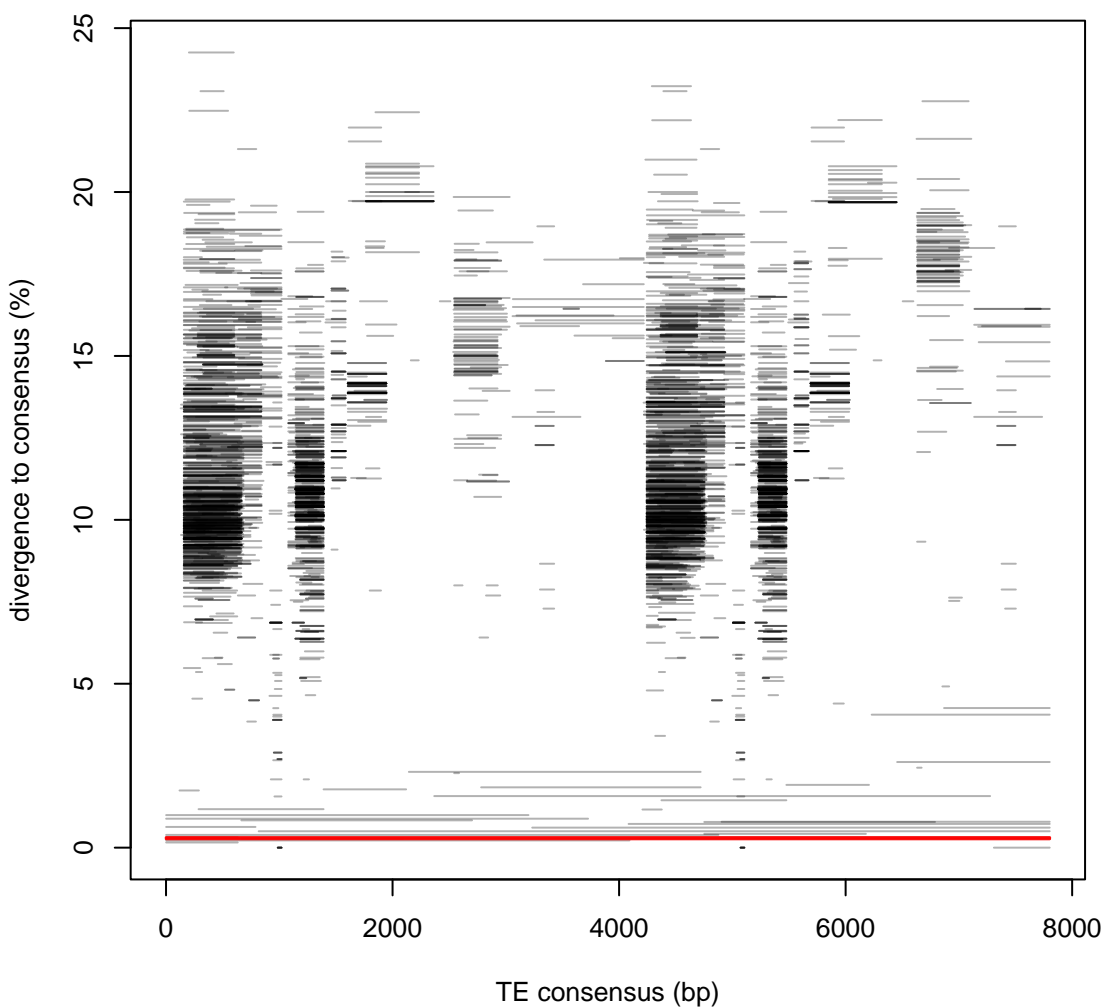
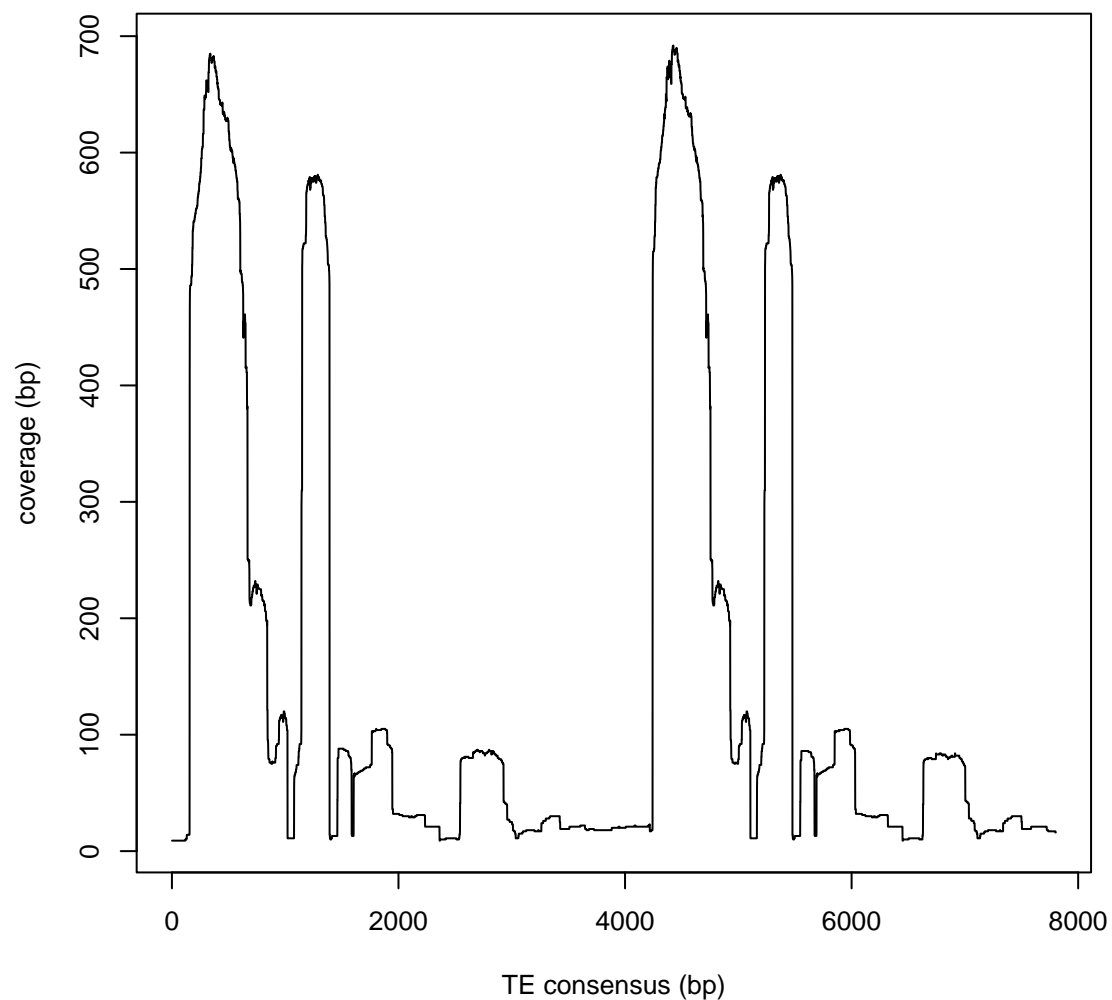


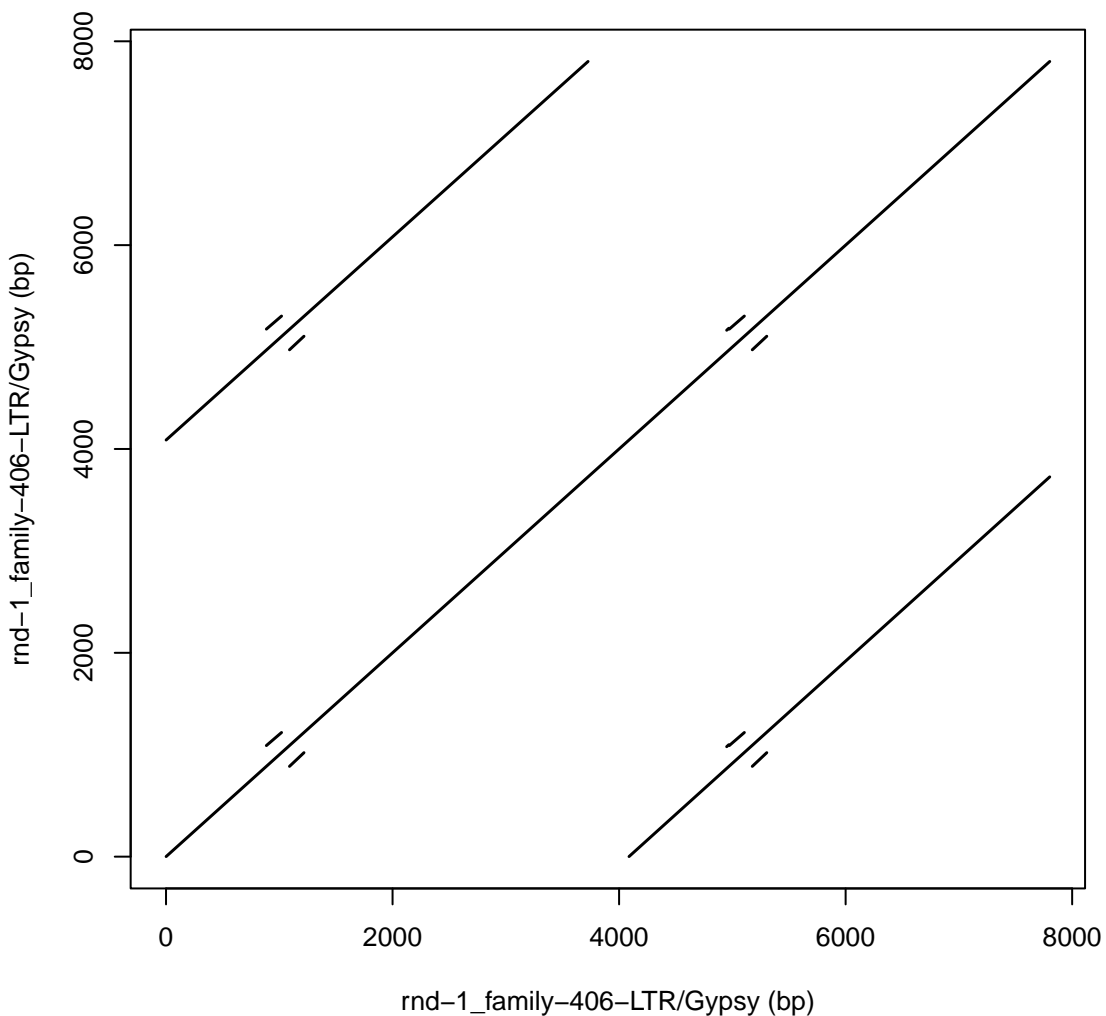
TE: rnd-1_family-406-LTR/Gypsy
consensus size: 7802bp; fragments: 3622; full length: 2 (≥ 7021.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

