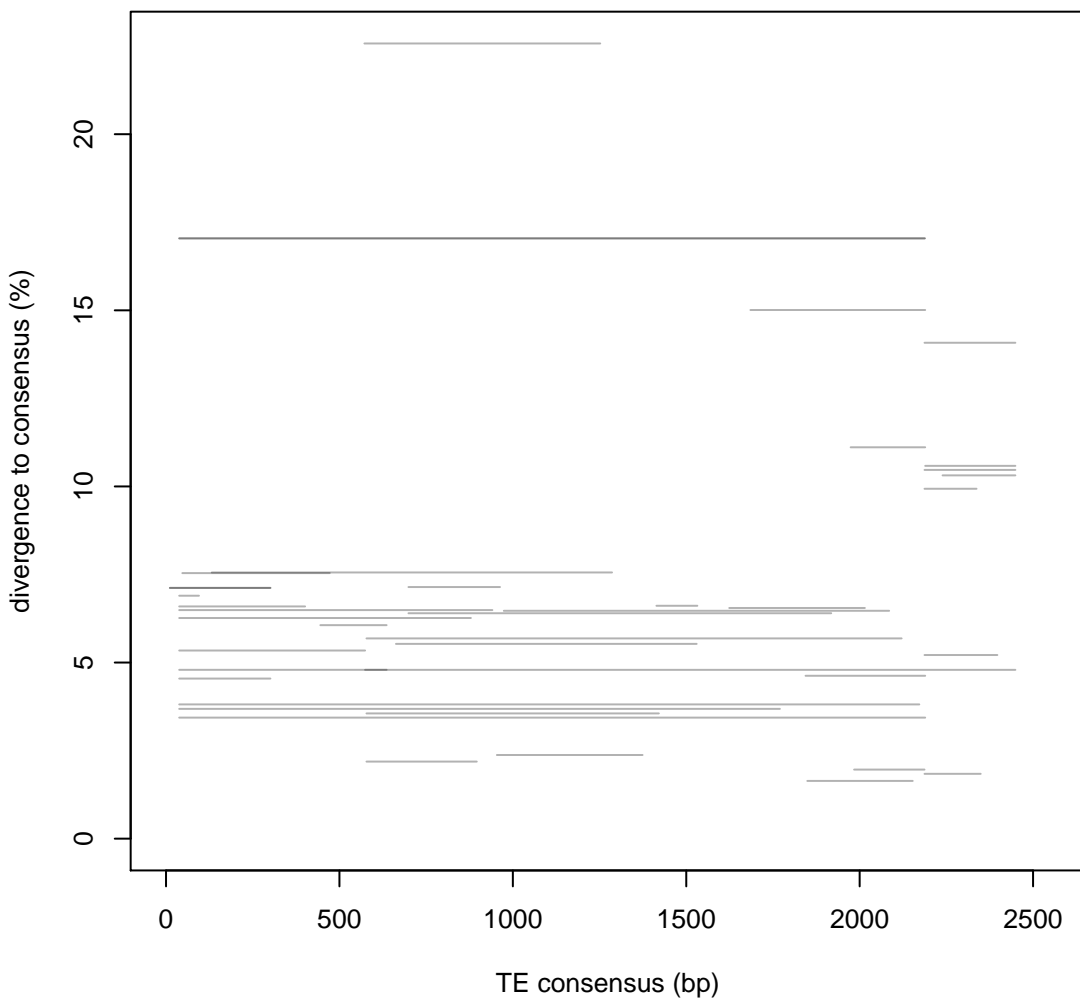
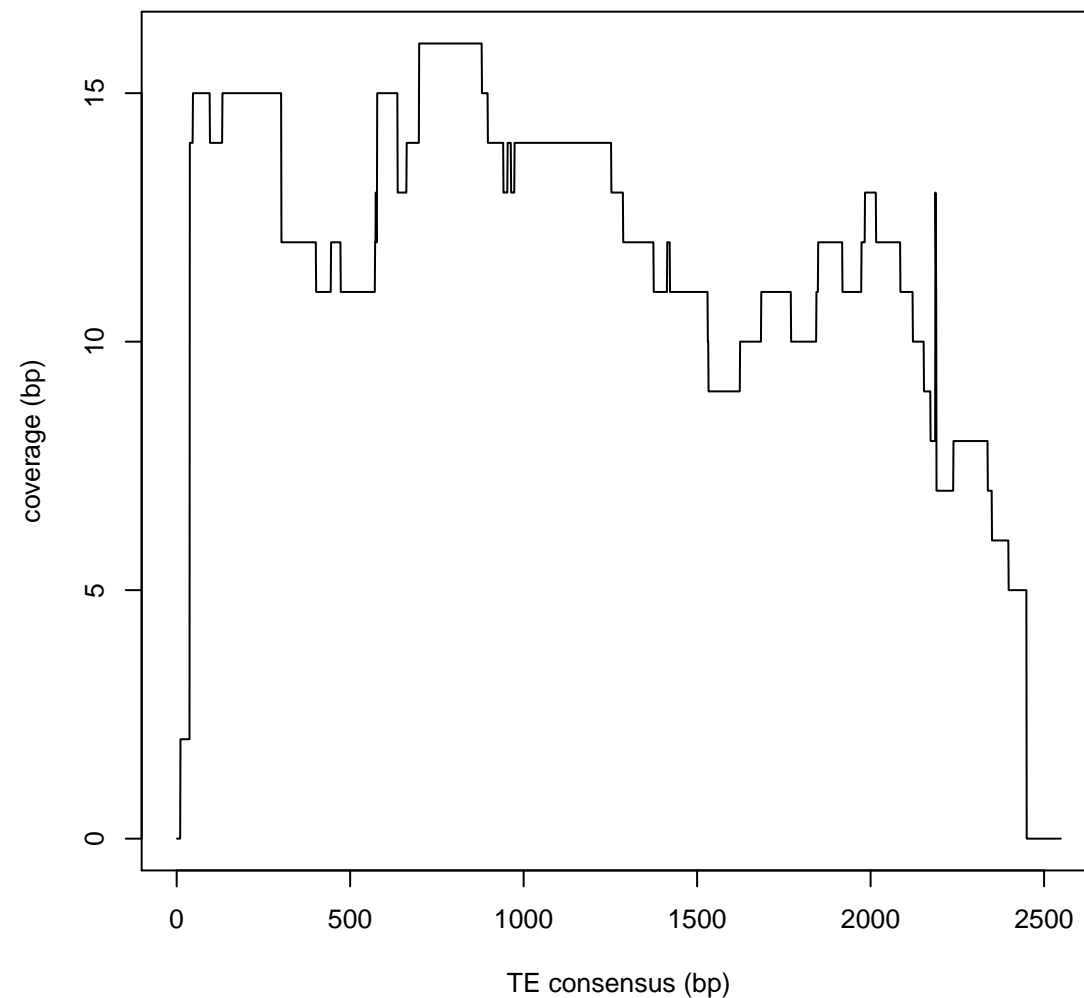


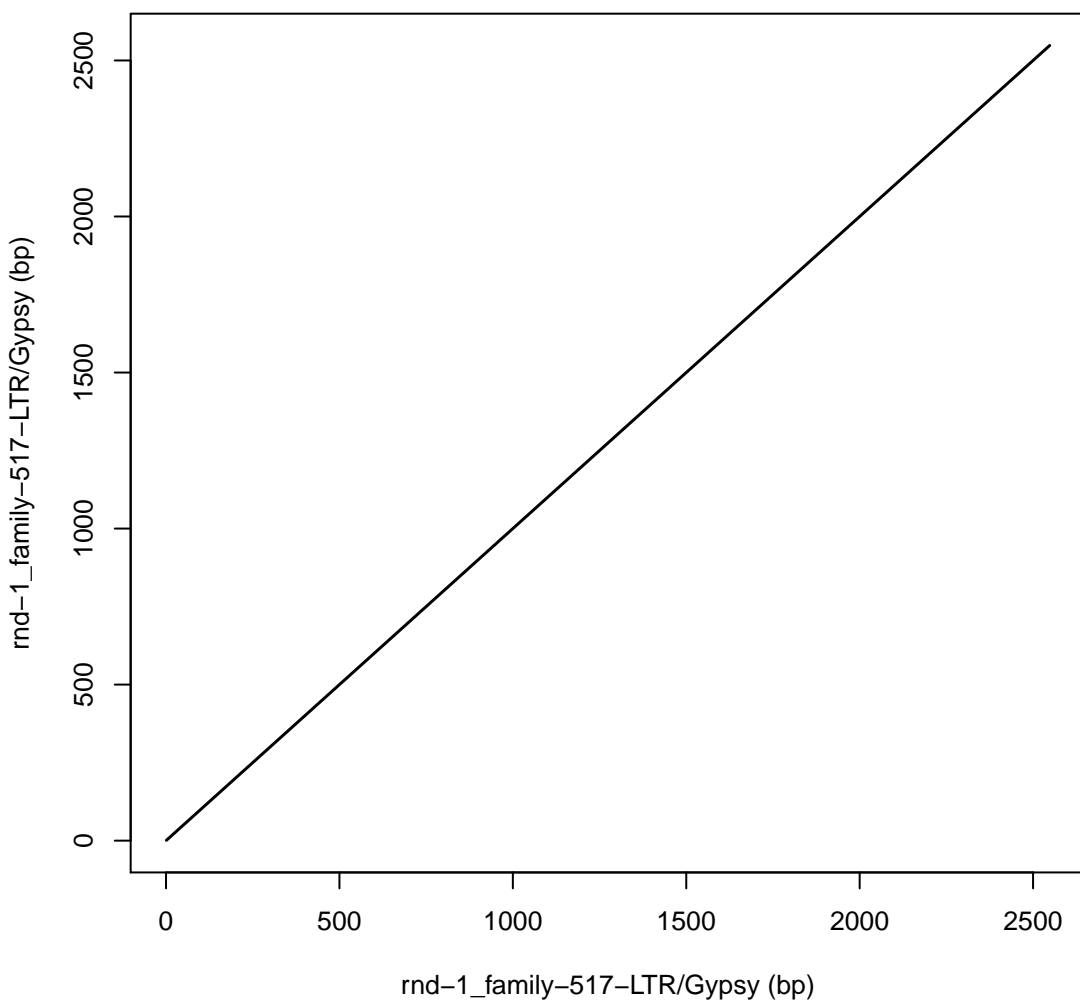
TE: rnd-1_family-517-LTR/Gypsy
consensus size: 2548bp; fragments: 41; full length: 0 (>=2293.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

