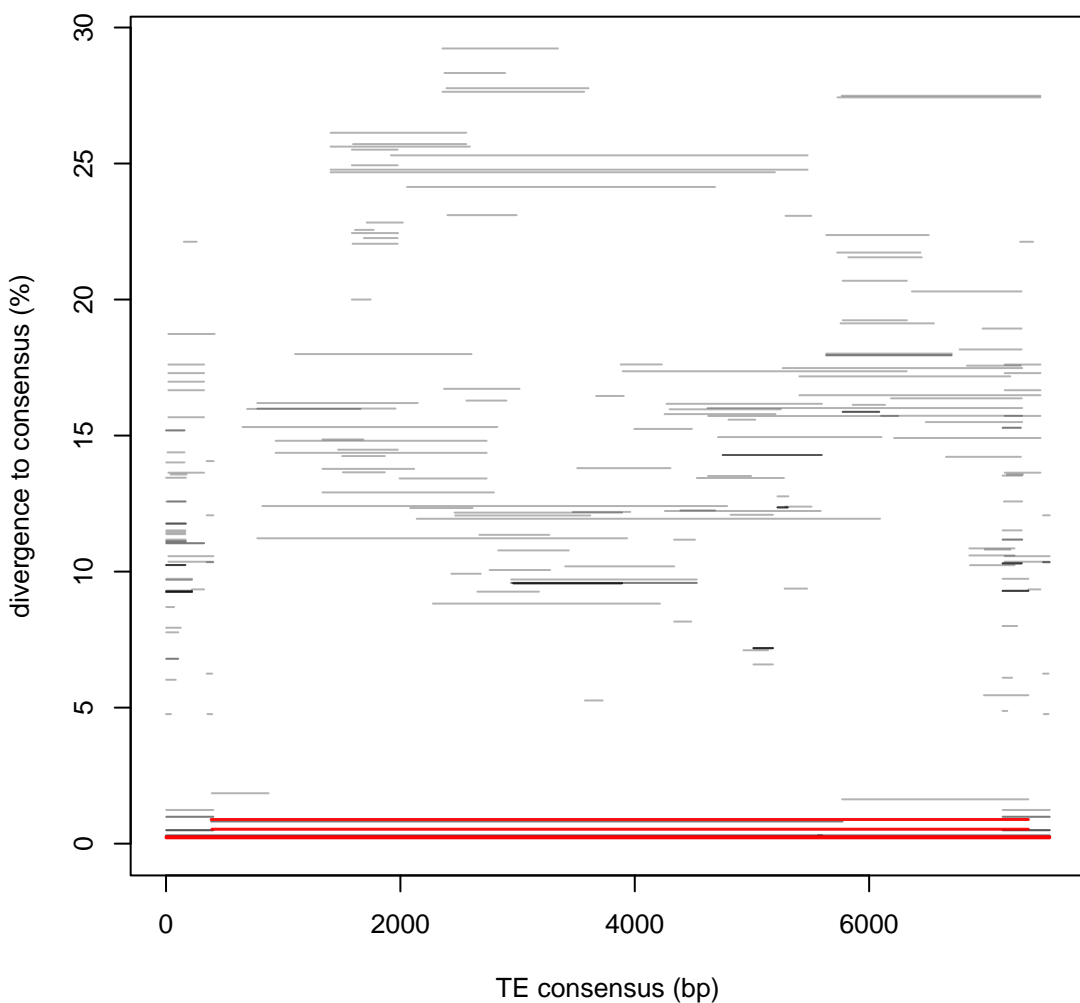
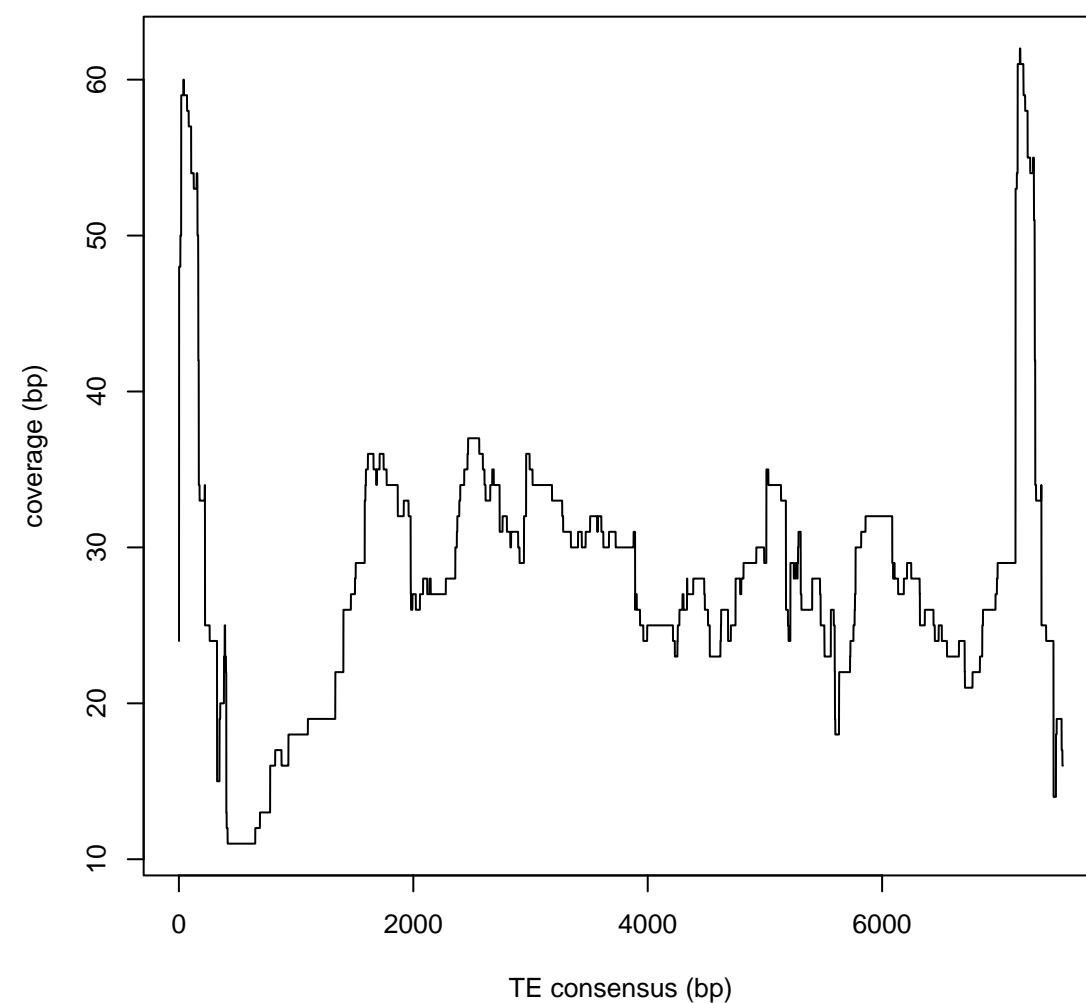


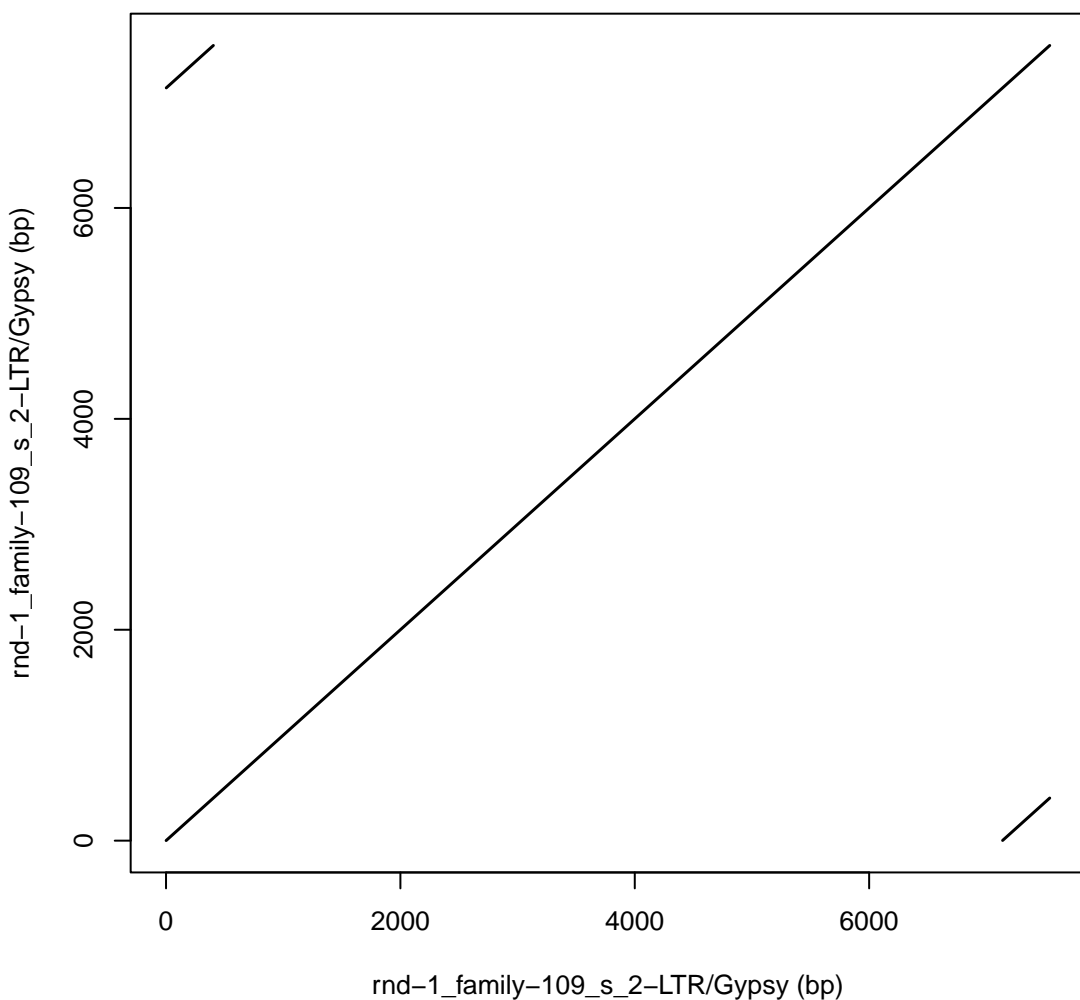
TE: rnd-1_family-109_s_2-LTR/Gypsy
consensus size: 7541bp; fragments: 243; full length: 5 (≥ 6786.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

