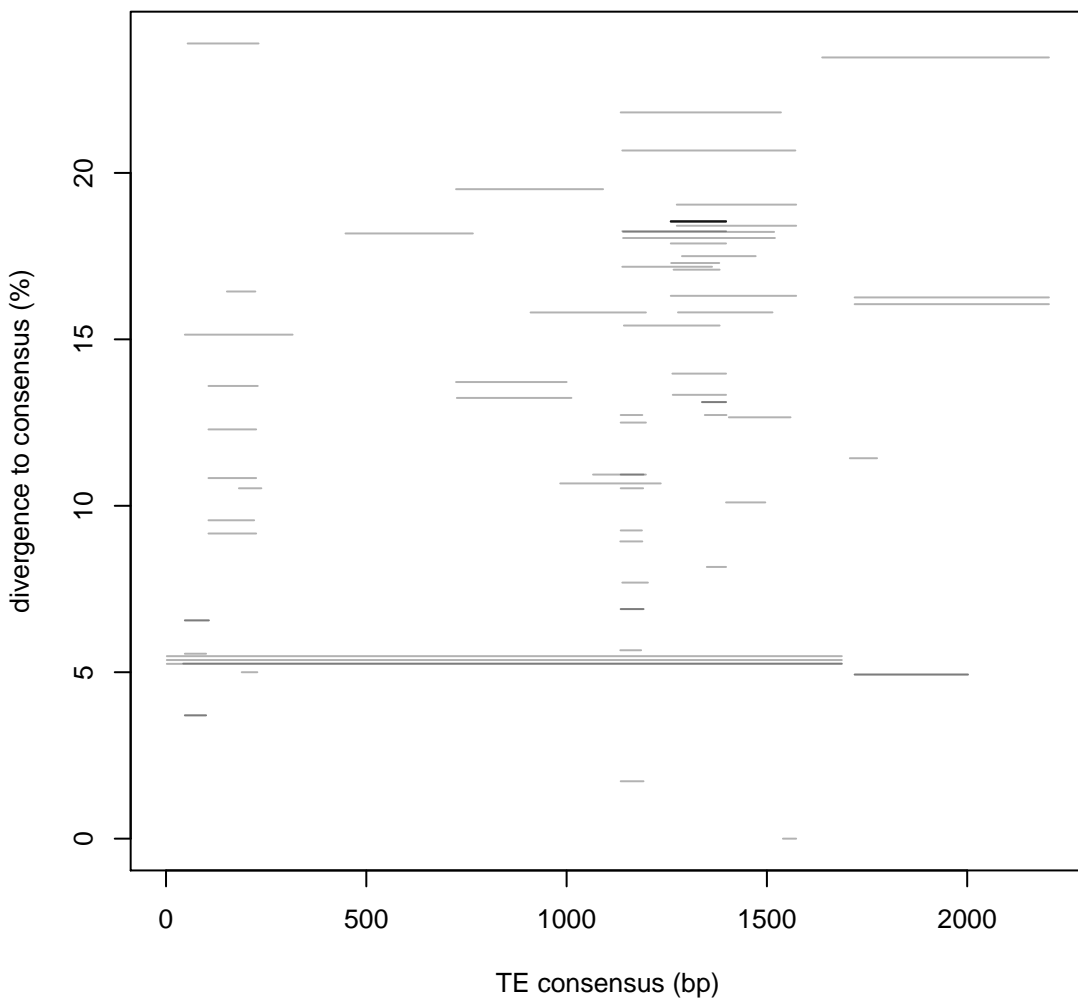
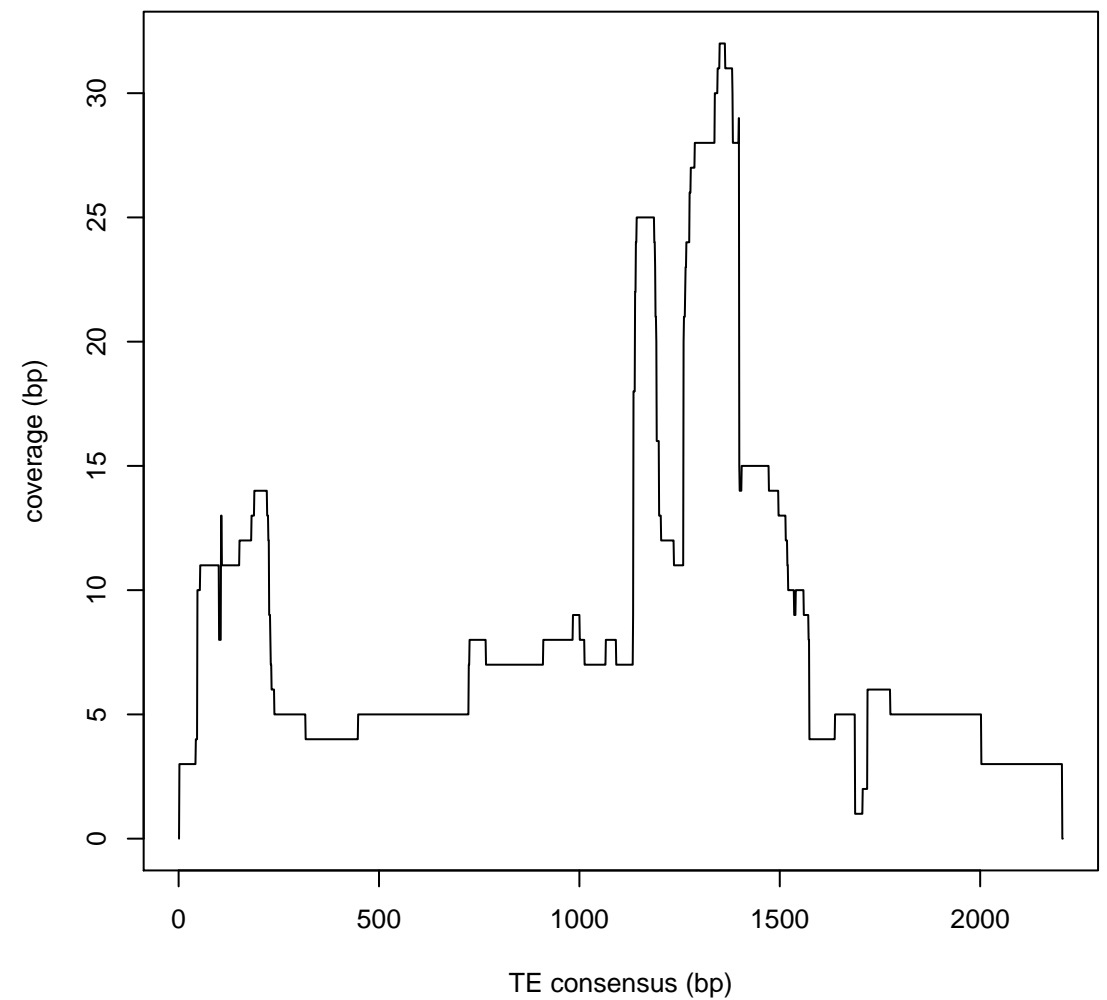


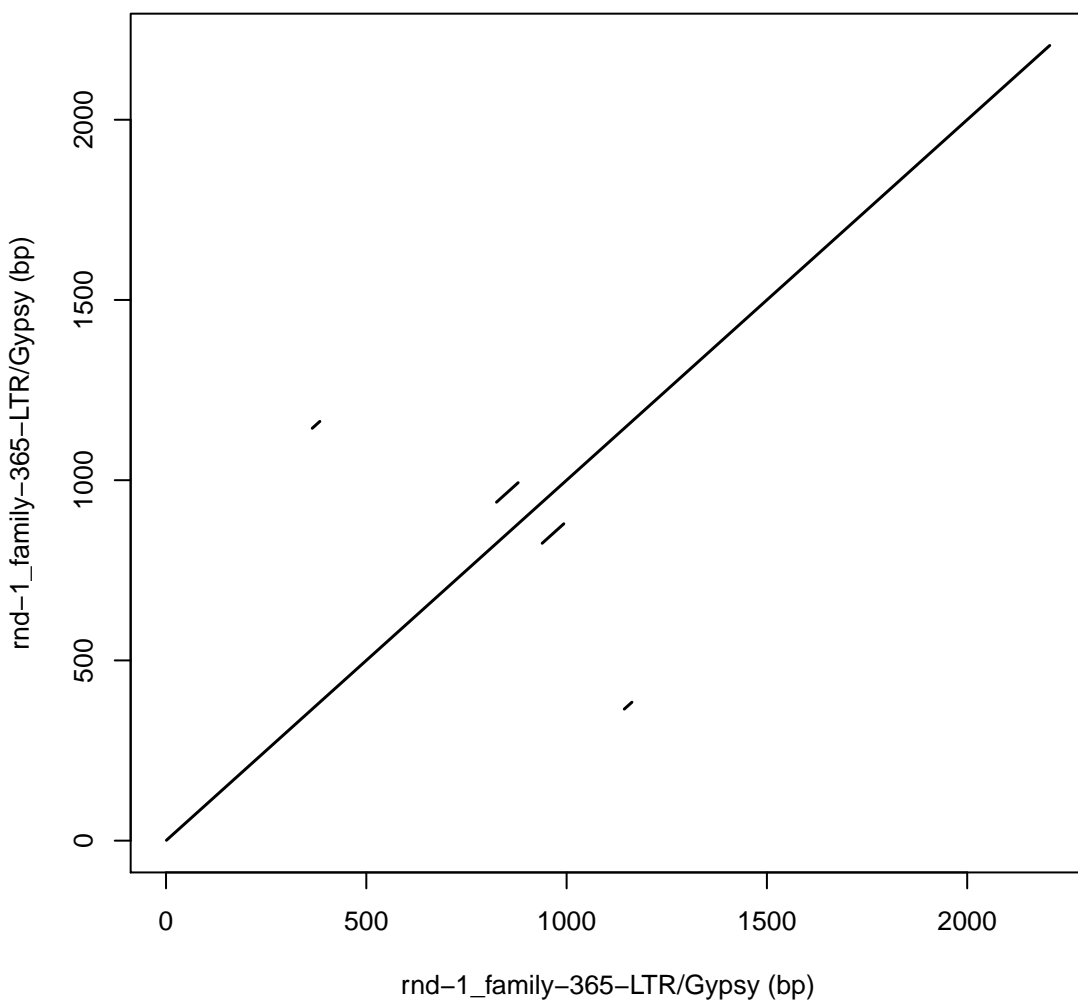
TE: rnd-1_family-365-LTR/Gypsy
consensus size: 2206bp; fragments: 74; full length: 0 (≥ 1985.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

