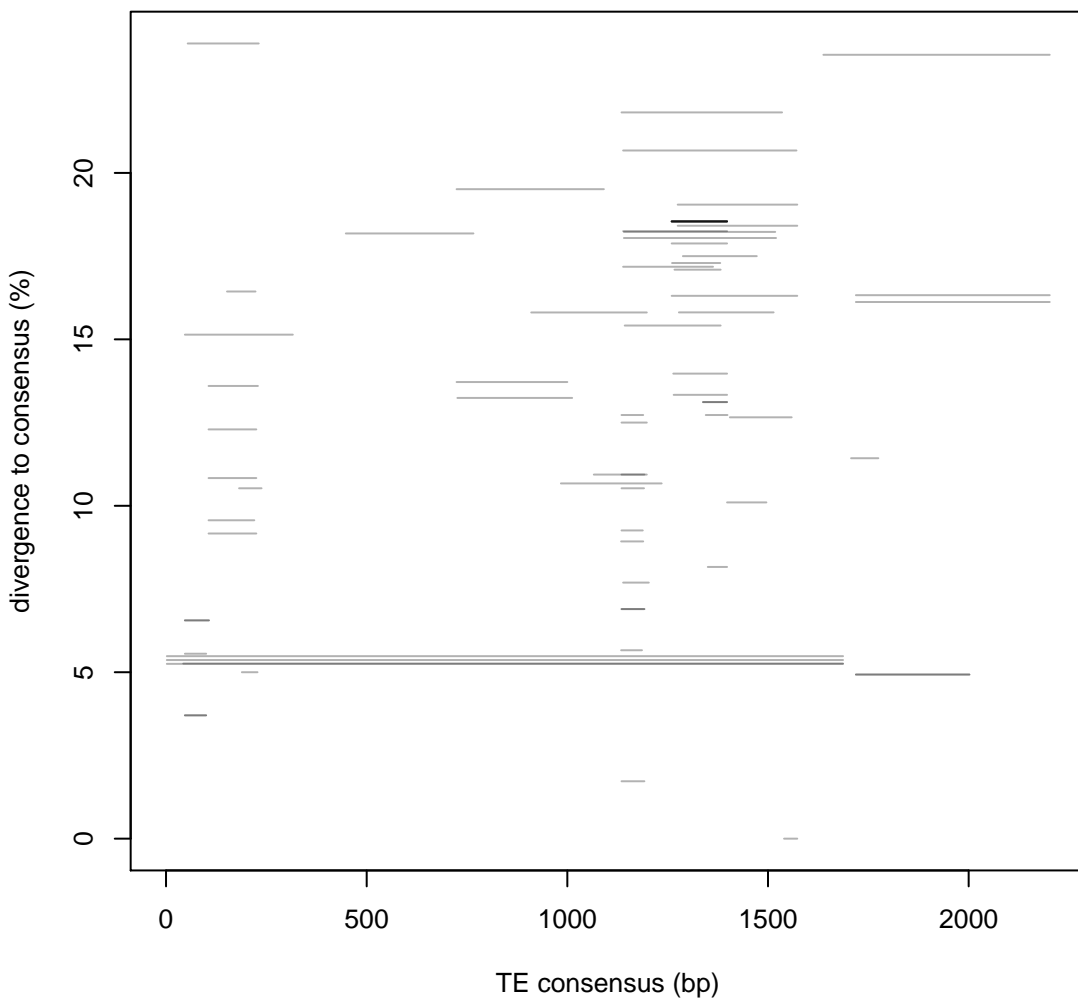
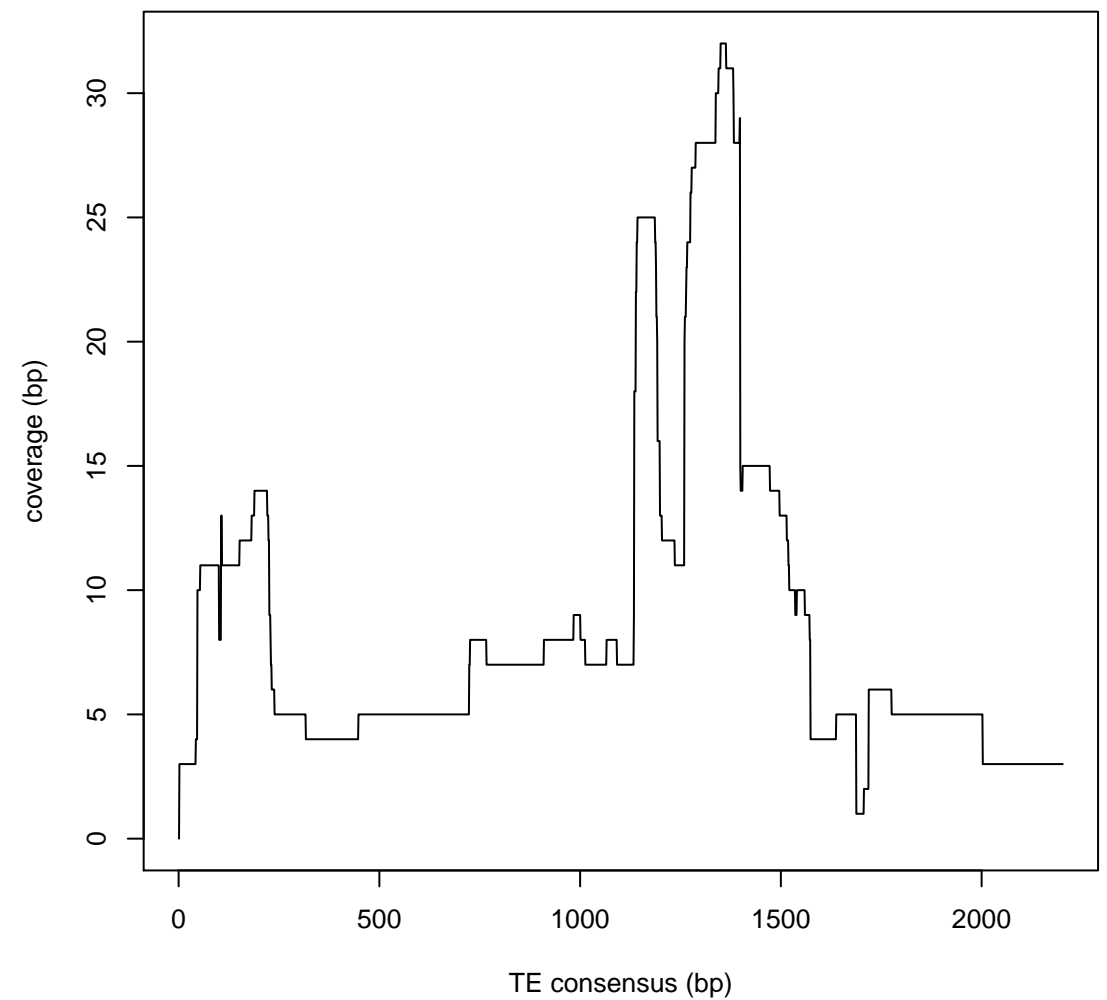


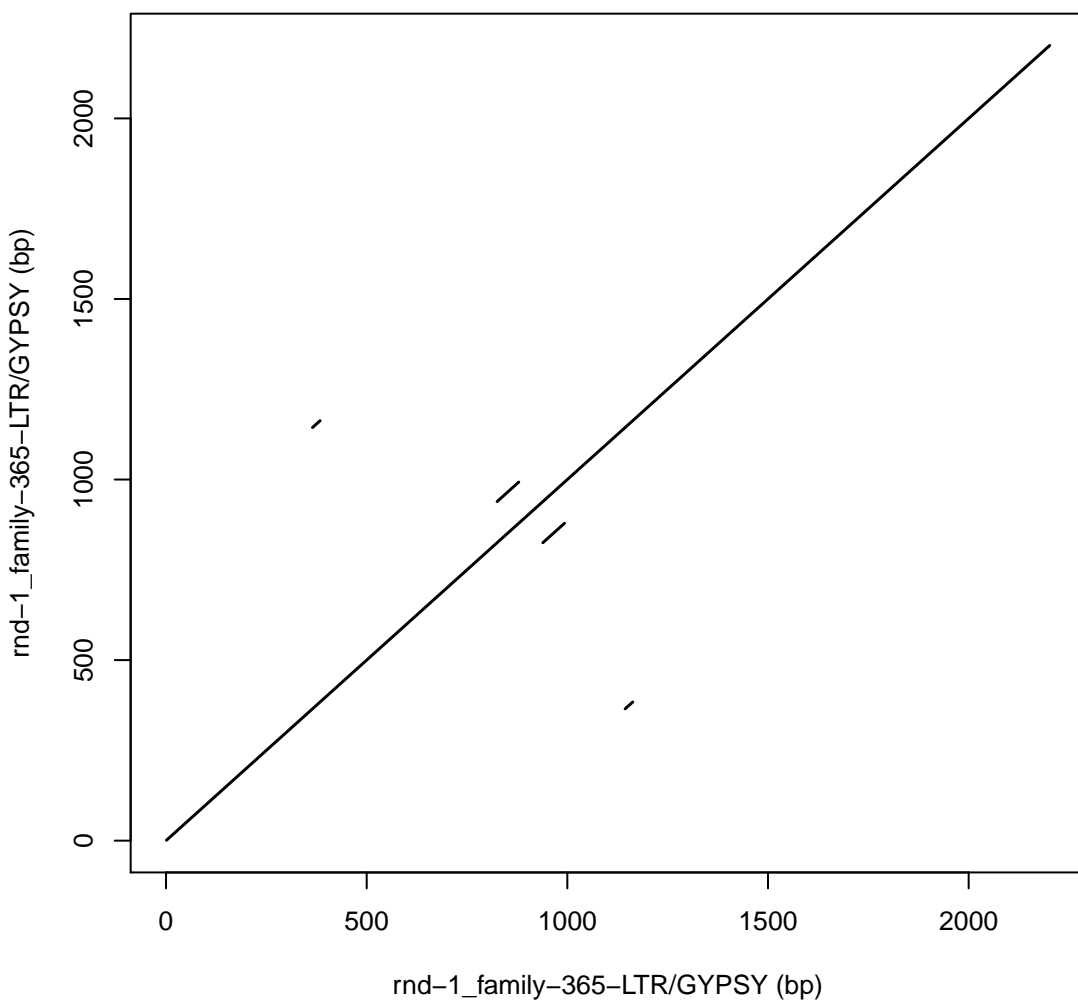
TE: rnd-1_family-365-LTR/GYPSY
consensus size: 2202bp; fragments: 74; full length: 0 (≥ 1981.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

