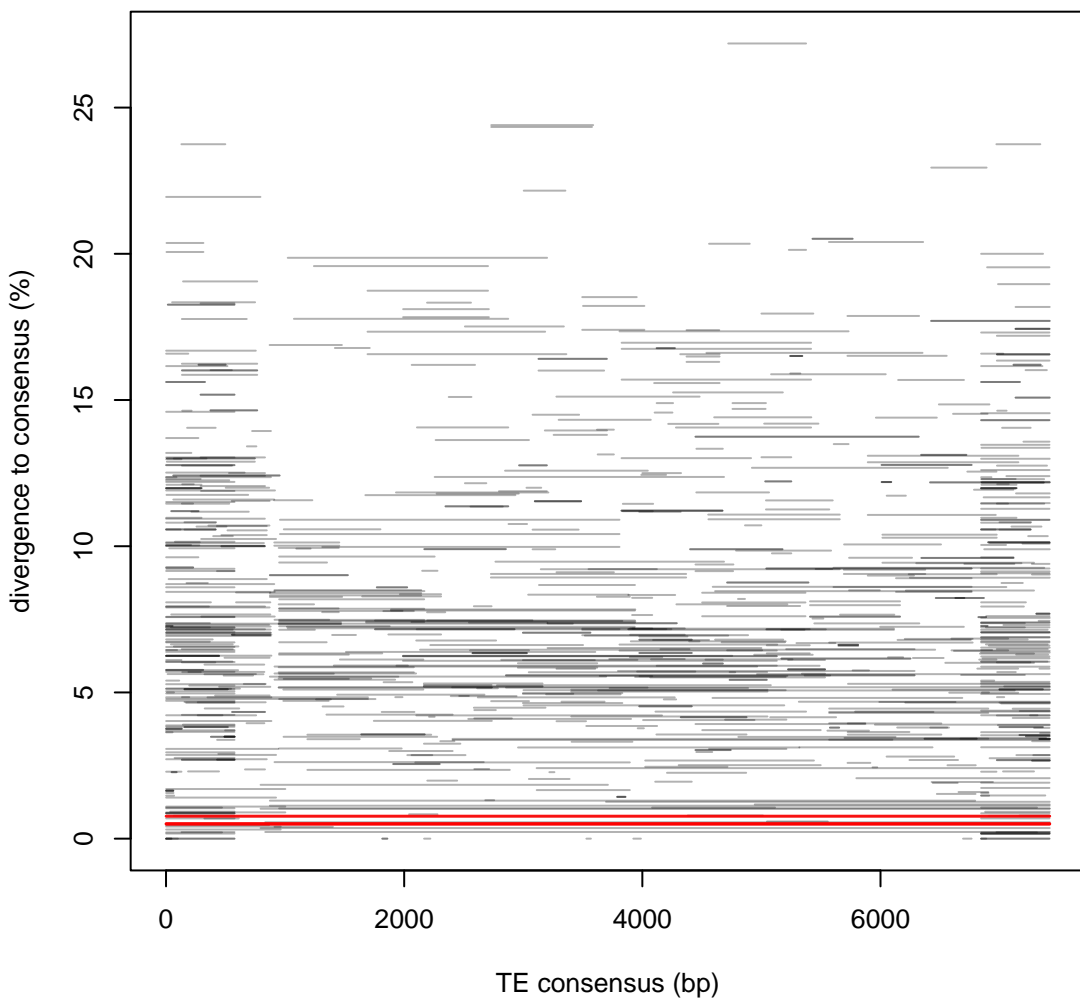
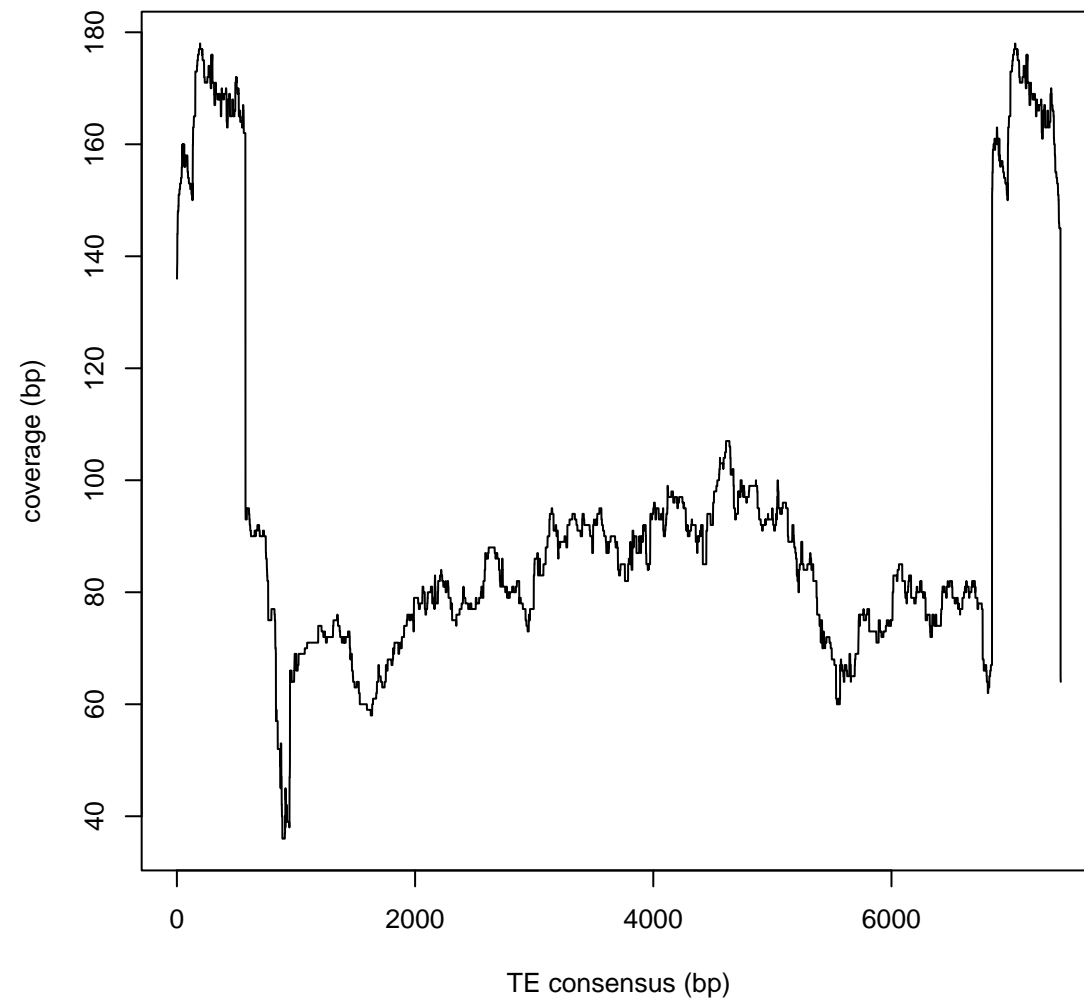


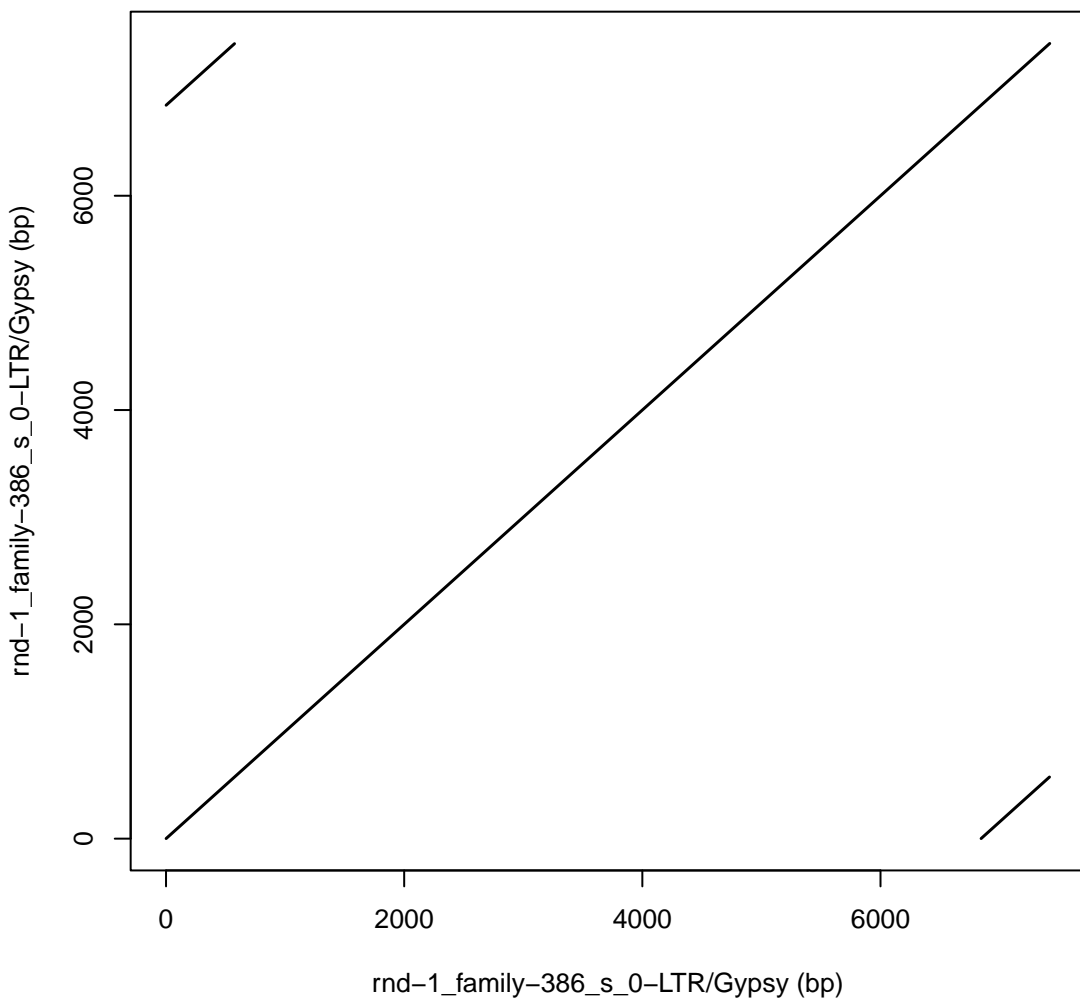
TE: rnd-1_family-386_s_0-LTR/Gypsy
consensus size: 7421bp; fragments: 973; full length: 3 (≥ 6678.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

