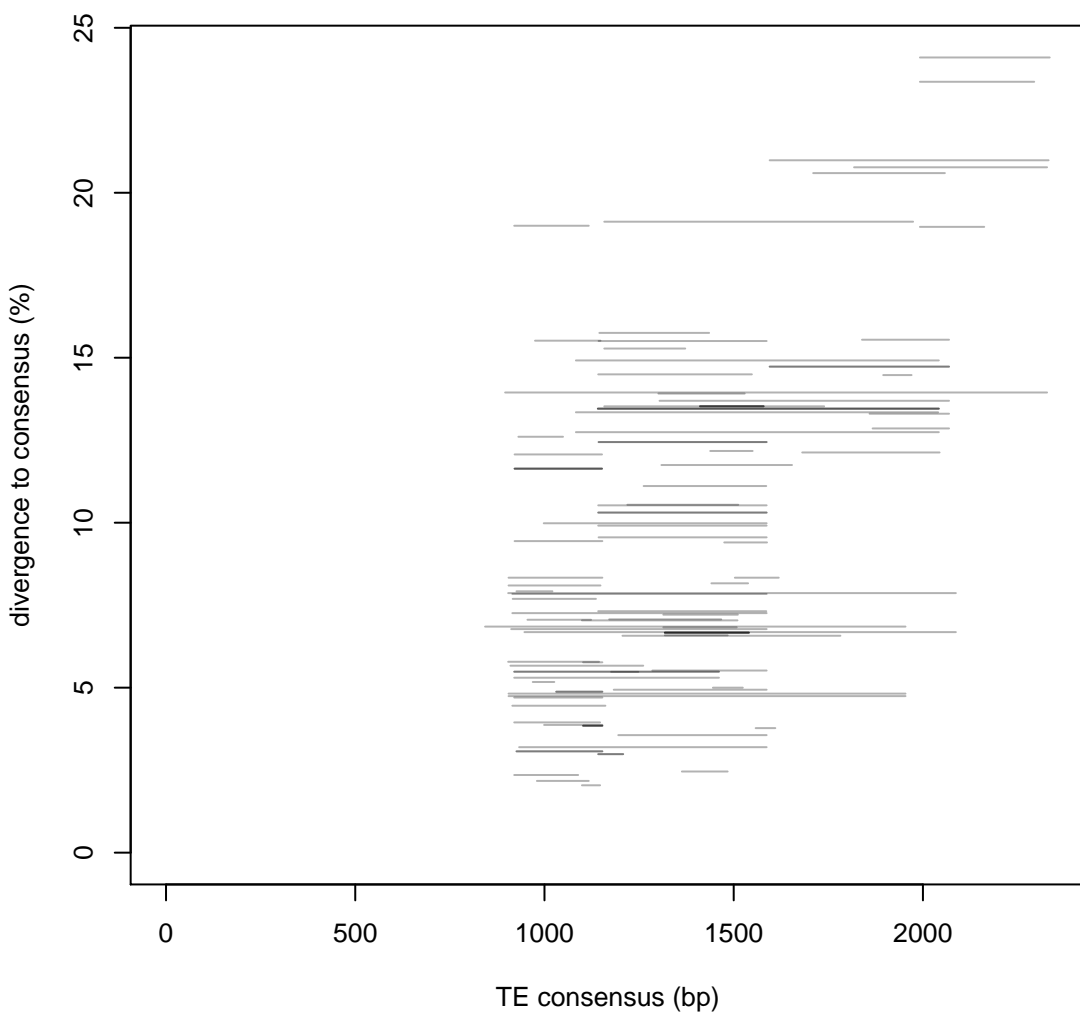
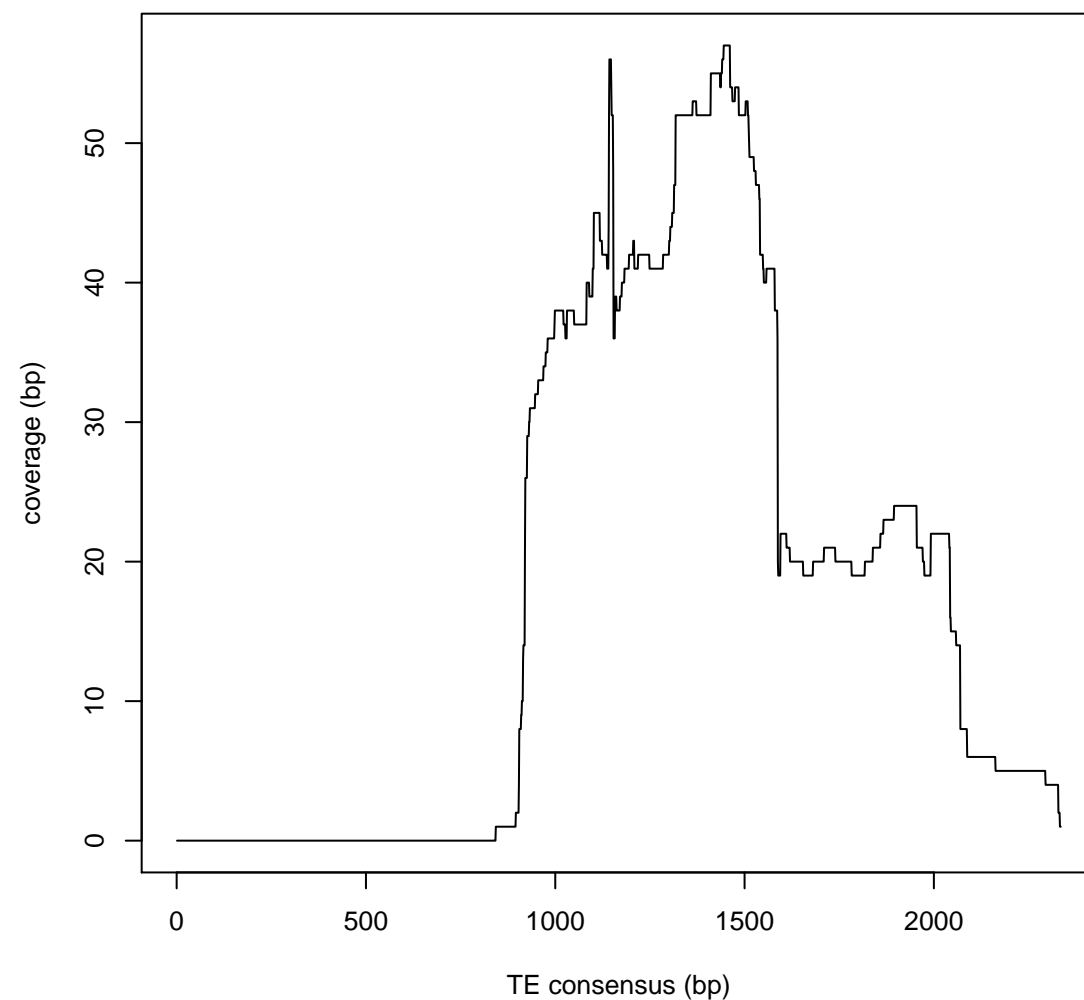


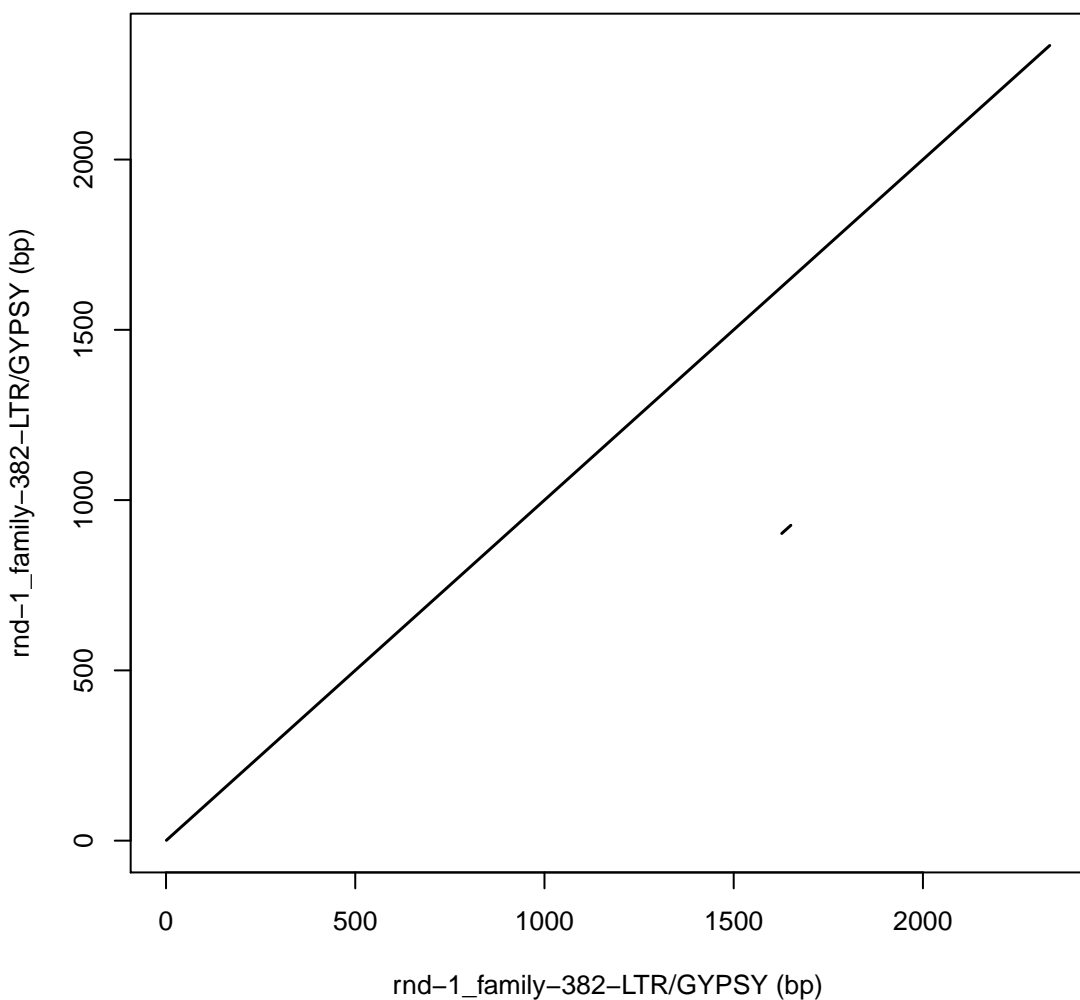
TE: rnd-1_family-382-LTR/GYPSY
consensus size: 2335bp; fragments: 109; full length: 0 (>=2101.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

