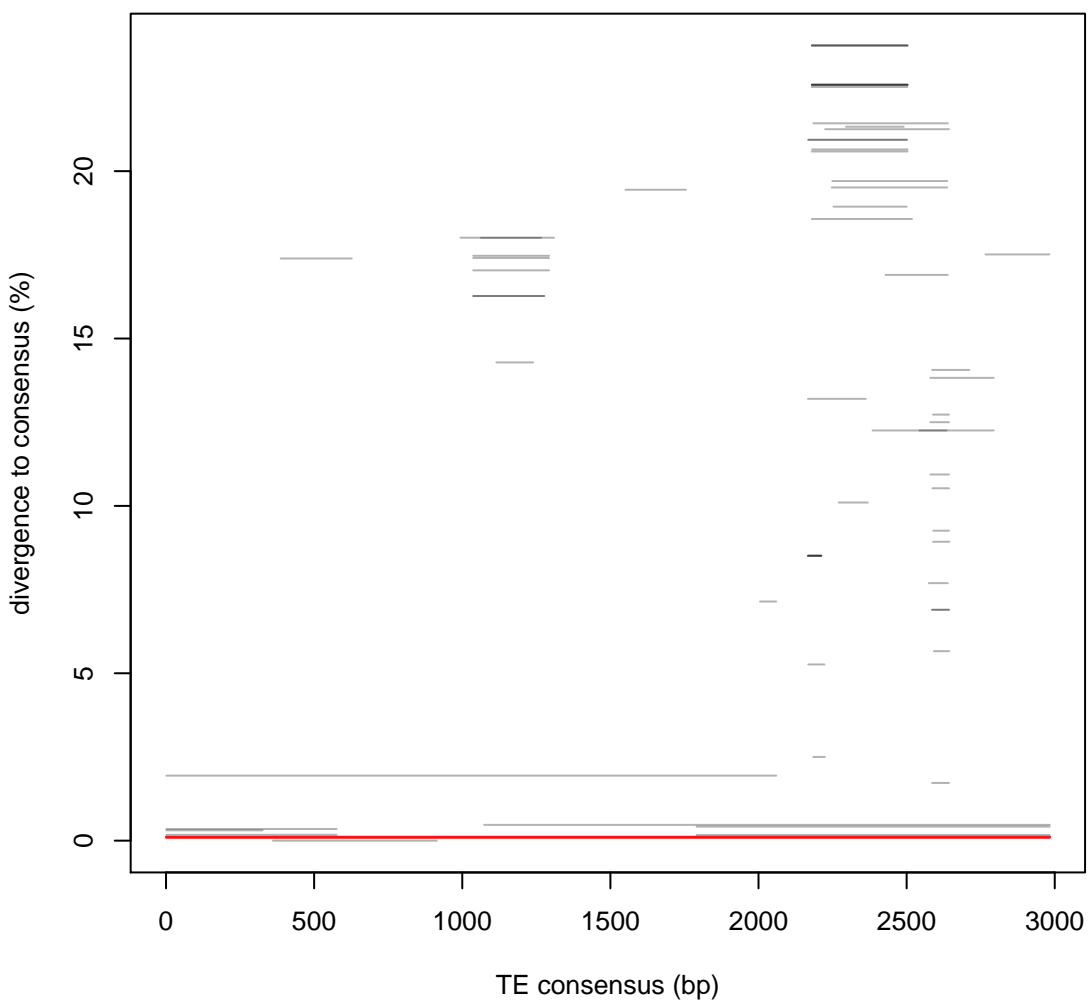
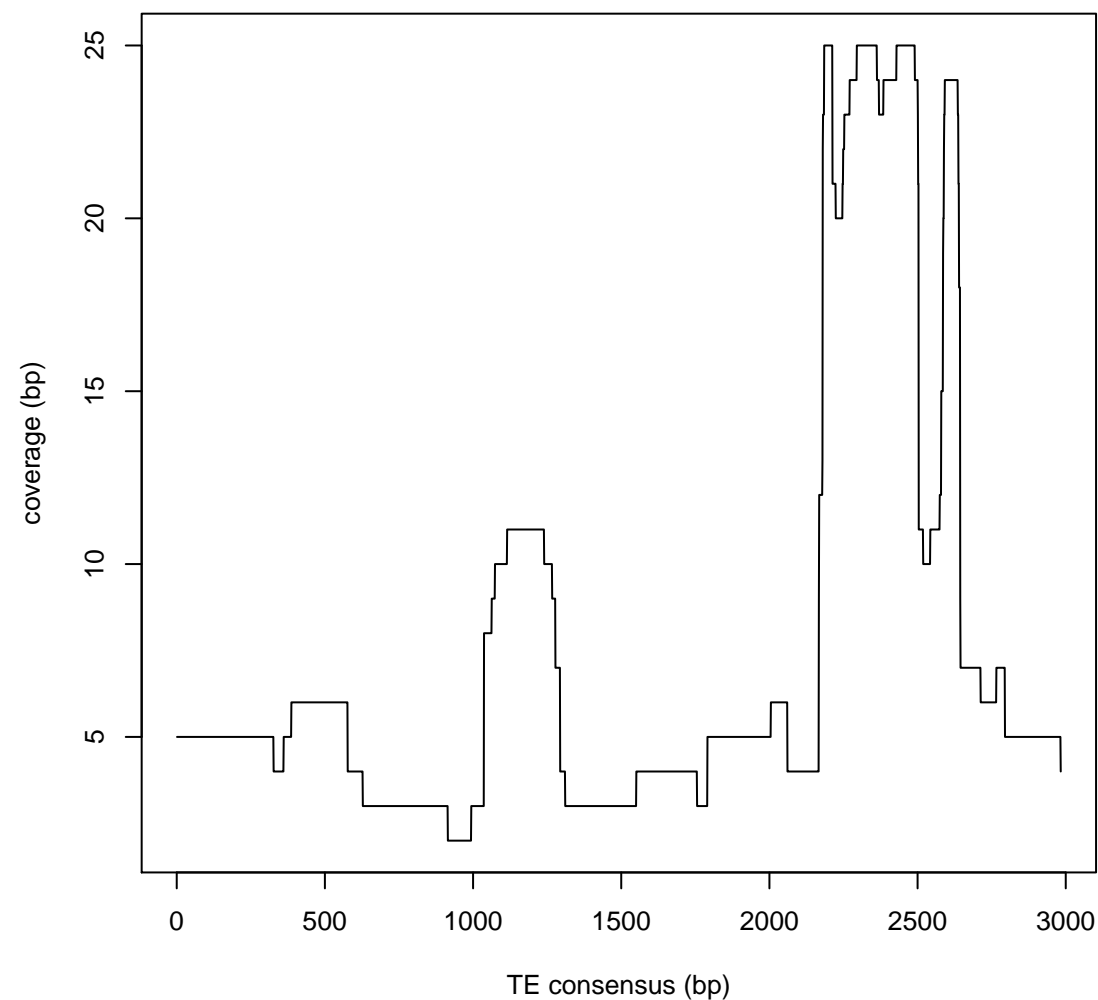


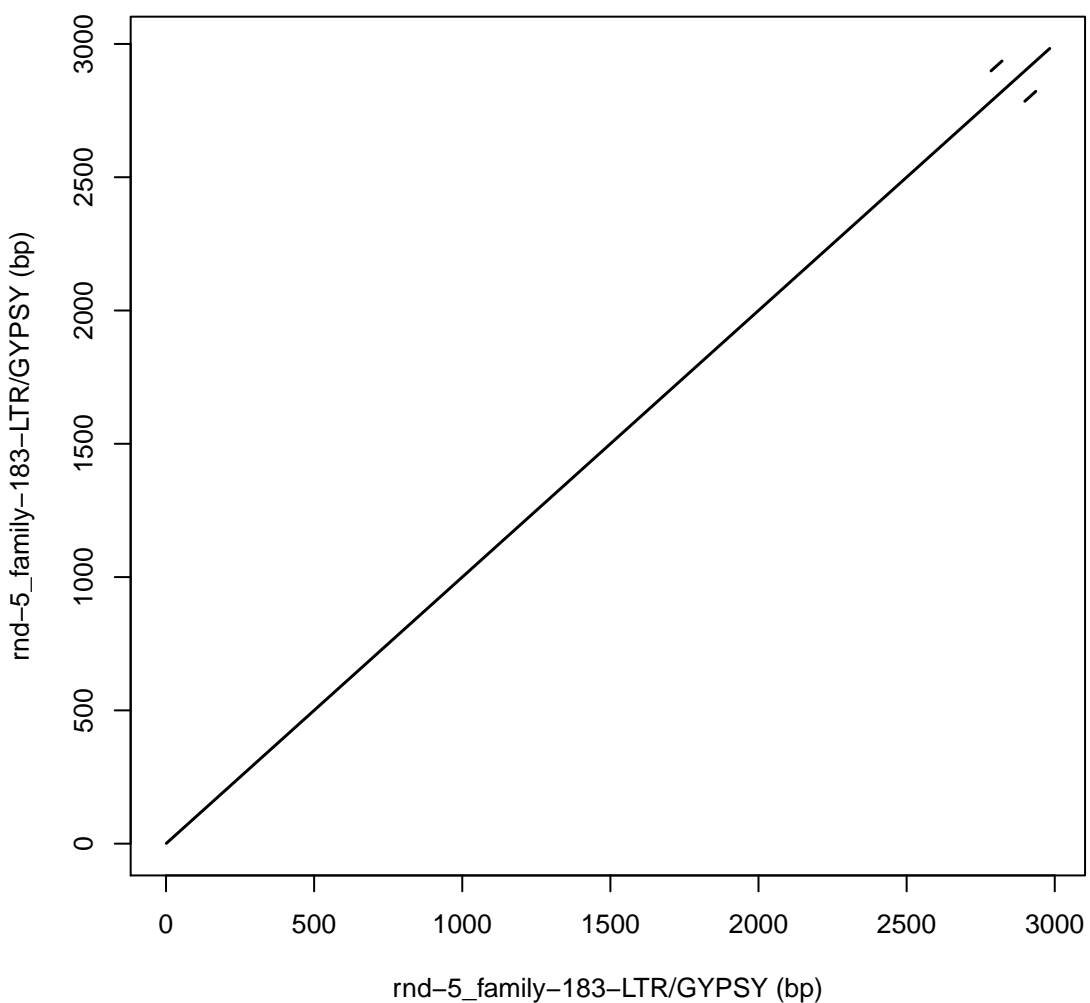
TE: rnd-5_family-183-LTR/GYPSY
consensus size: 2983bp; fragments: 64; full length: 1 (>=2684.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

