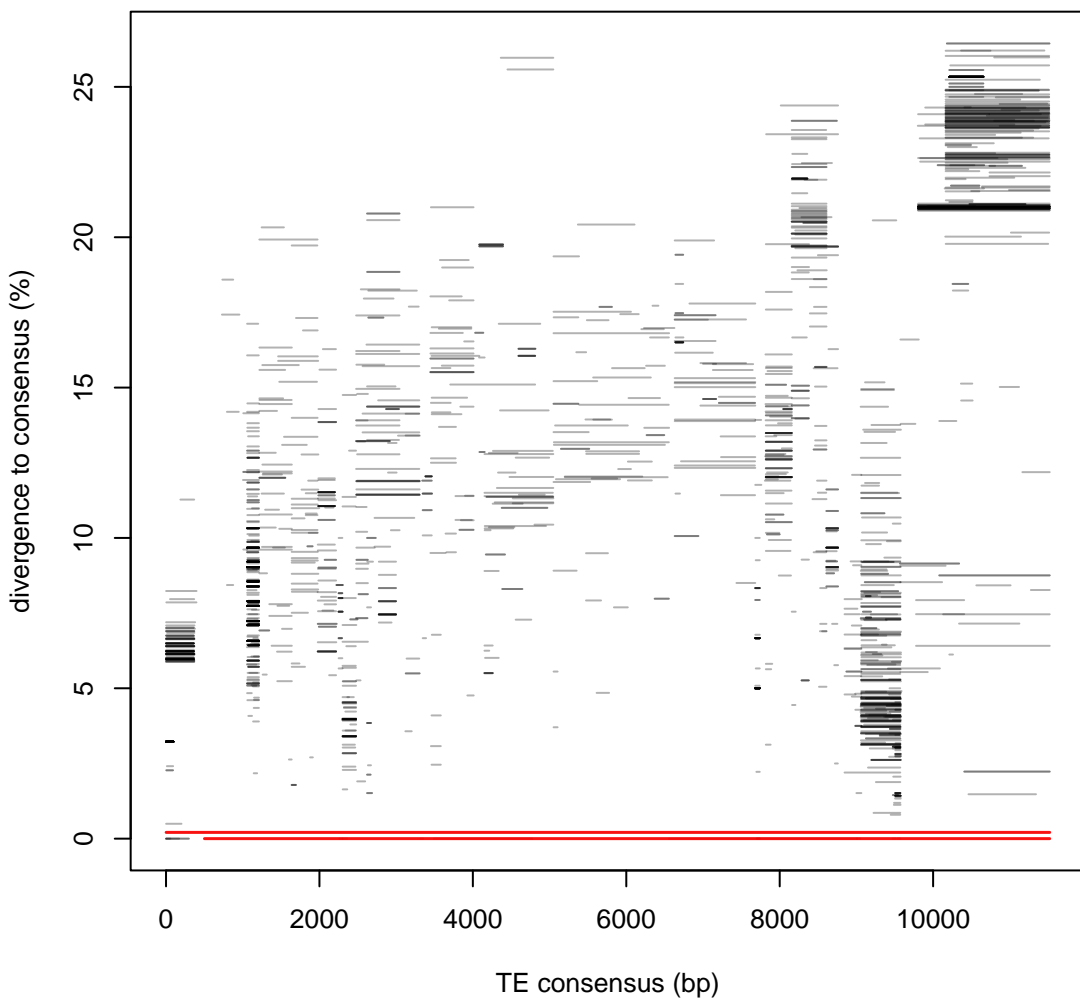
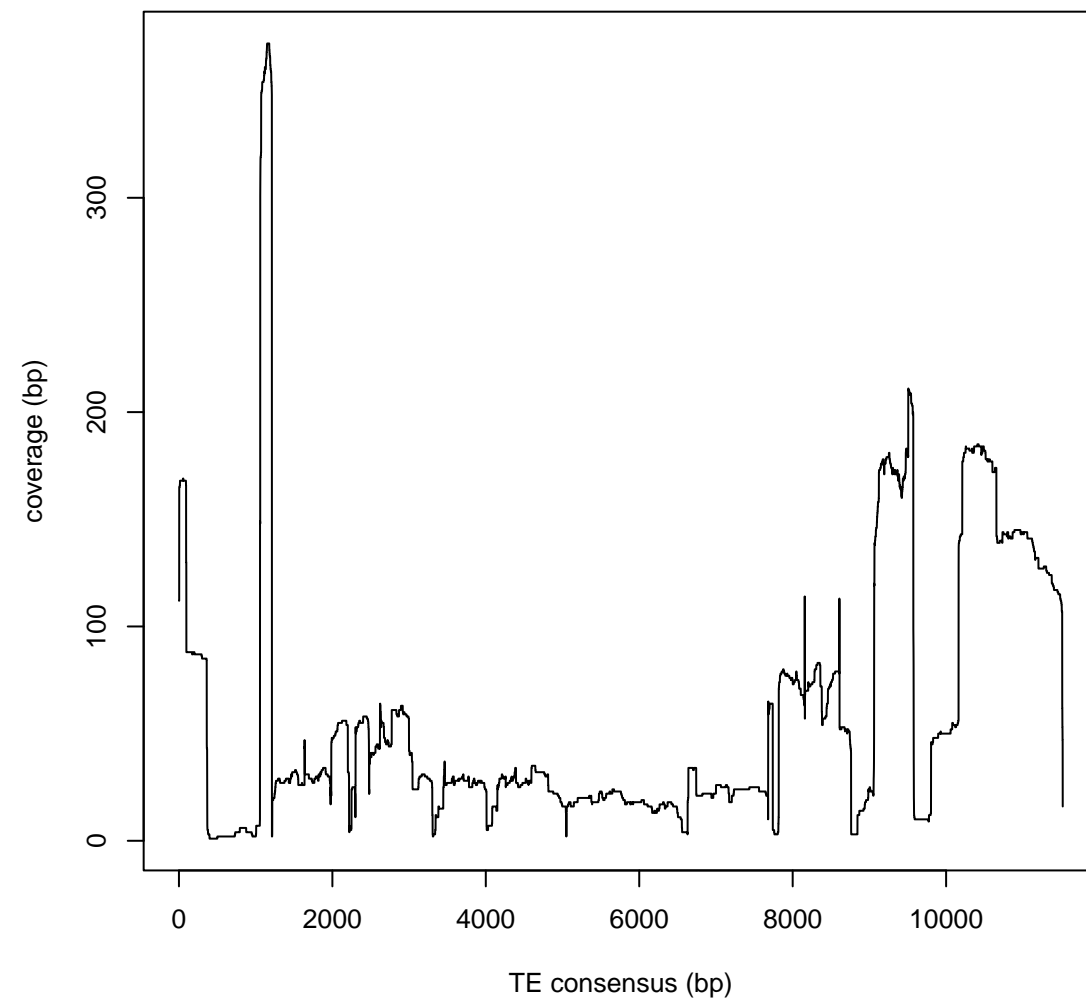


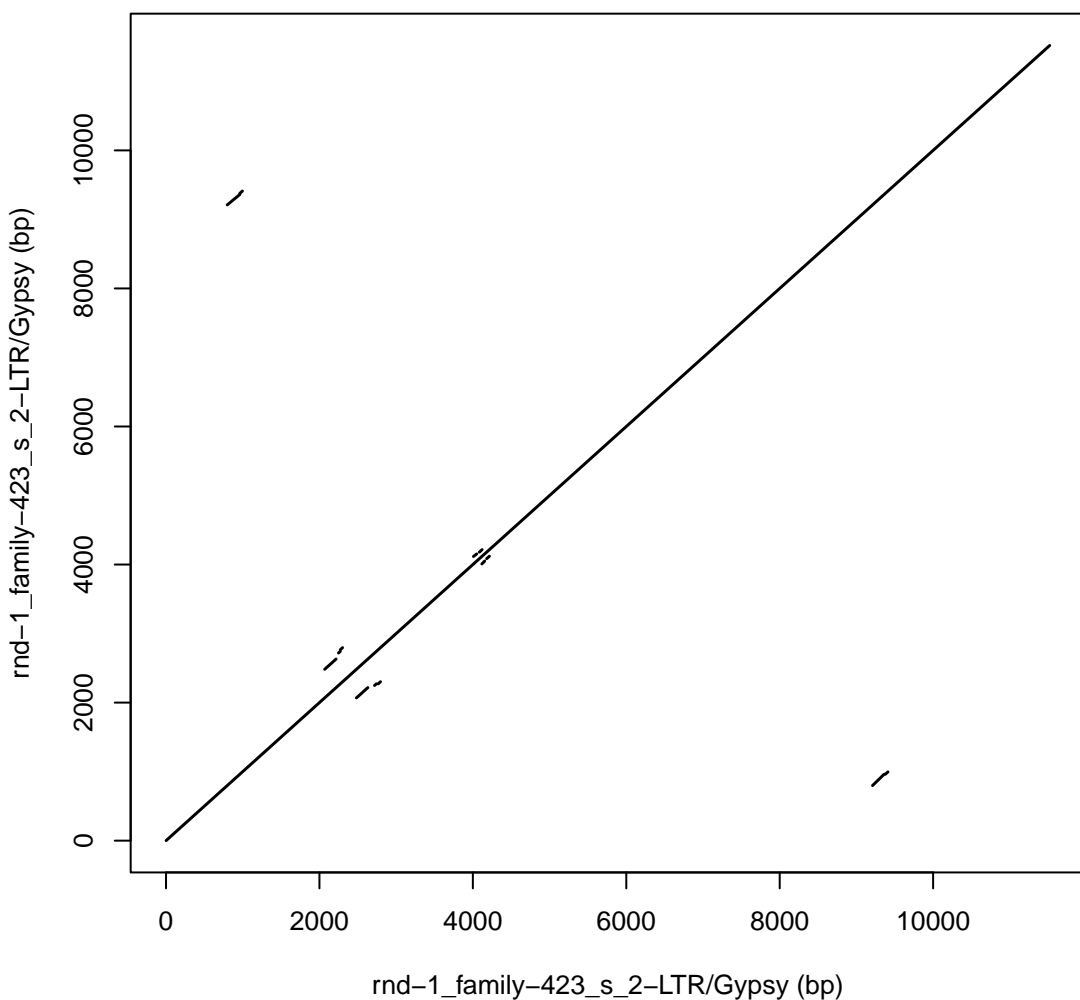
TE: rnd-1_family-423_s_2-LTR/Gypsy
consensus size: 11520bp; fragments: 1879; full length: 2 (>=10368bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

