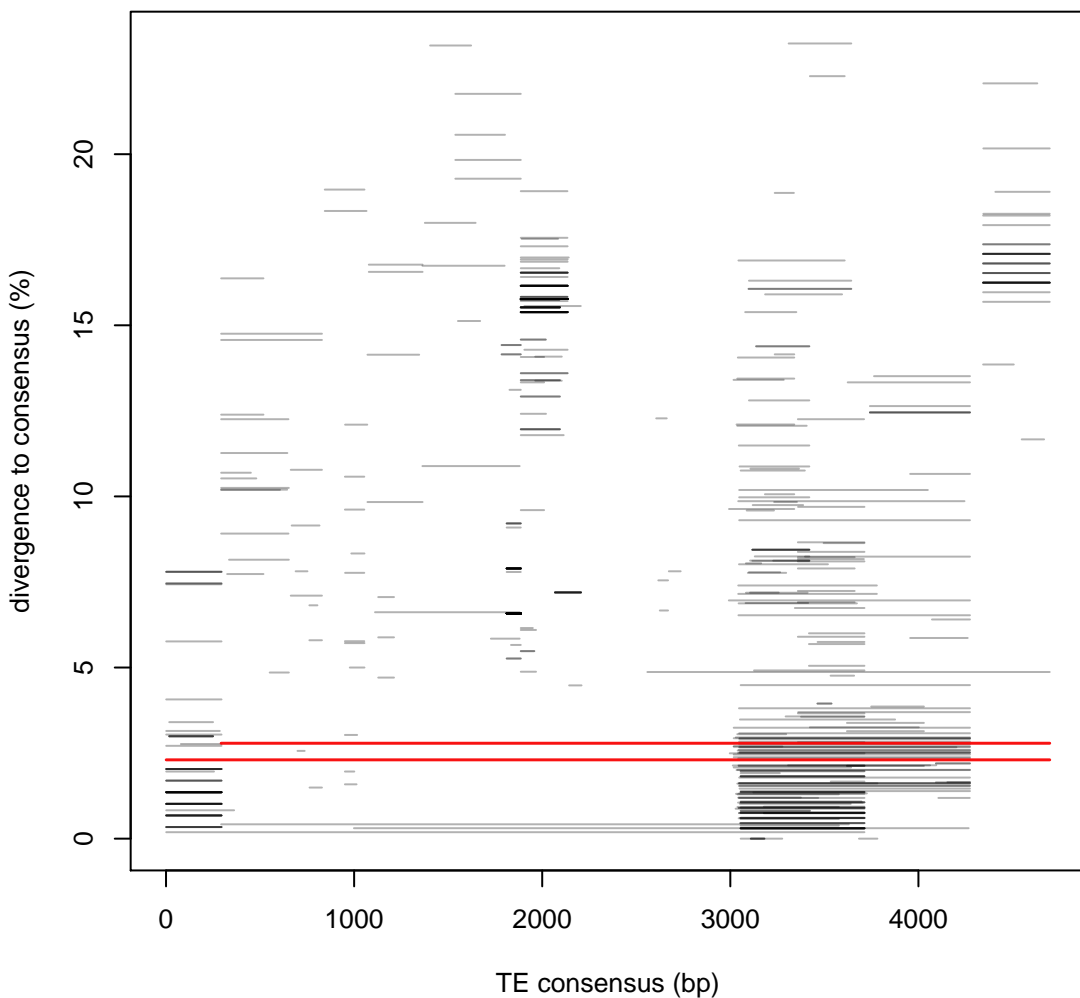
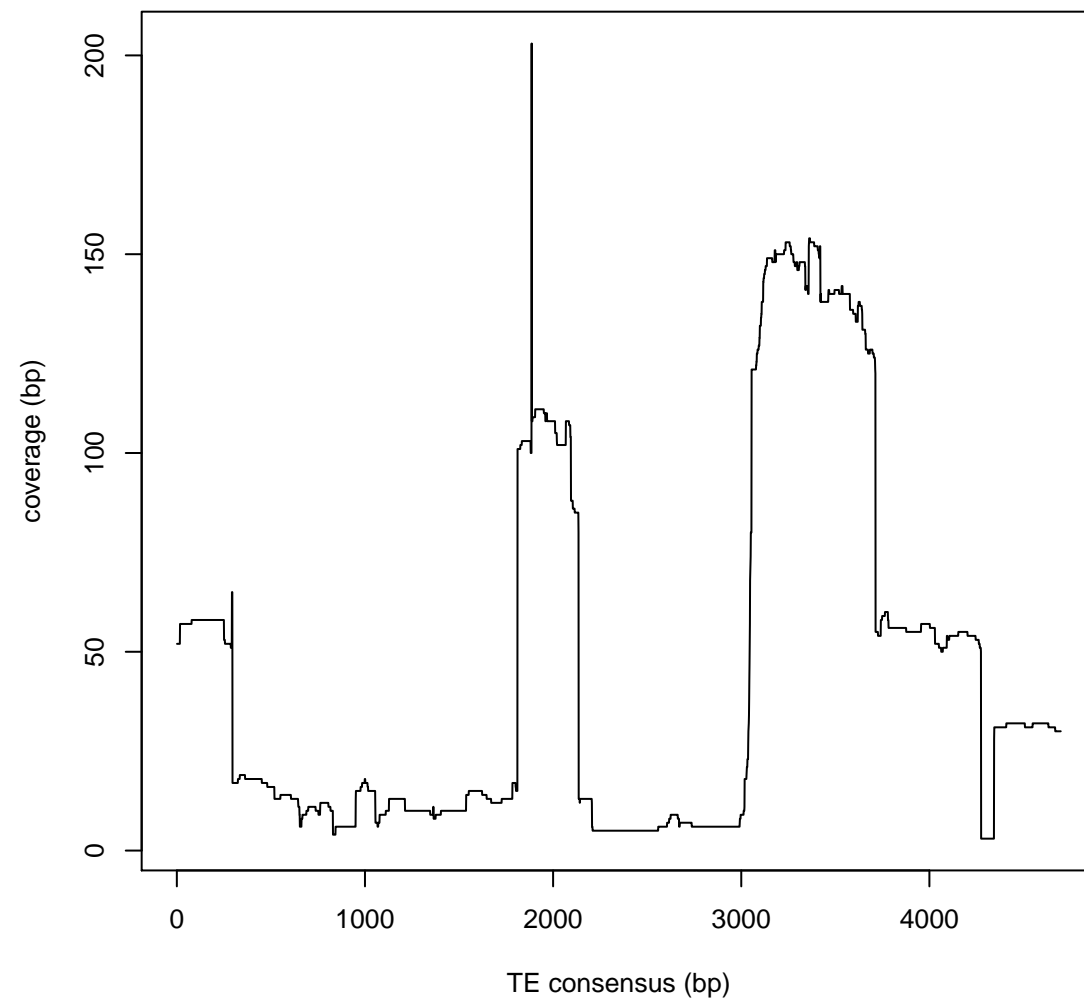


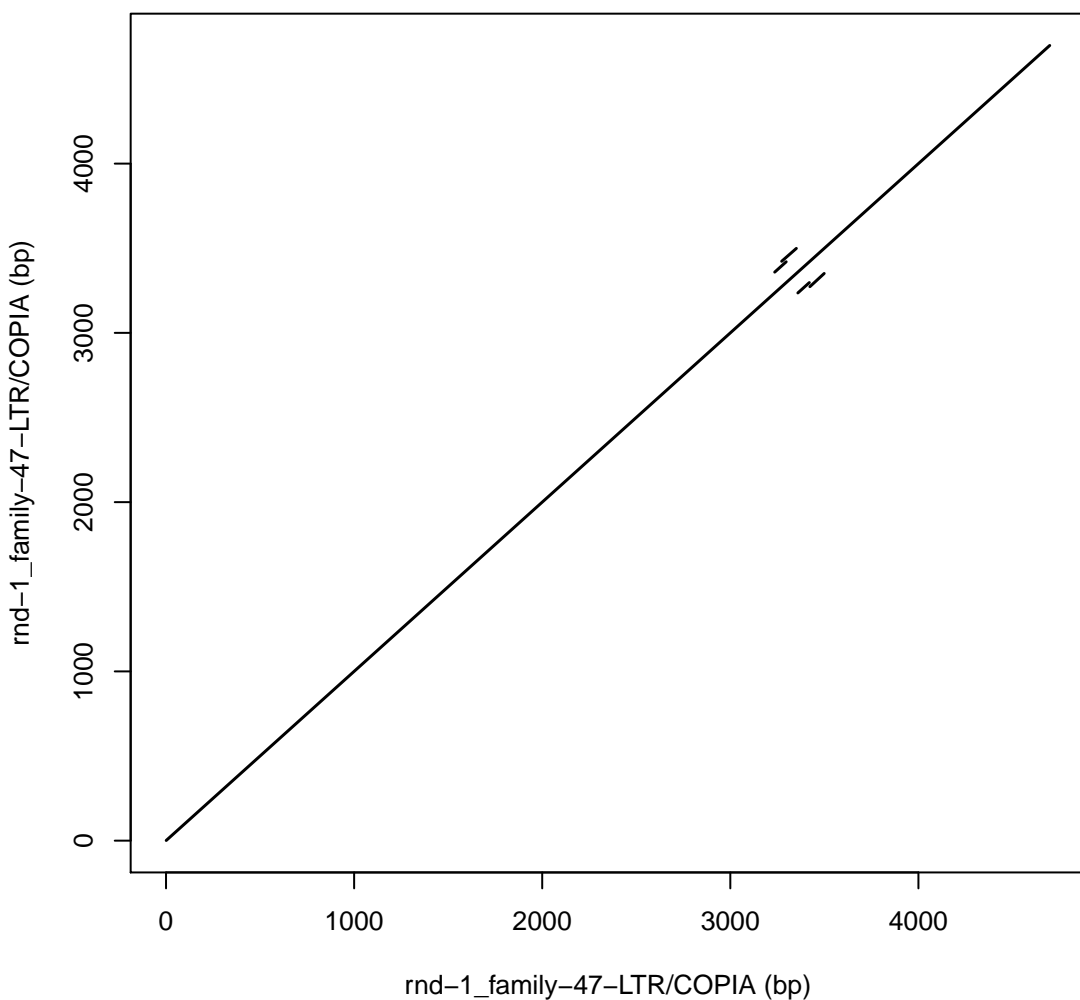
TE: rnd-1_family-47-LTR/COPIA
consensus size: 4698bp; fragments: 560; full length: 2 (>=4228.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

