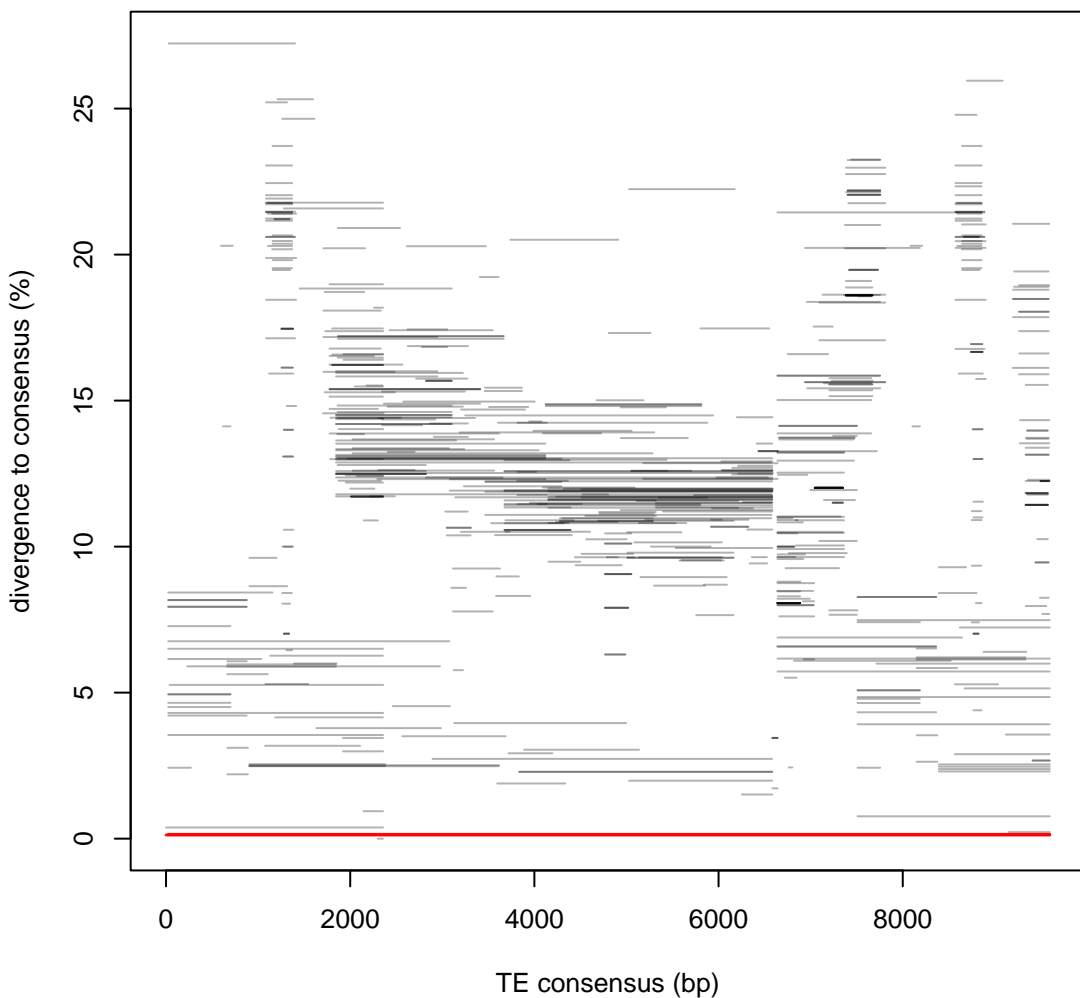
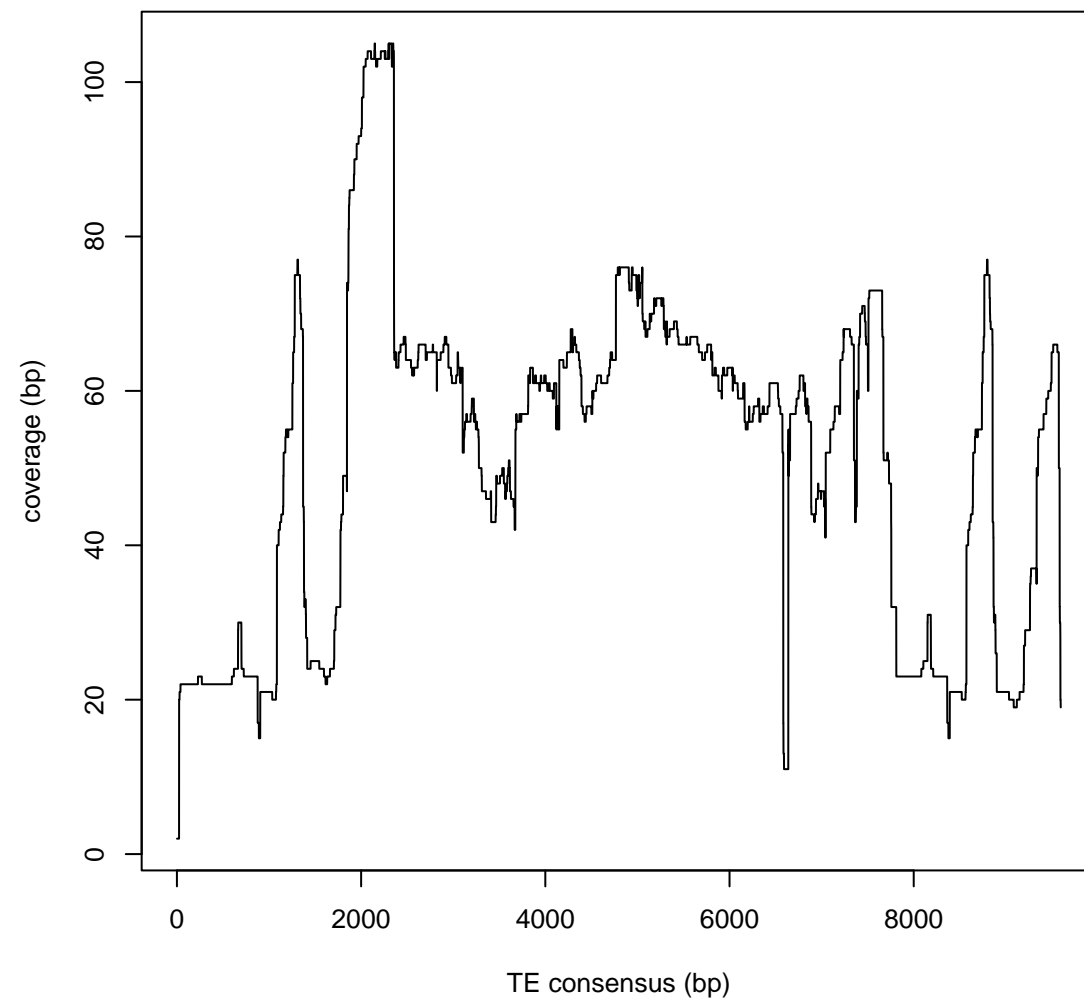


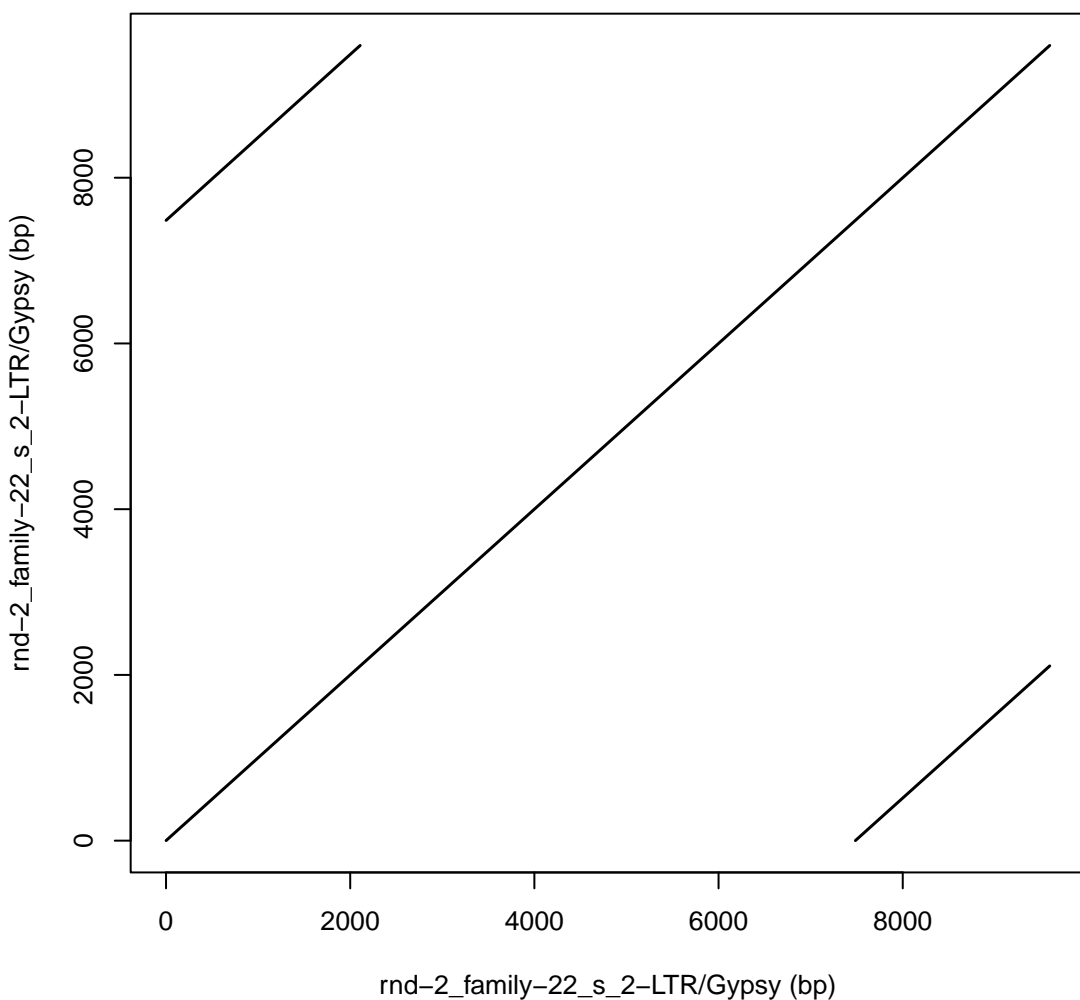
TE: rnd-2_family-22_s_2-LTR/Gypsy
consensus size: 9596bp; fragments: 651; full length: 2 (≥ 8636.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

