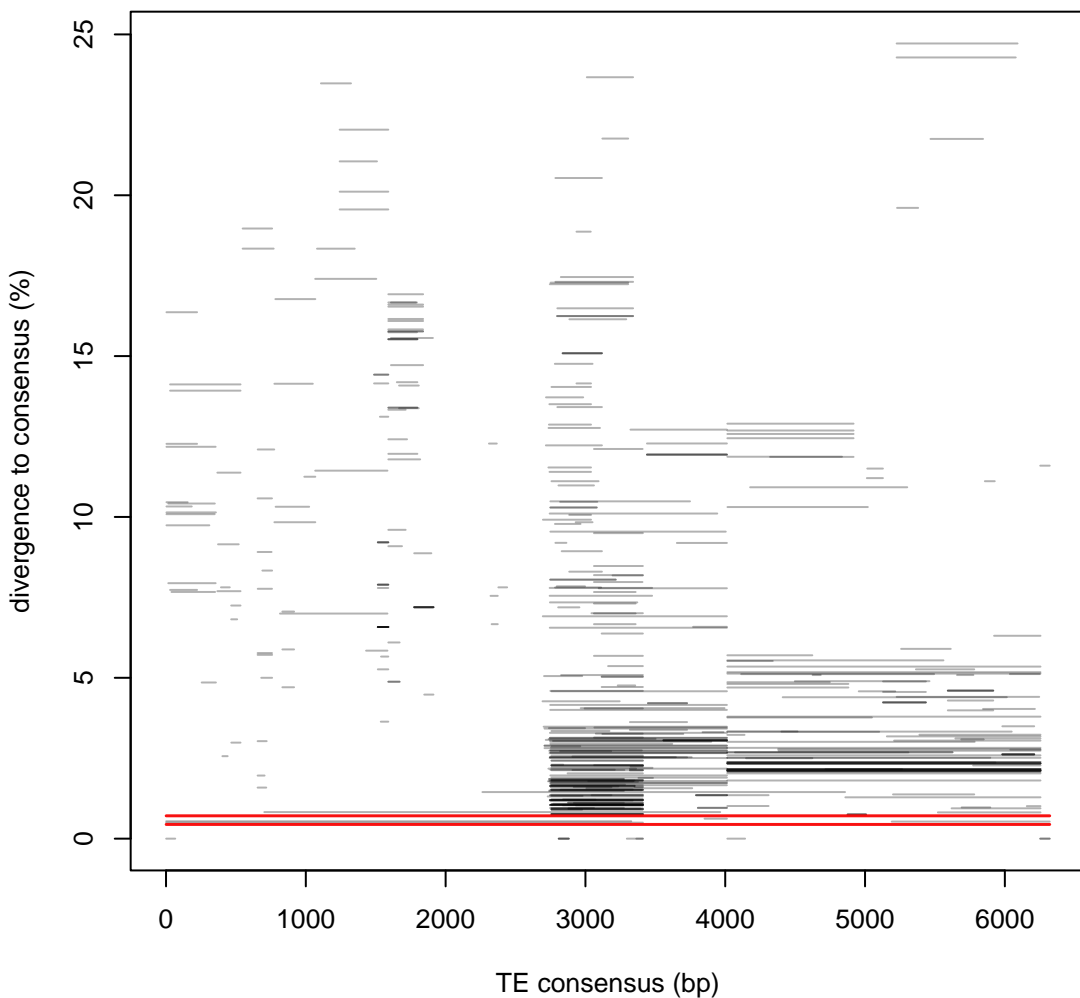
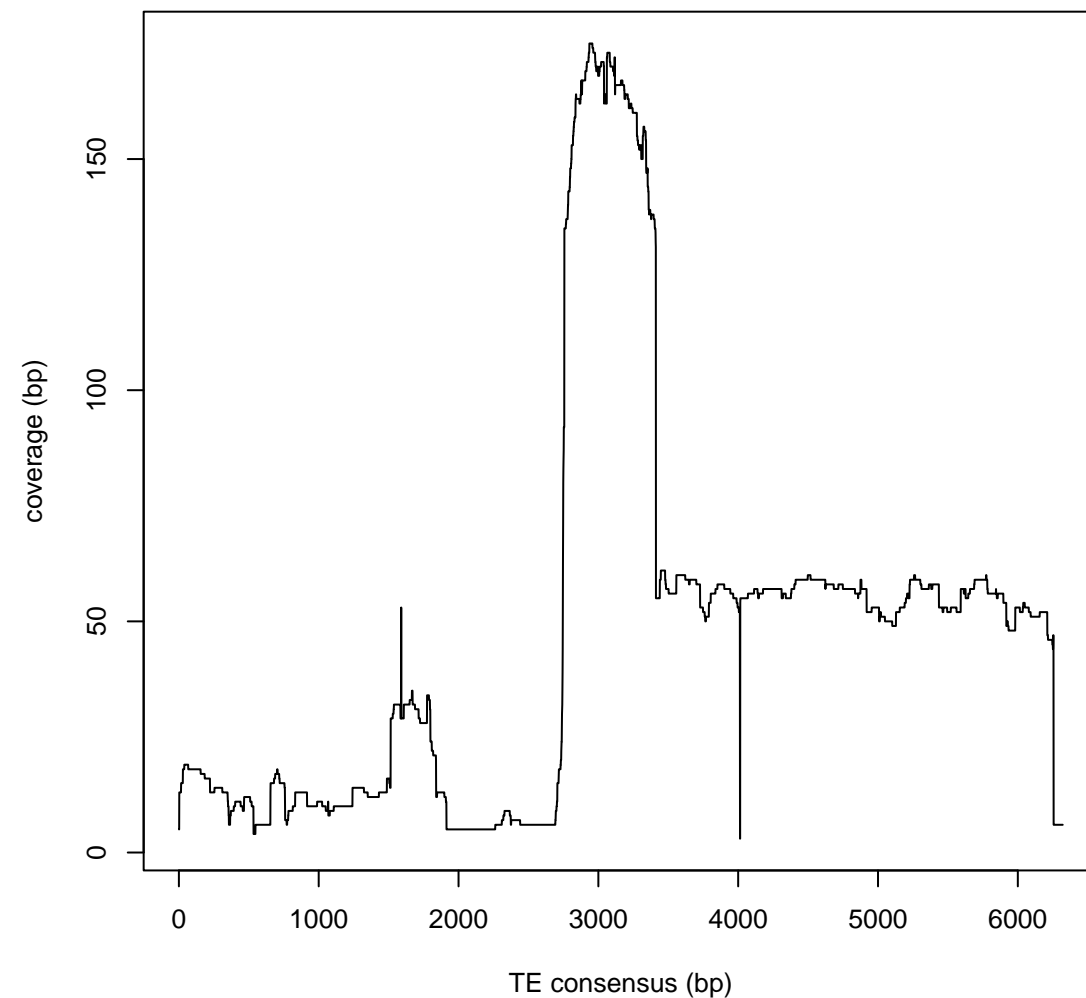


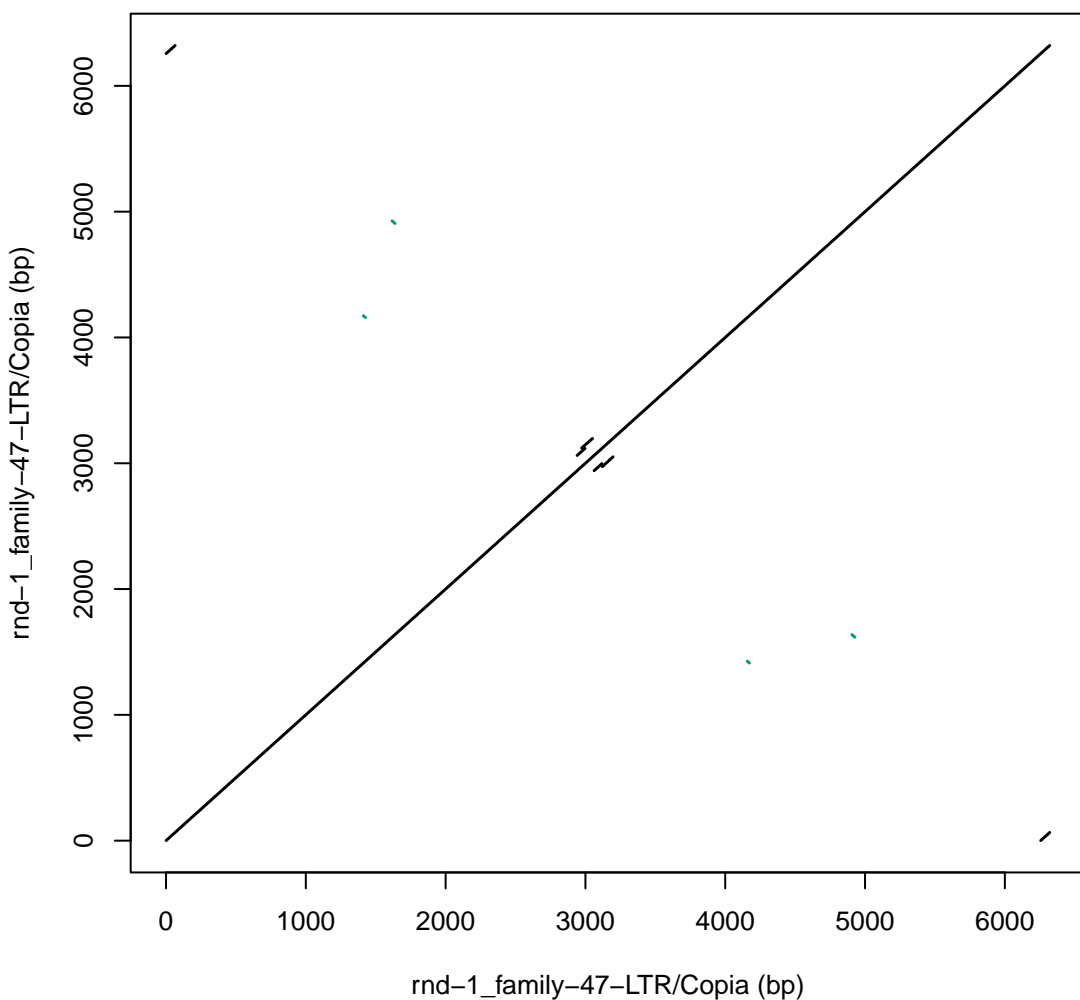
TE: rnd-1_family-47-LTR/Copia
consensus size: 6321bp; fragments: 472; full length: 2 (≥ 5688.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

