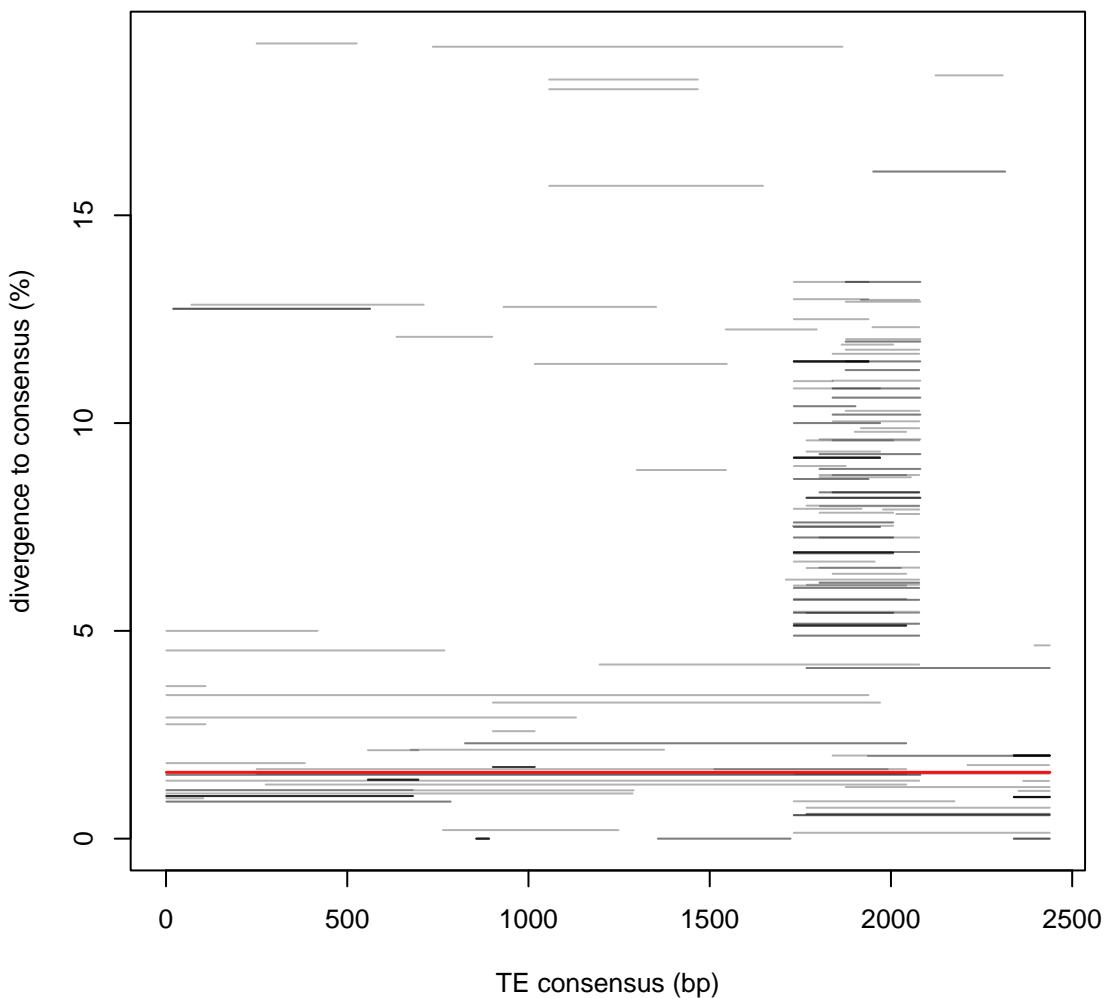
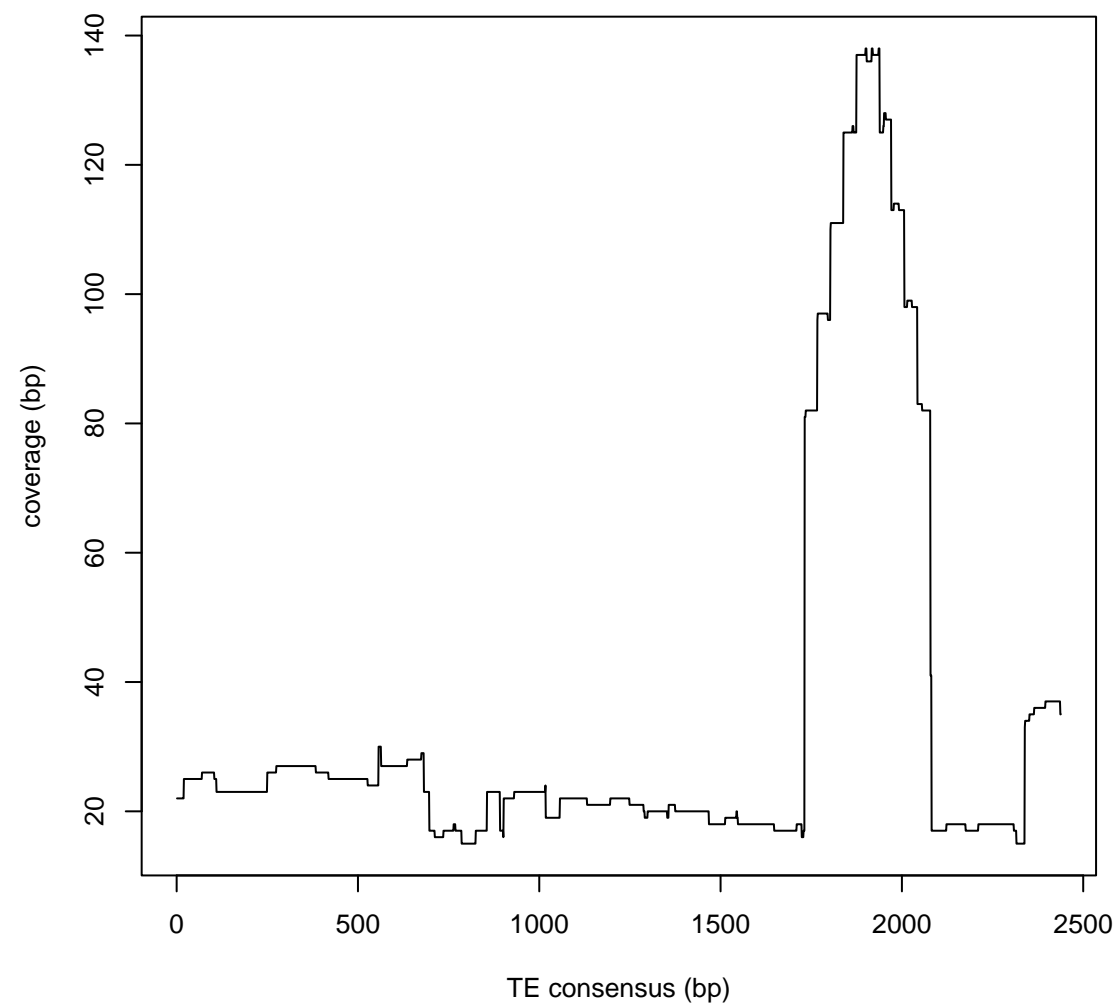


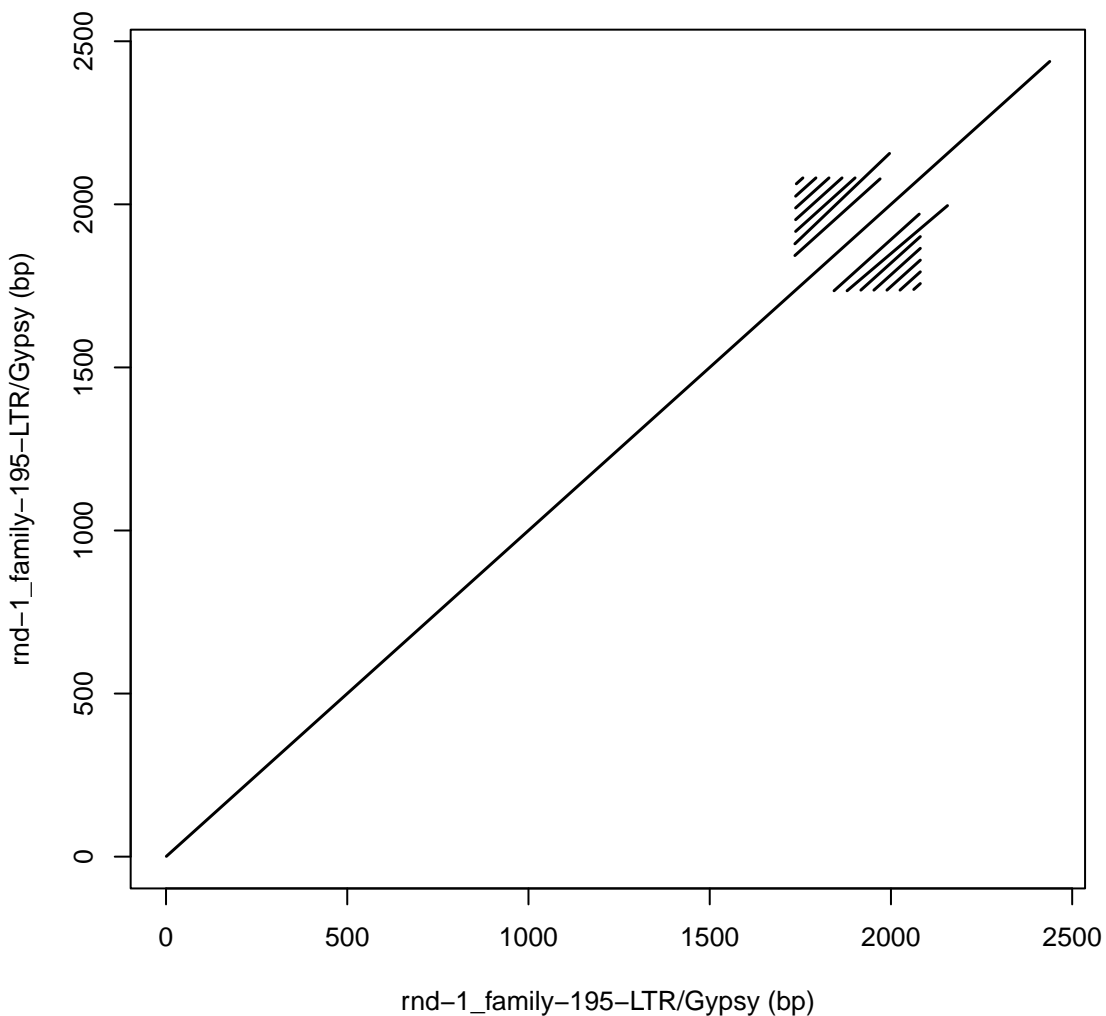
TE: rnd-1_family-195-LTR/Gypsy
consensus size: 2438bp; fragments: 224; full length: 1 (>=2194.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

