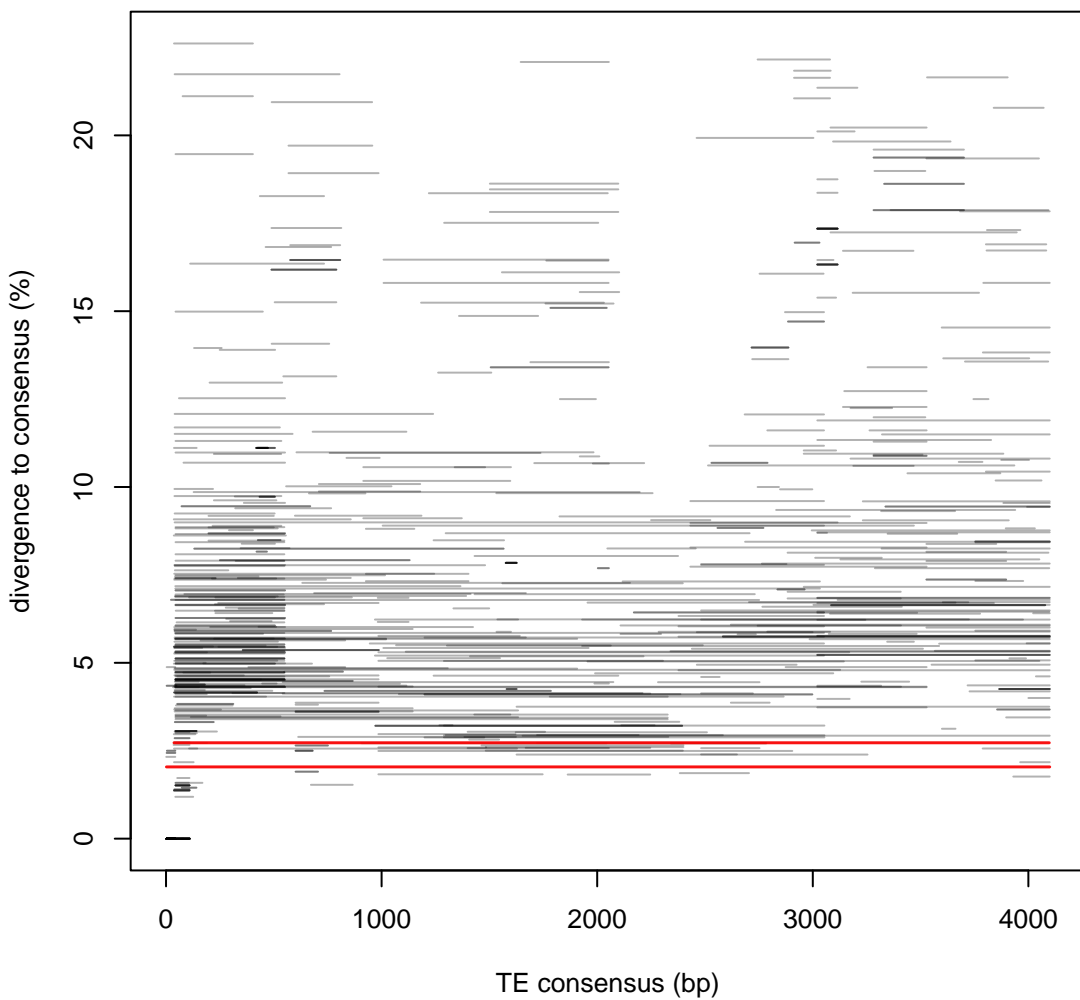
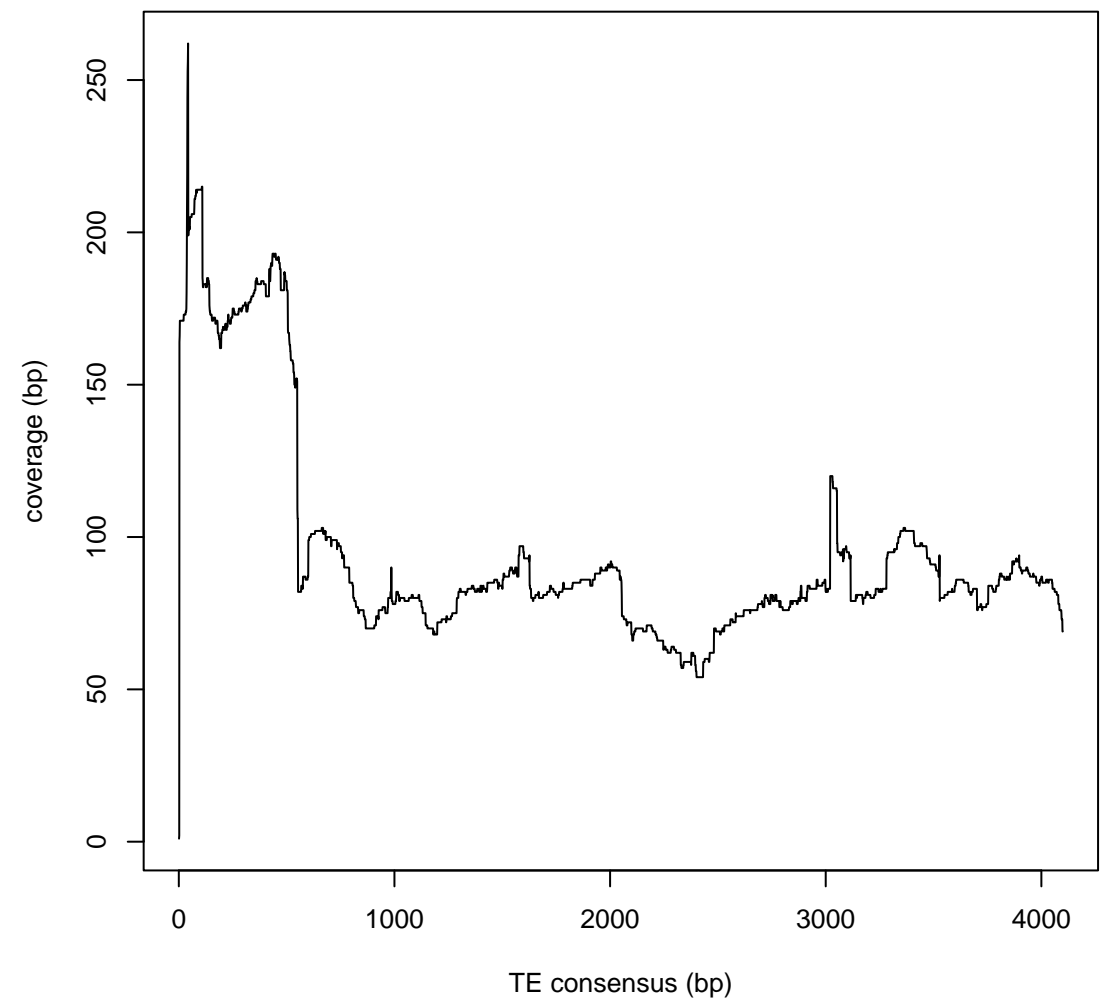


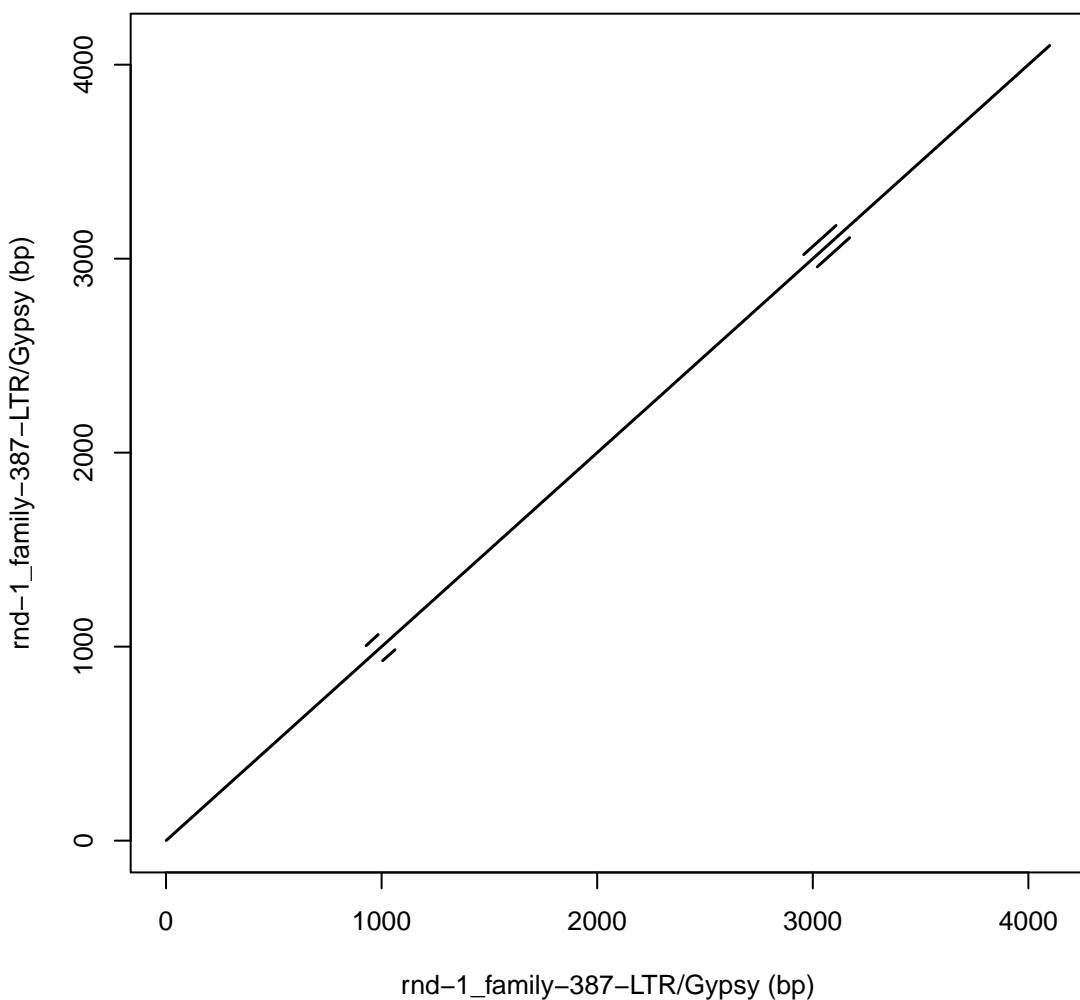
TE: rnd-1_family-387-LTR/Gypsy
consensus size: 4099bp; fragments: 831; full length: 2 (≥ 3689.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

