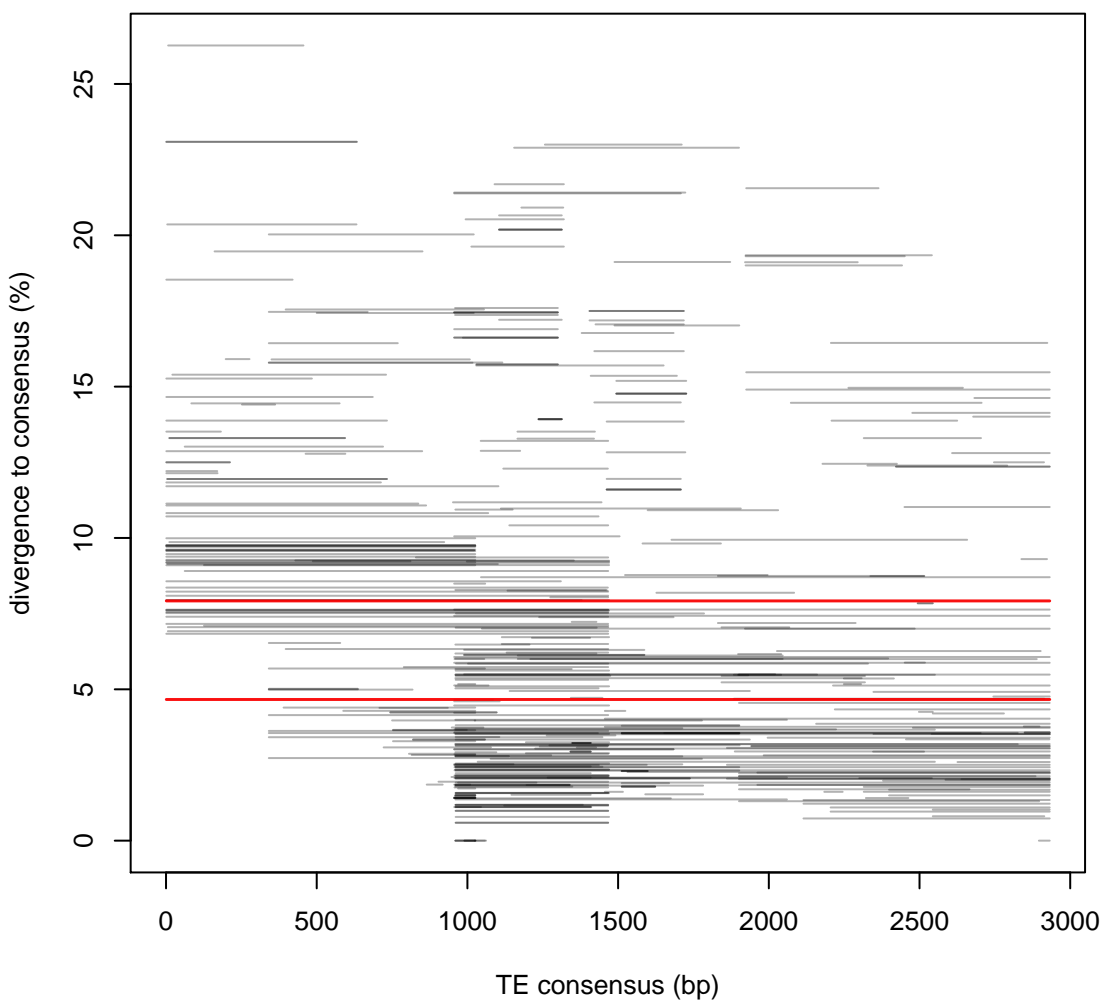
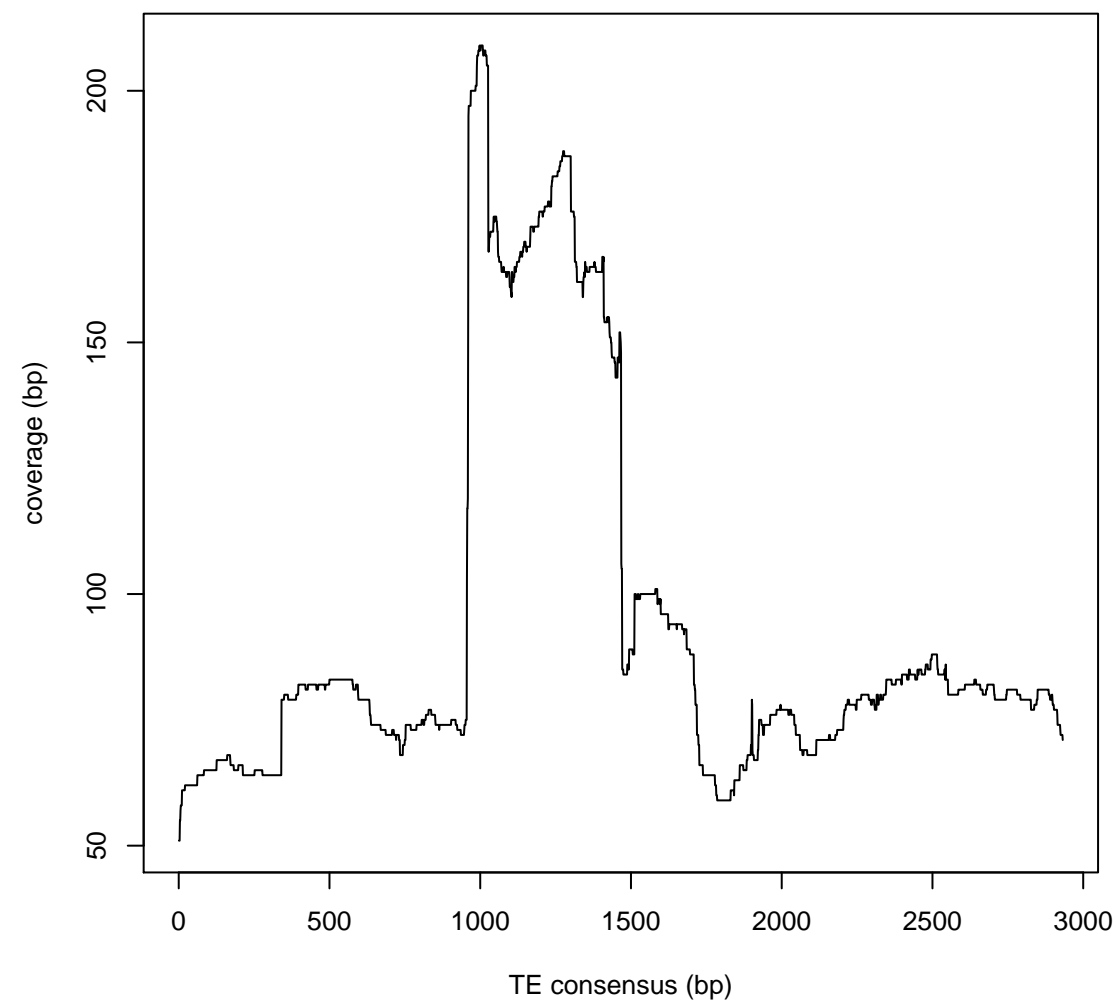


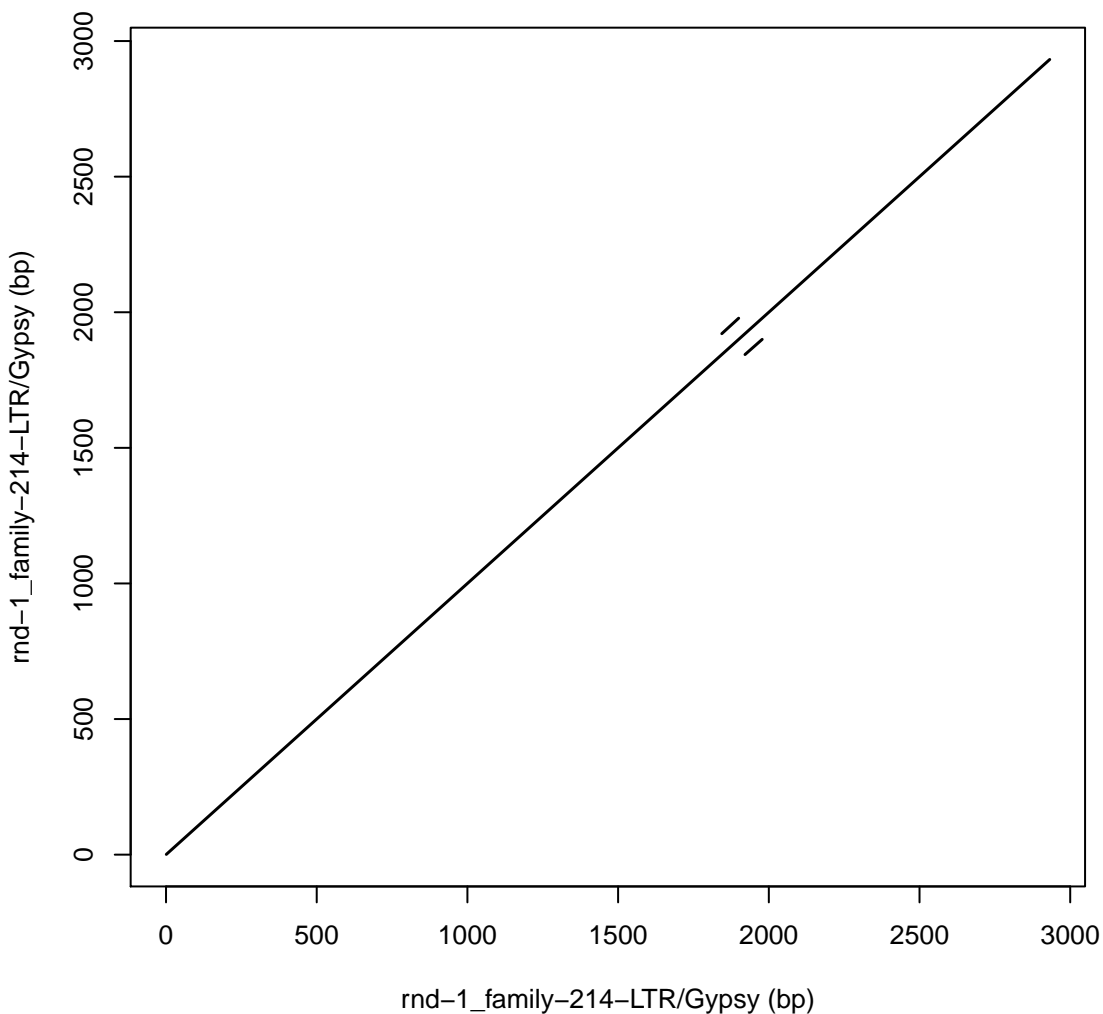
TE: rnd-1_family-214-LTR/Gypsy
consensus size: 2932bp; fragments: 446; full length: 2 (≥ 2638.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

