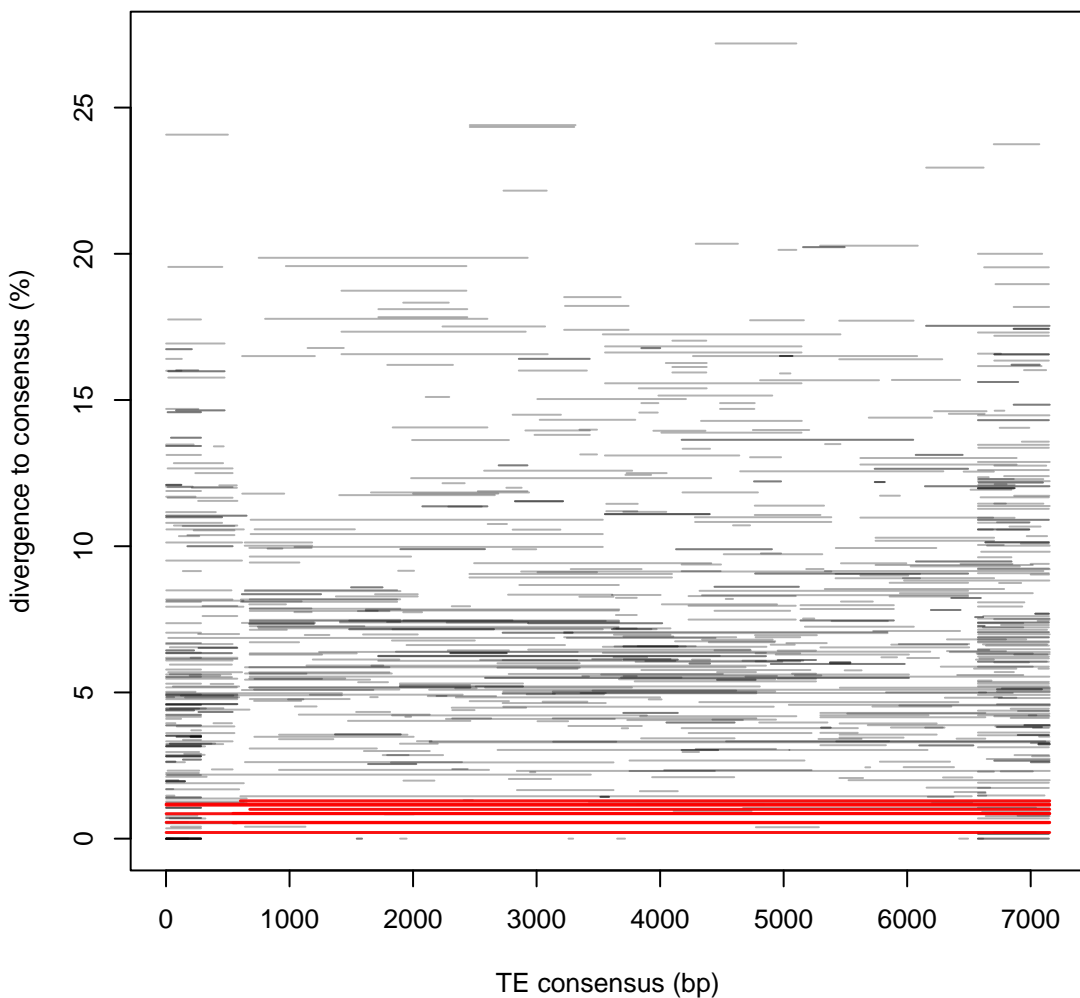
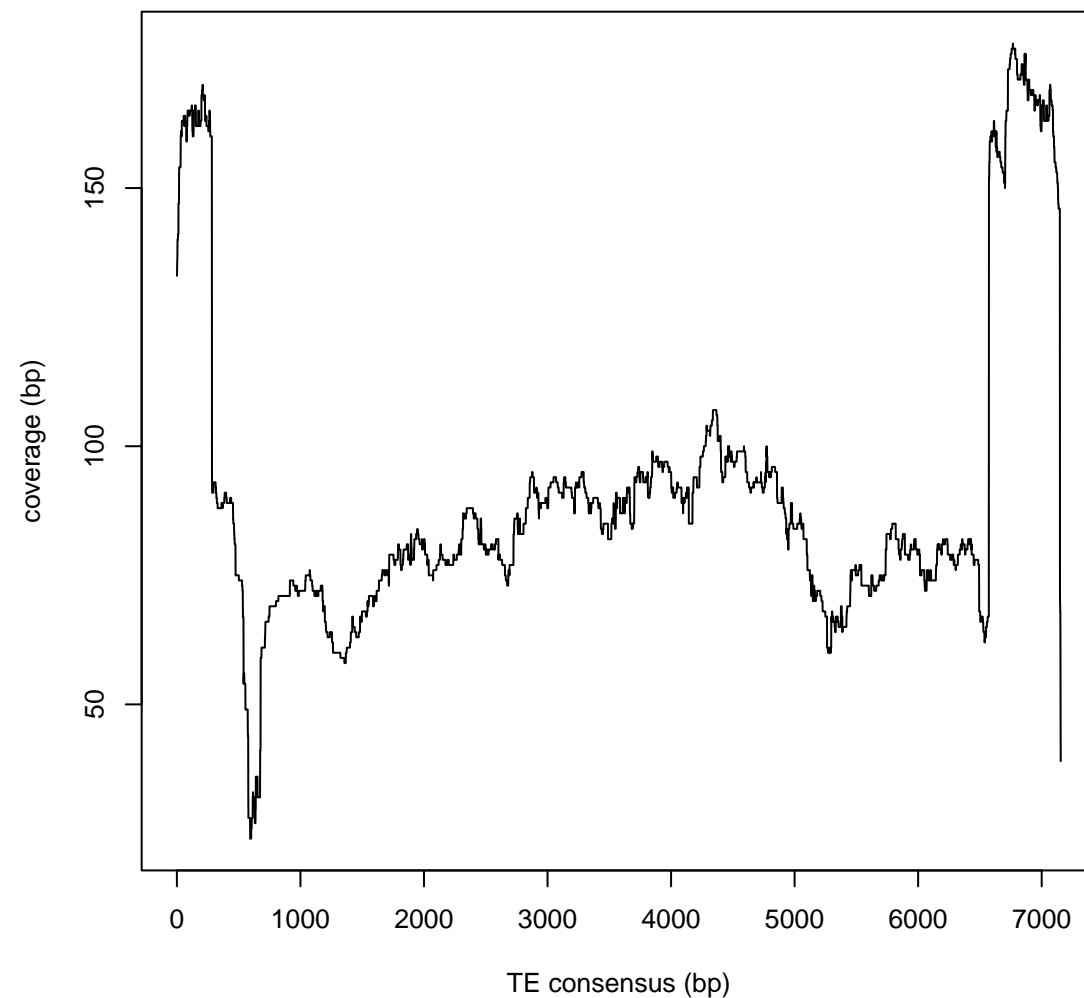


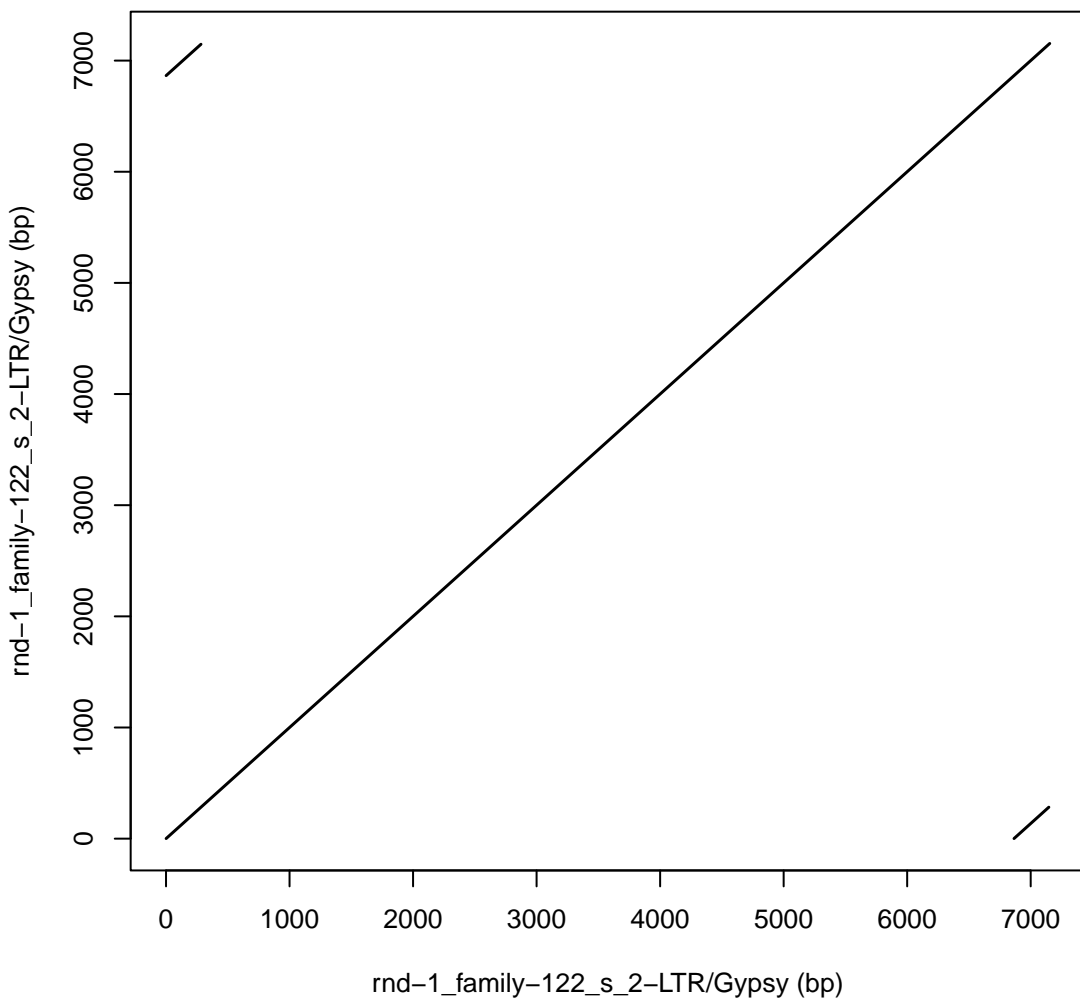
TE: rnd-1_family-122_s_2-LTR/Gypsy
consensus size: 7154bp; fragments: 913; full length: 9 (≥ 6438.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

