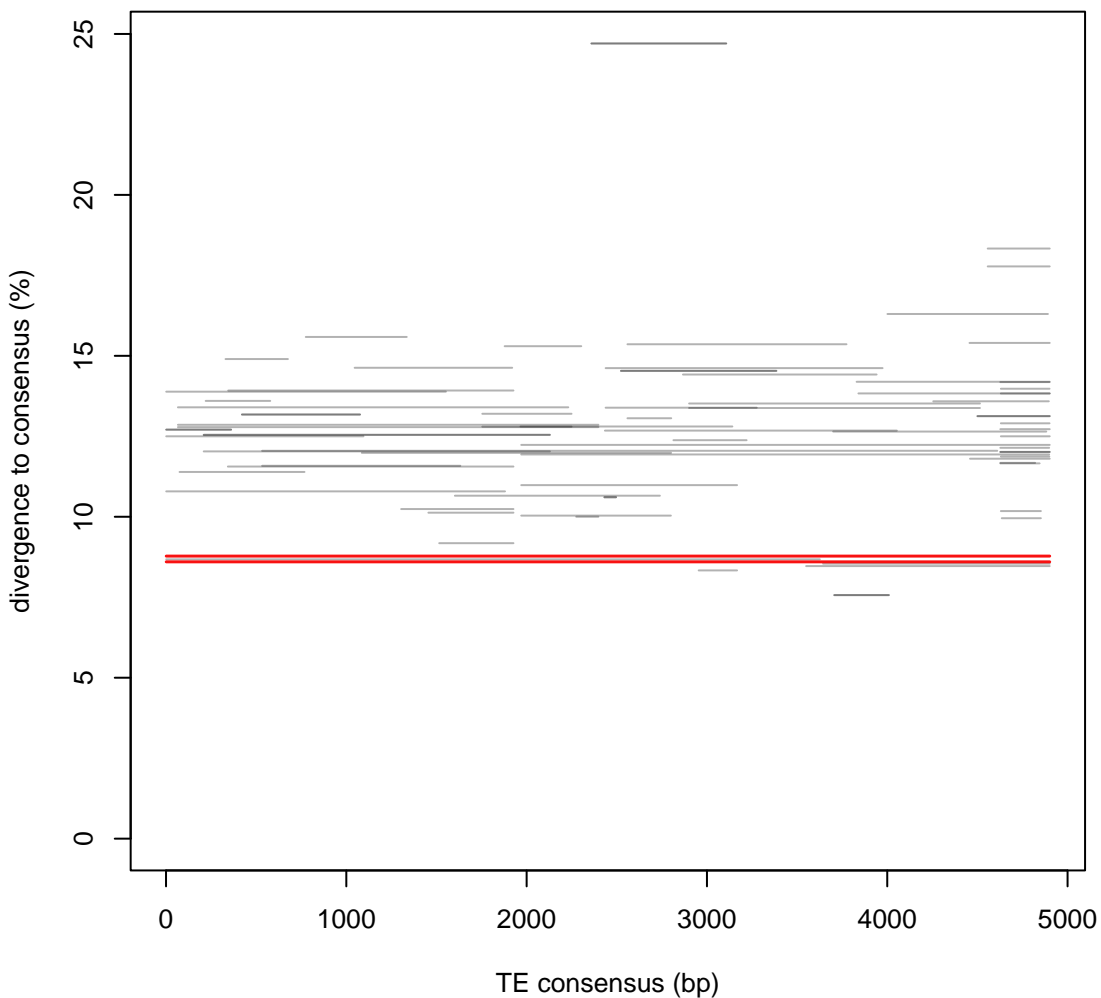
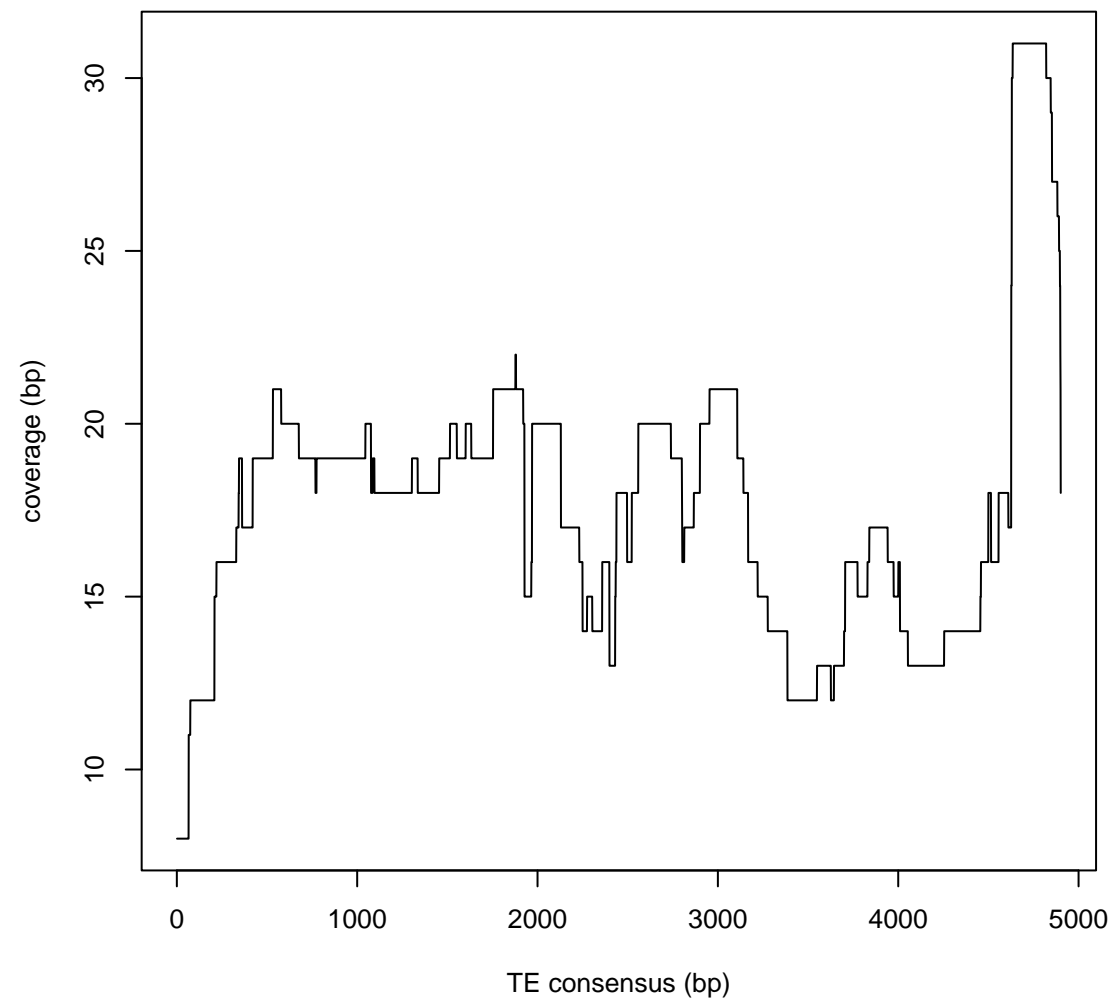


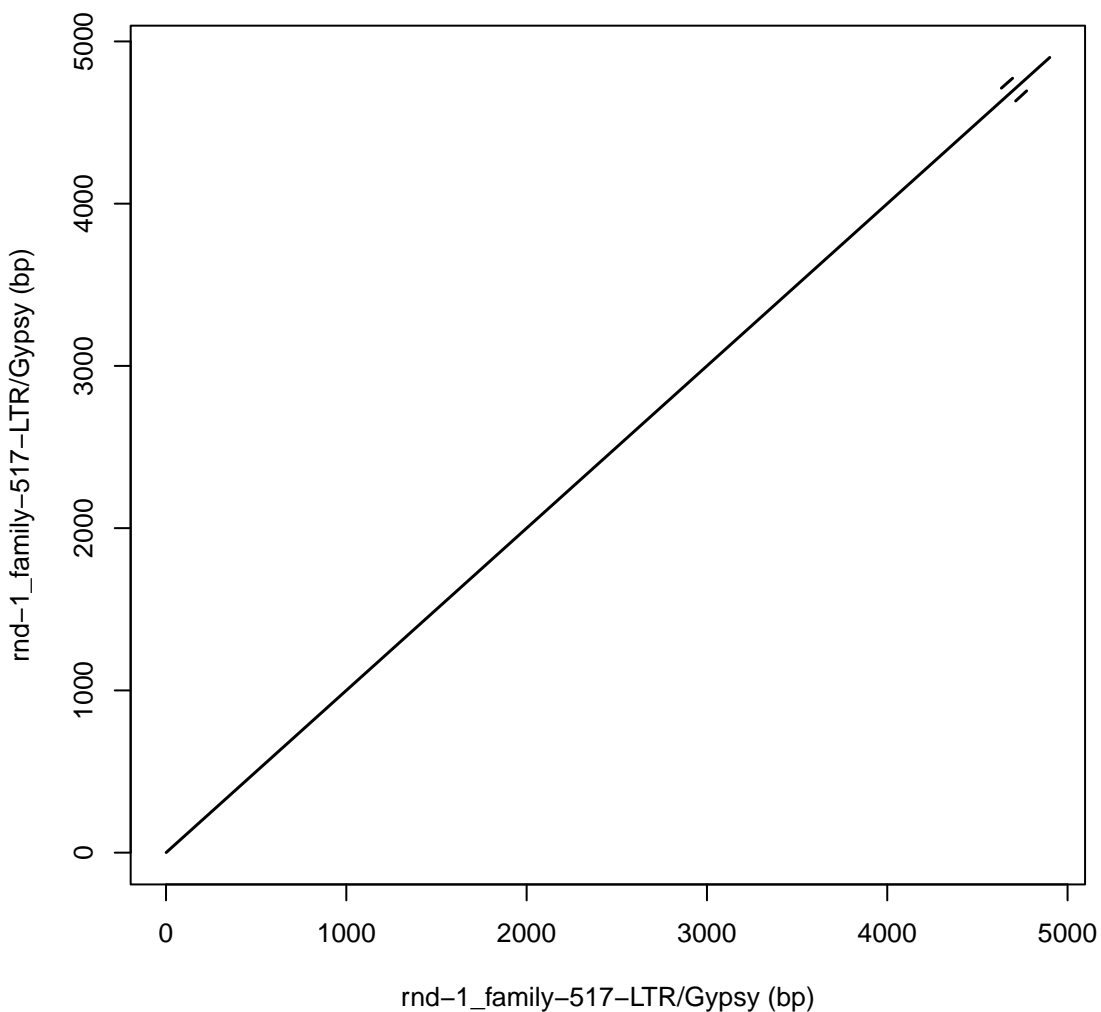
TE: rnd-1_family-517-LTR/Gypsy
consensus size: 4901bp; fragments: 84; full length: 2 (≥ 4410.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

