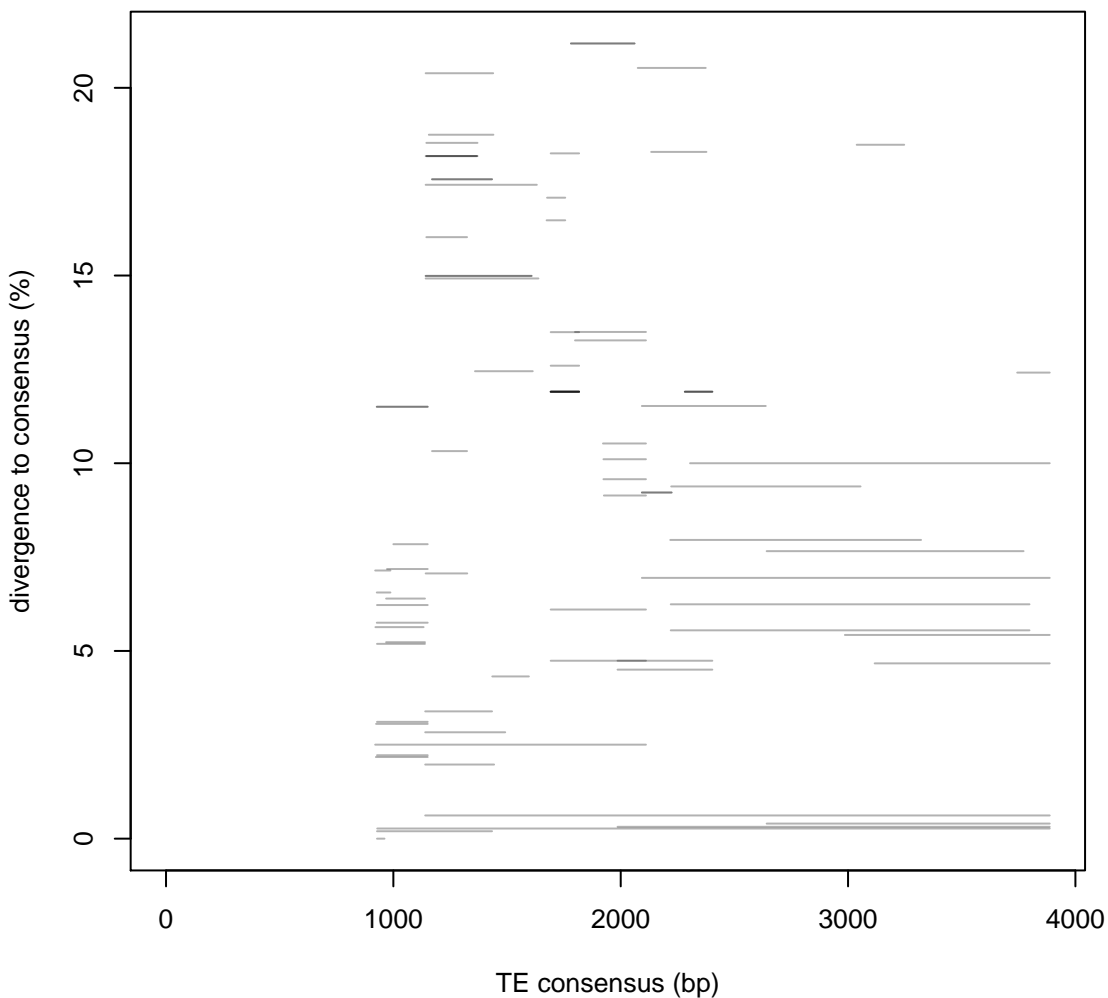
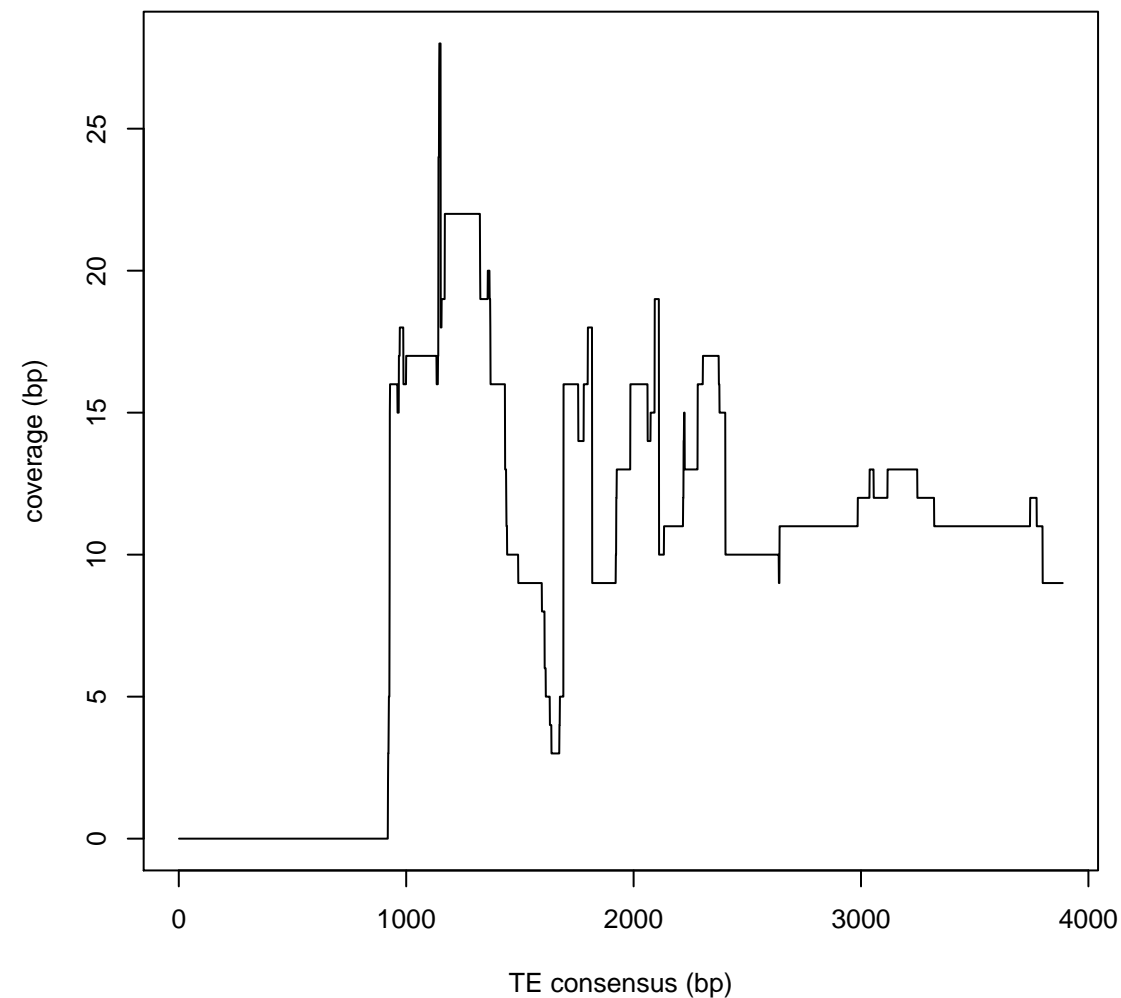


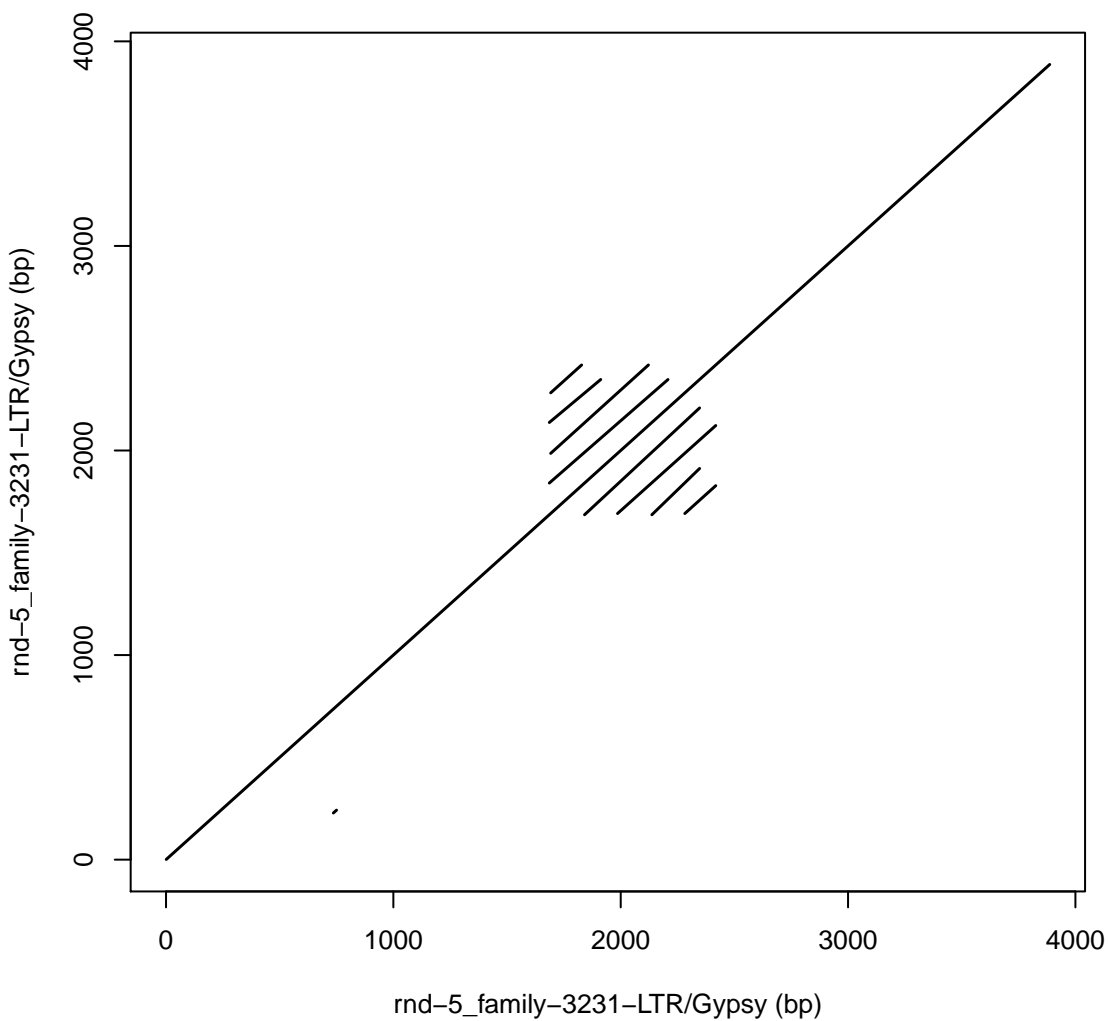
TE: rnd-5_family-3231-LTR/Gypsy
consensus size: 3887bp; fragments: 85; full length: 0 (≥ 3498.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

