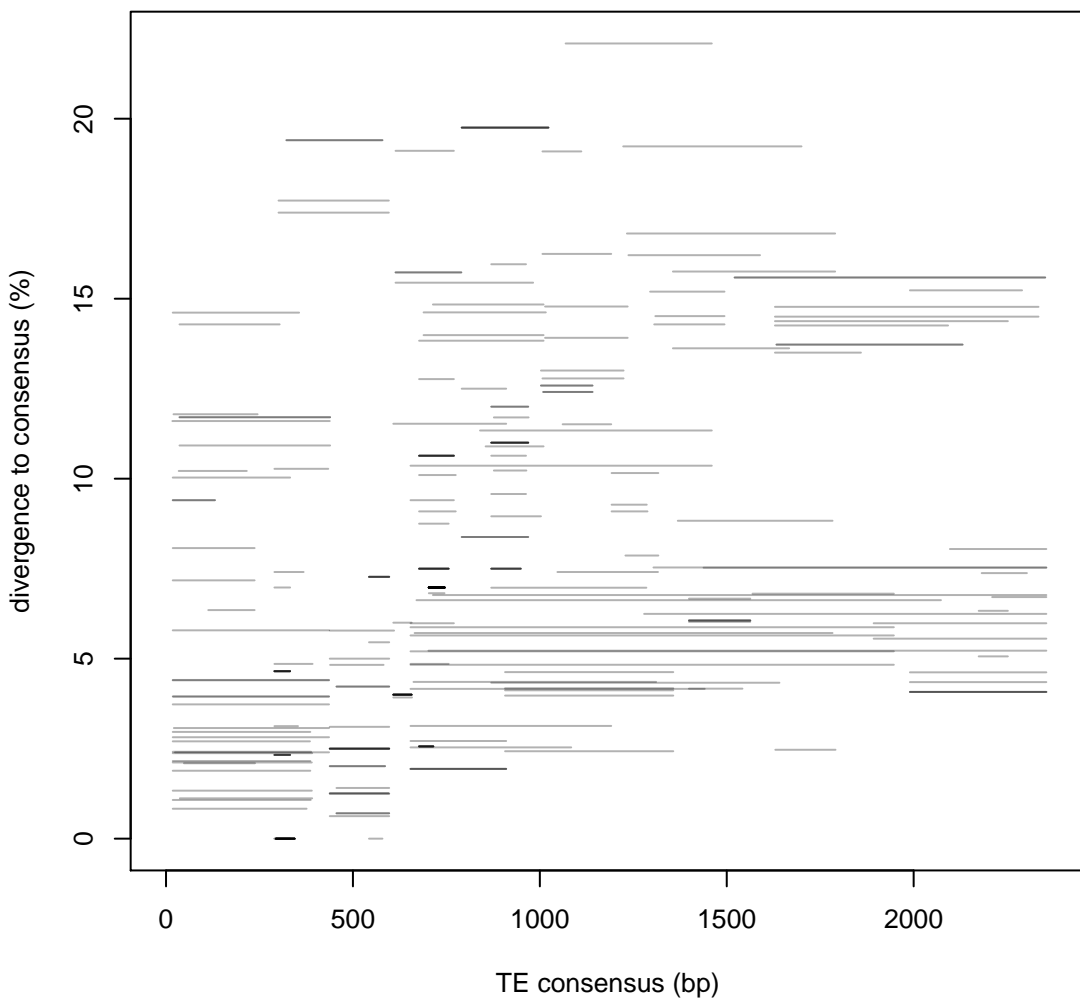
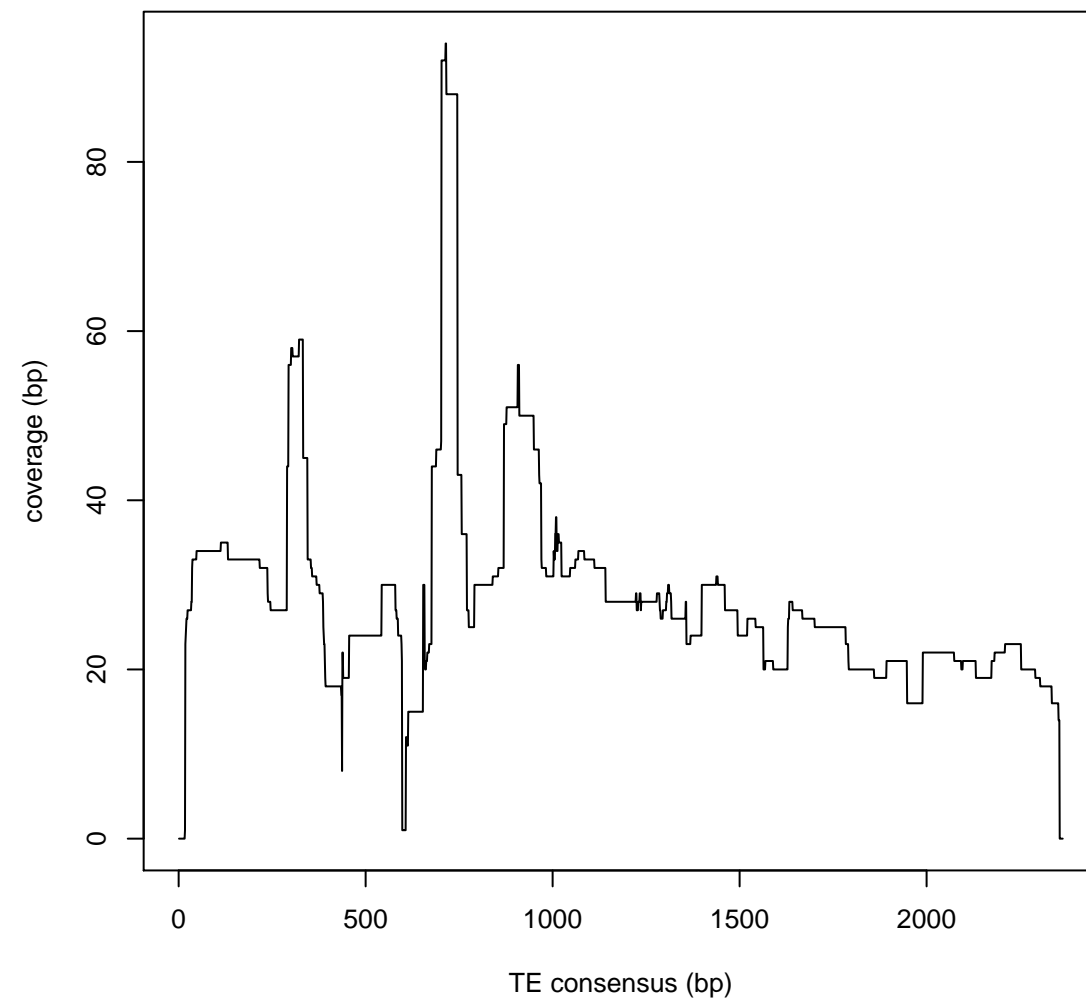


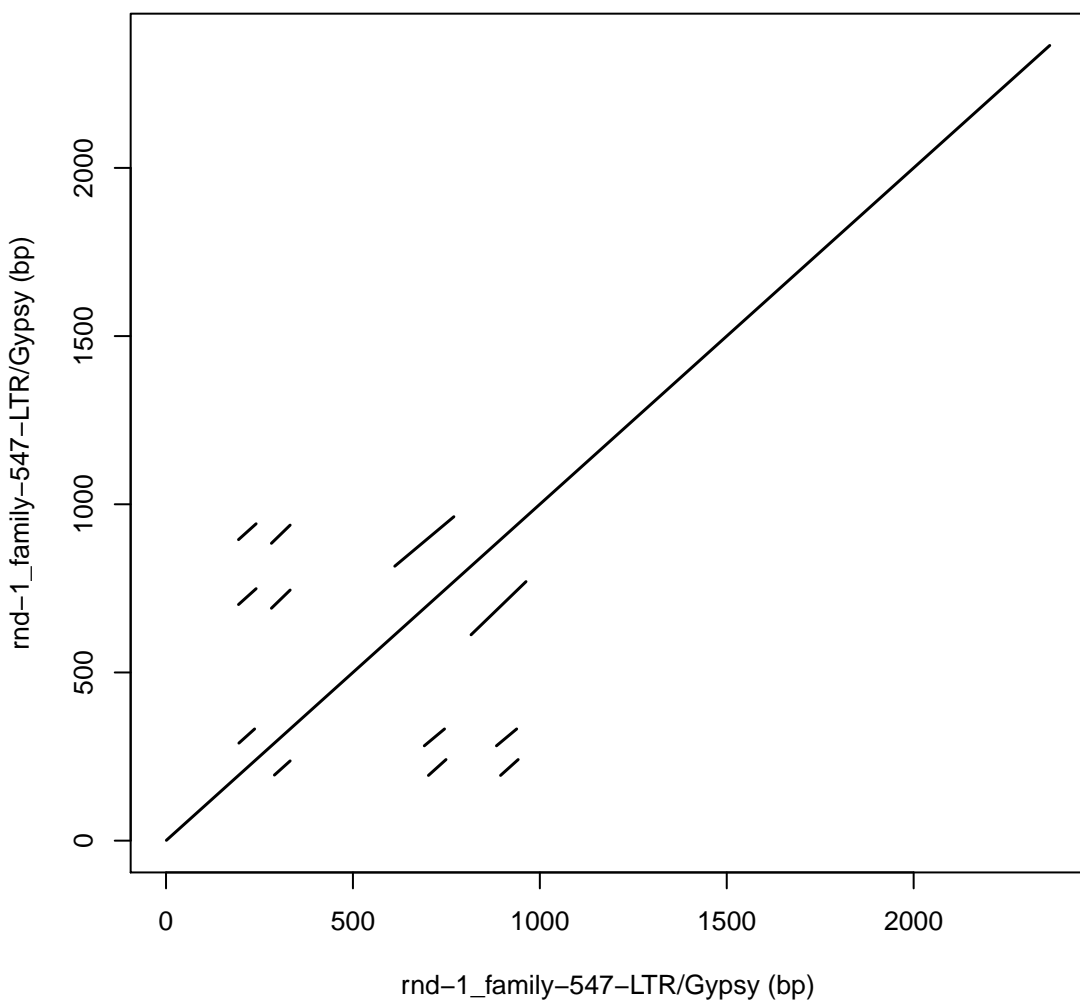
TE: rnd-1_family-547-LTR/Gypsy
consensus size: 2364bp; fragments: 290; full length: 0 (≥ 2127.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

