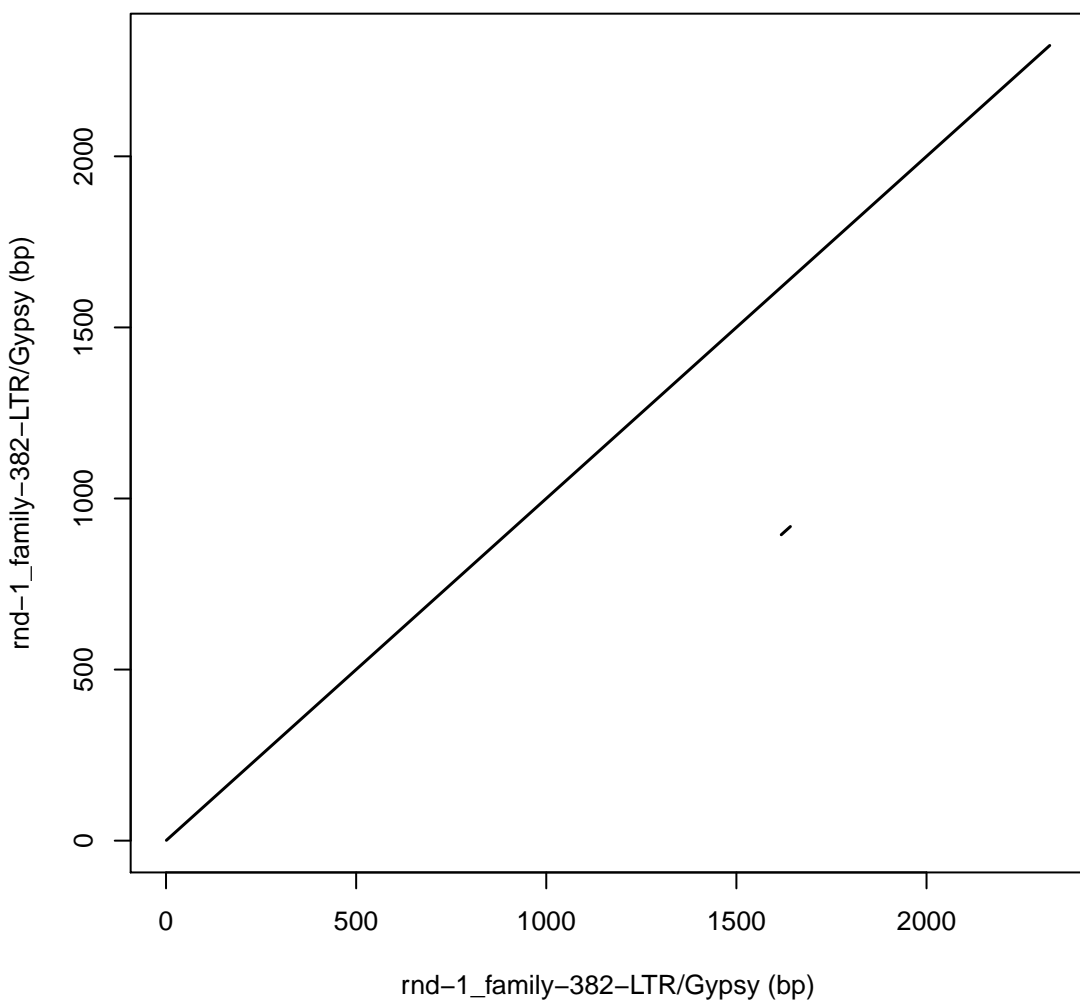
TE: rnd-1_family-382-LTR/Gypsy
consensus size: 2324bp; fragments: 109; full length: 0 (>=2091.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

no orf >400 bp detected

