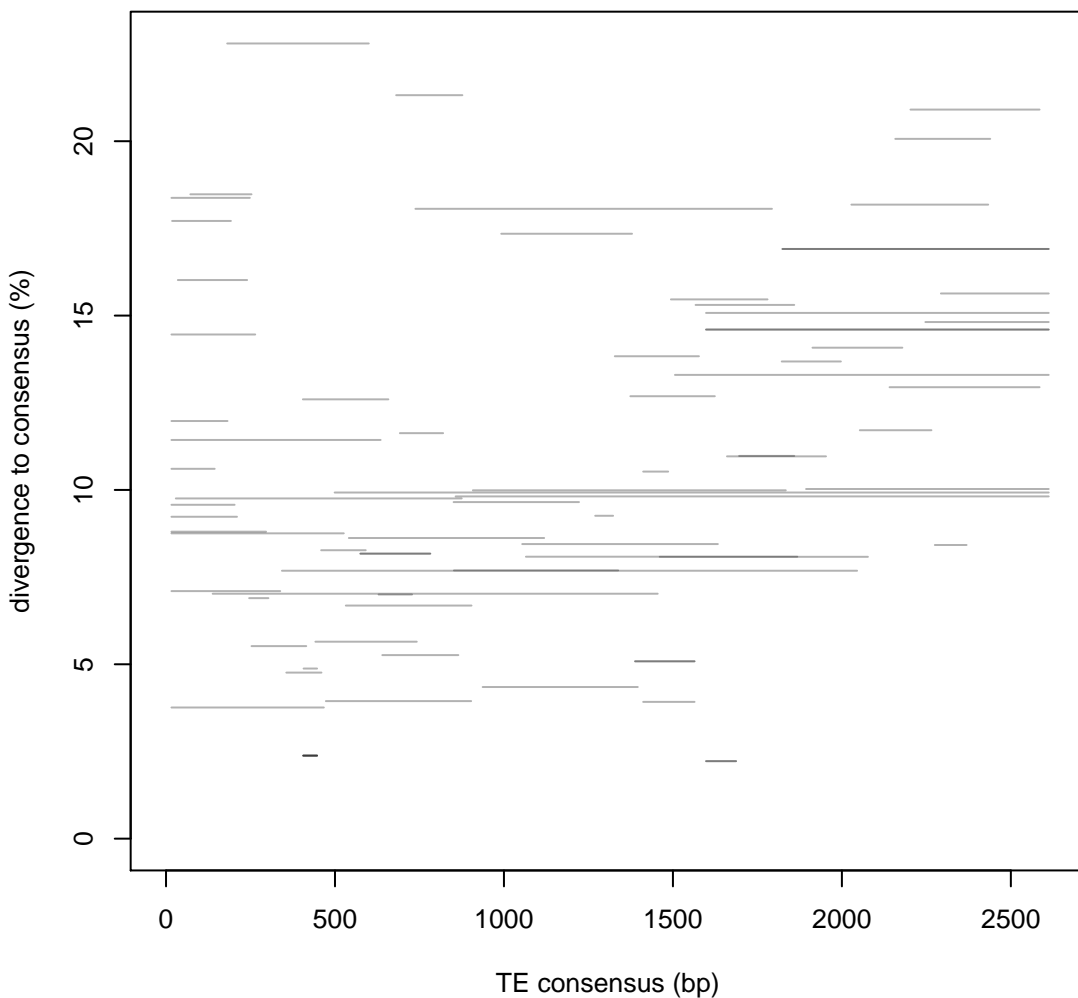
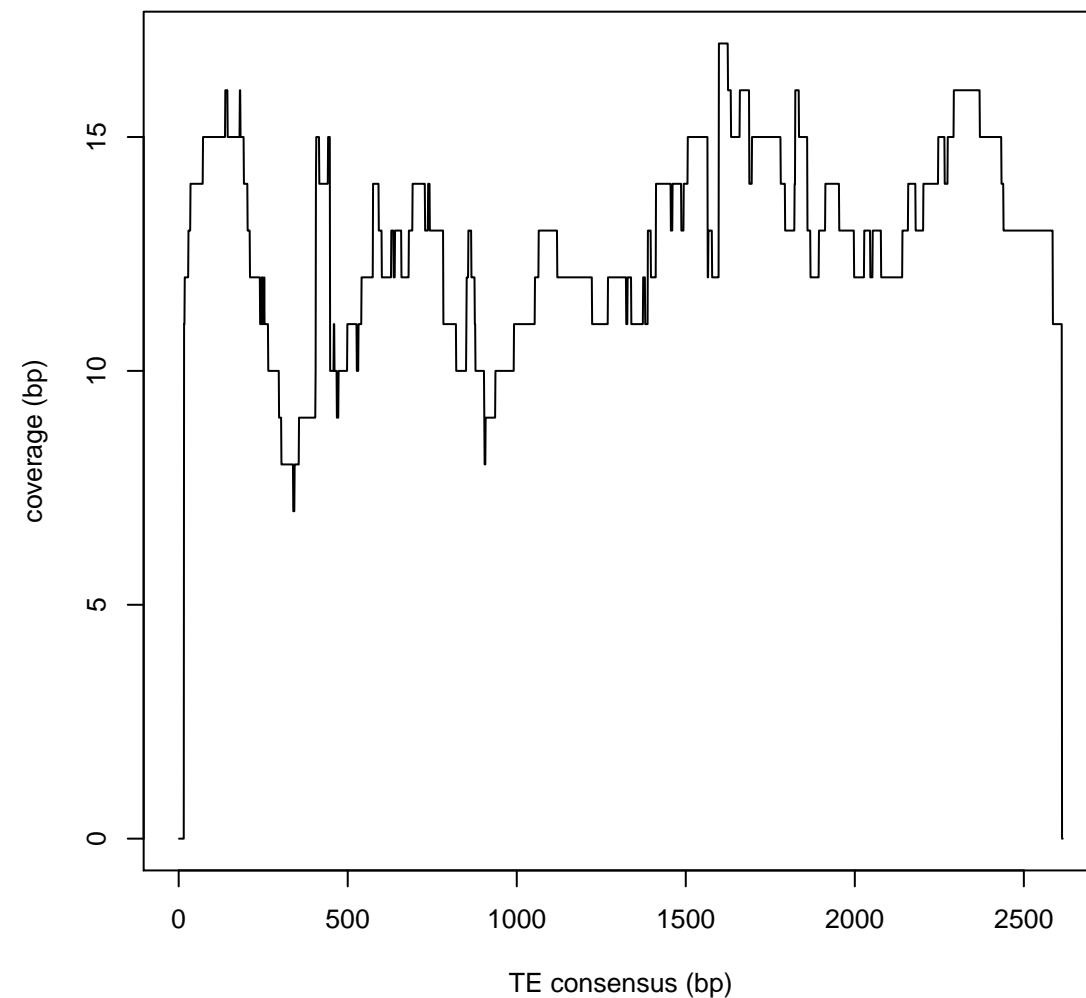


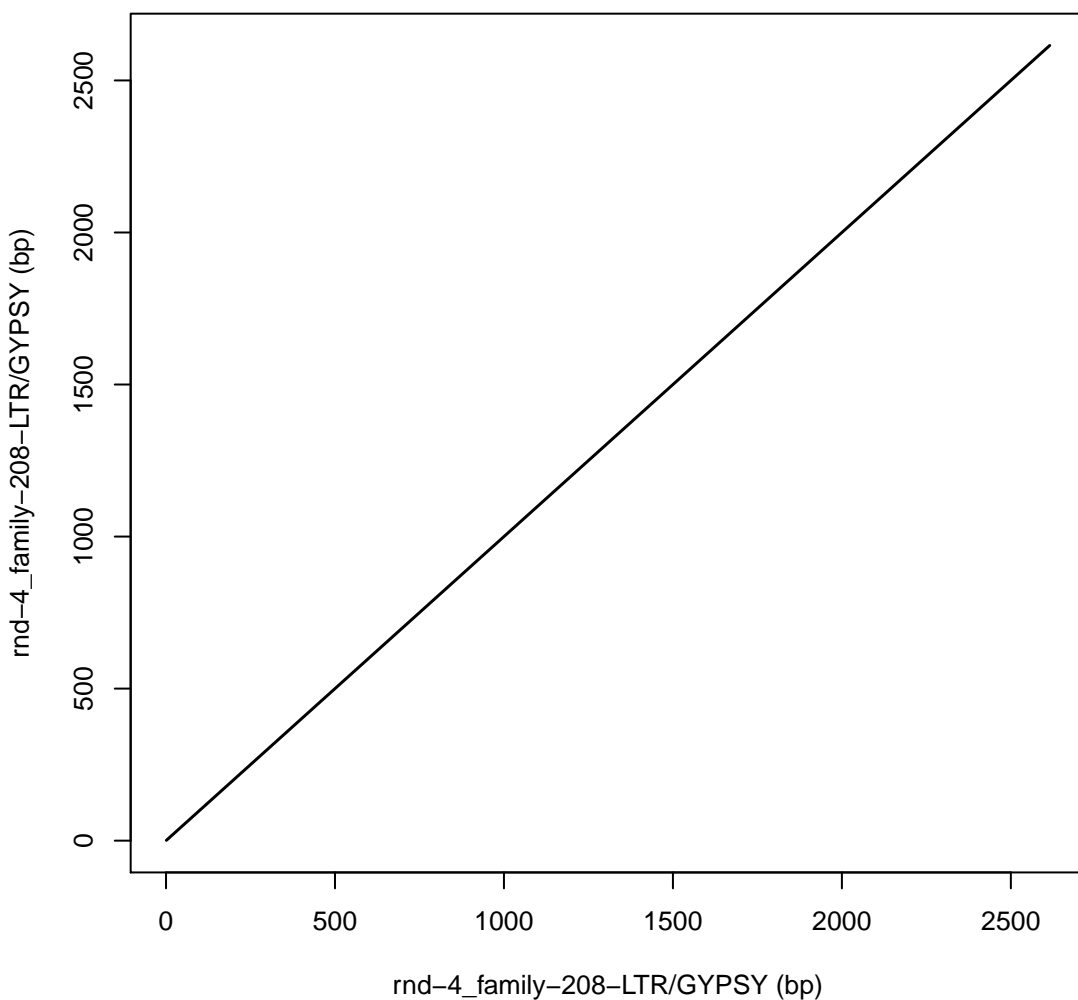
TE: rnd-4_family-208-LTR/GYPSY
consensus size: 2615bp; fragments: 79; full length: 0 (>=2353.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

