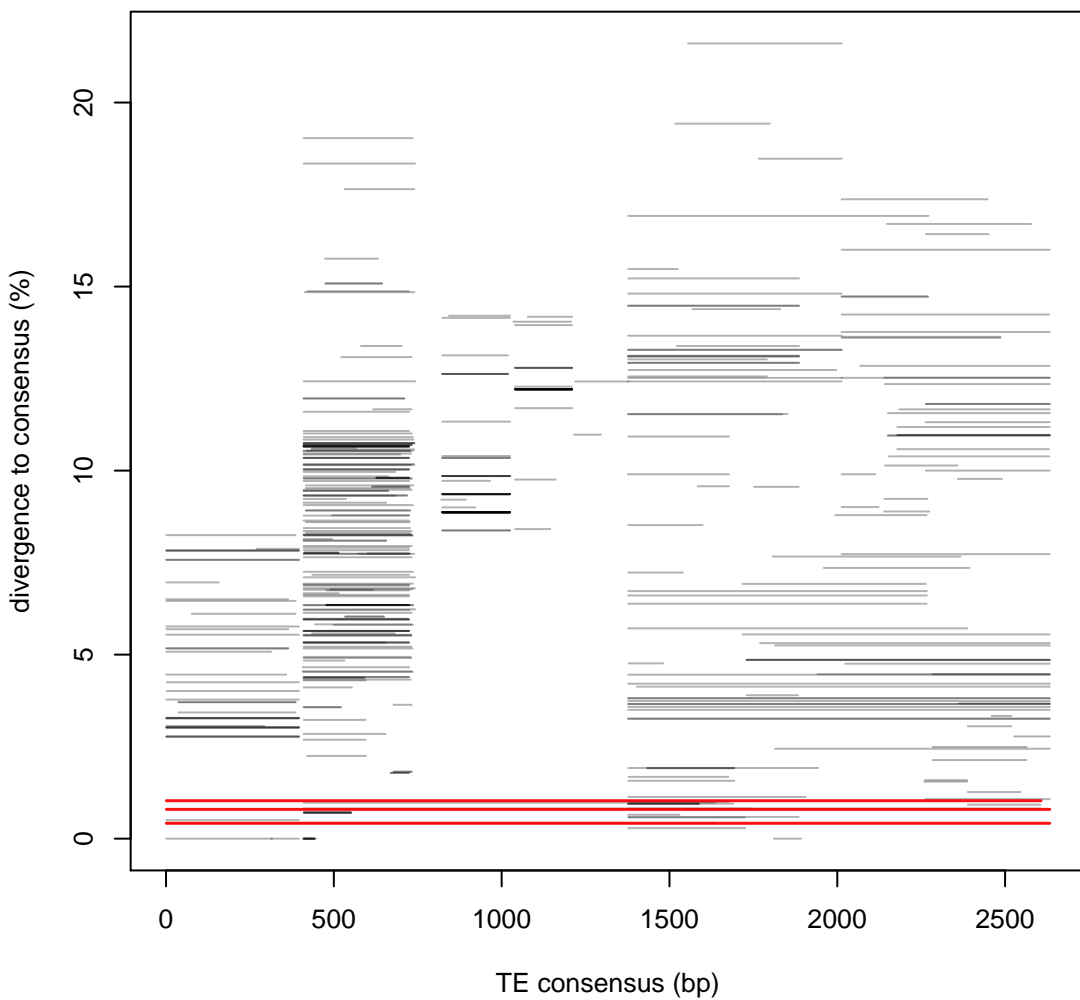
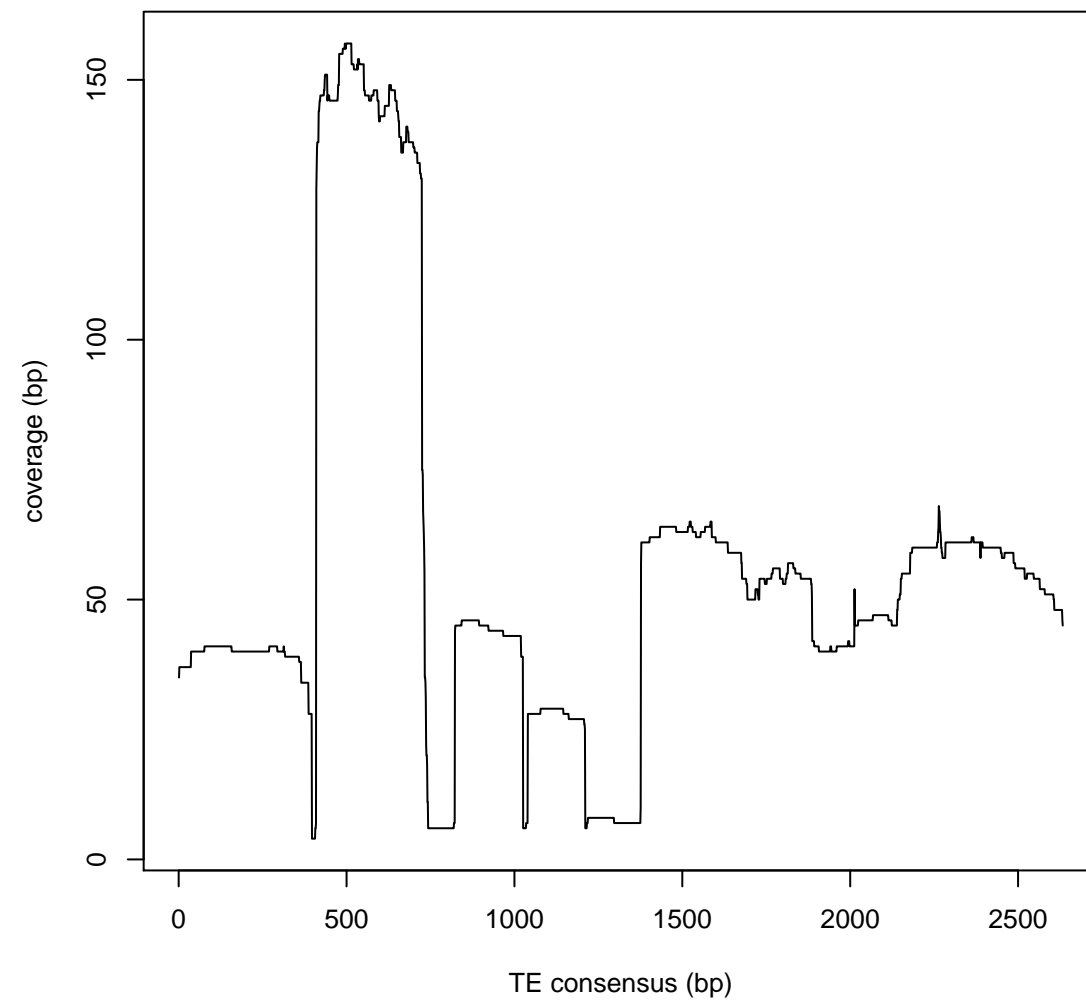


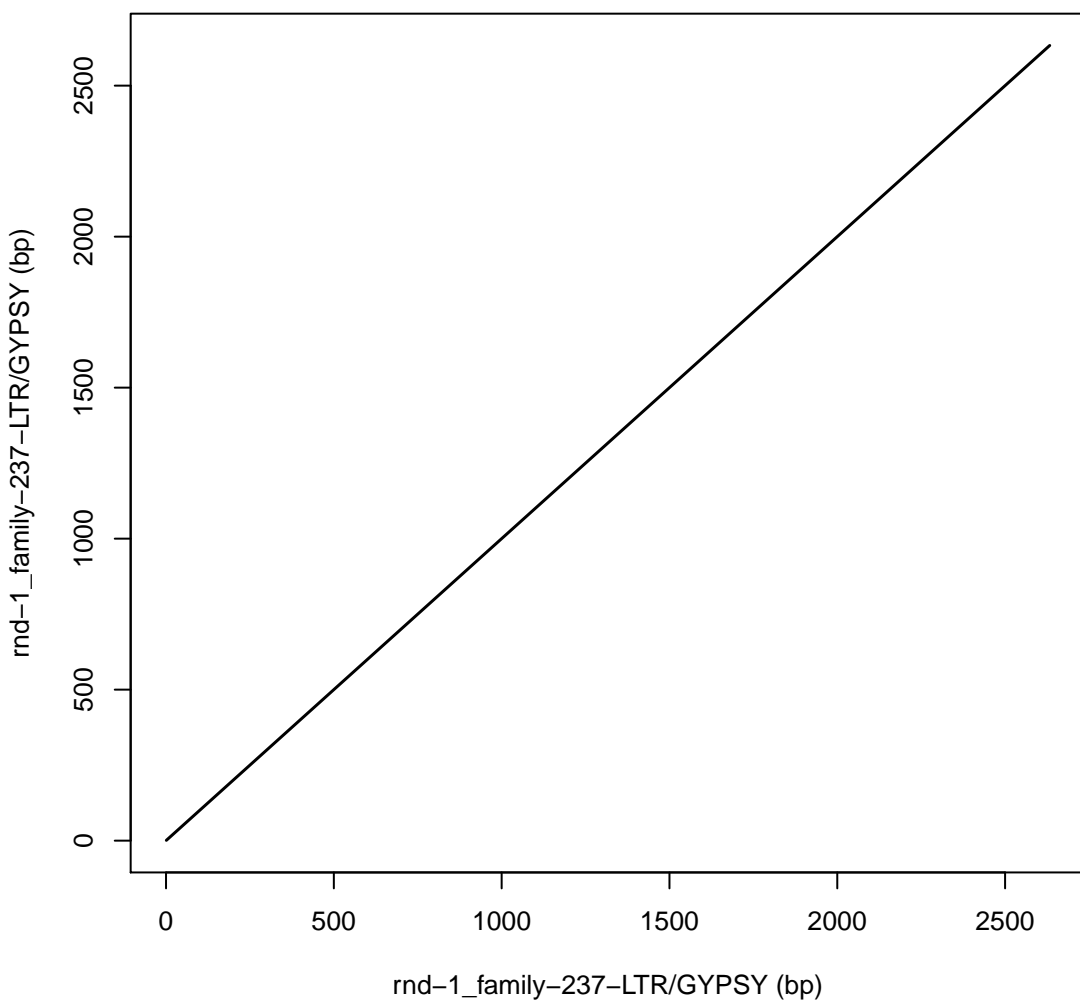
TE: rnd-1_family-237-LTR/GYPSY
consensus size: 2633bp; fragments: 413; full length: 3 (≥ 2369.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

