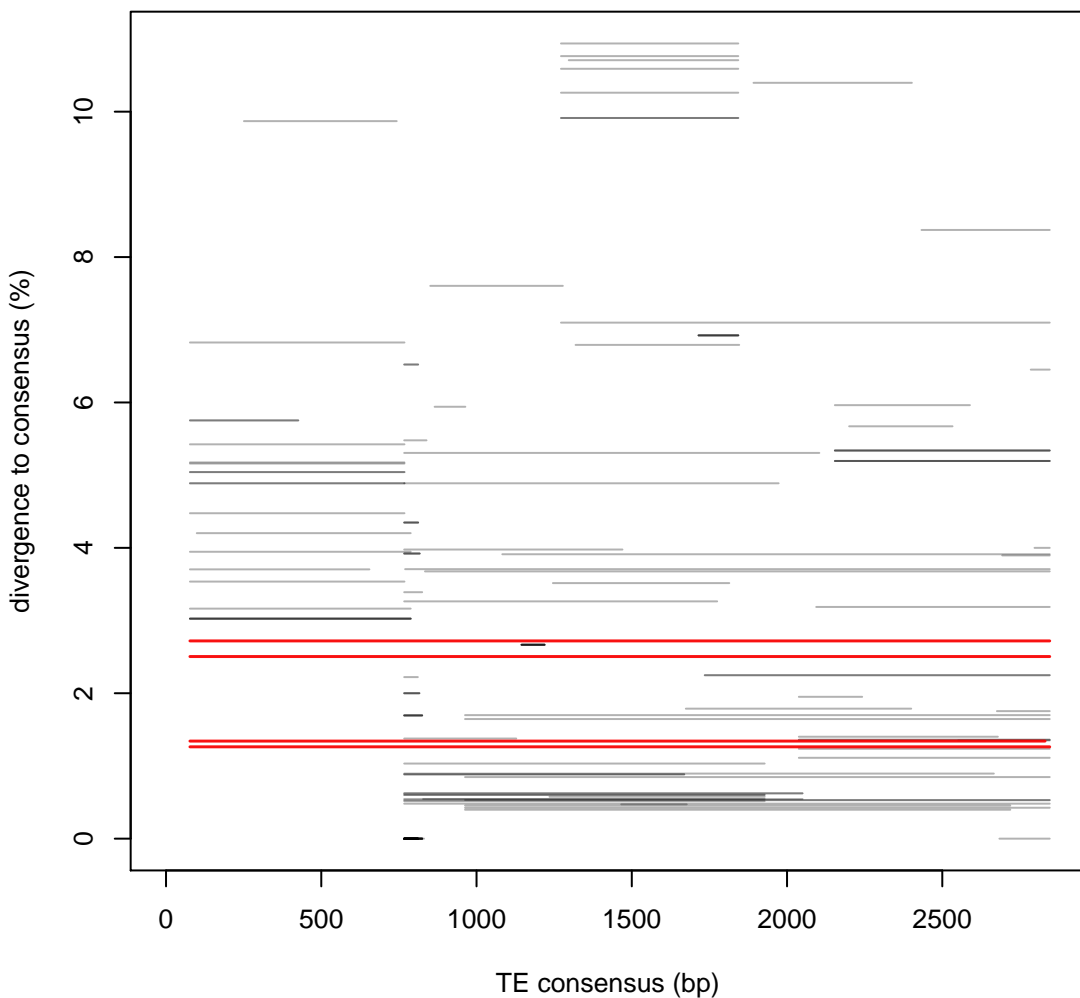
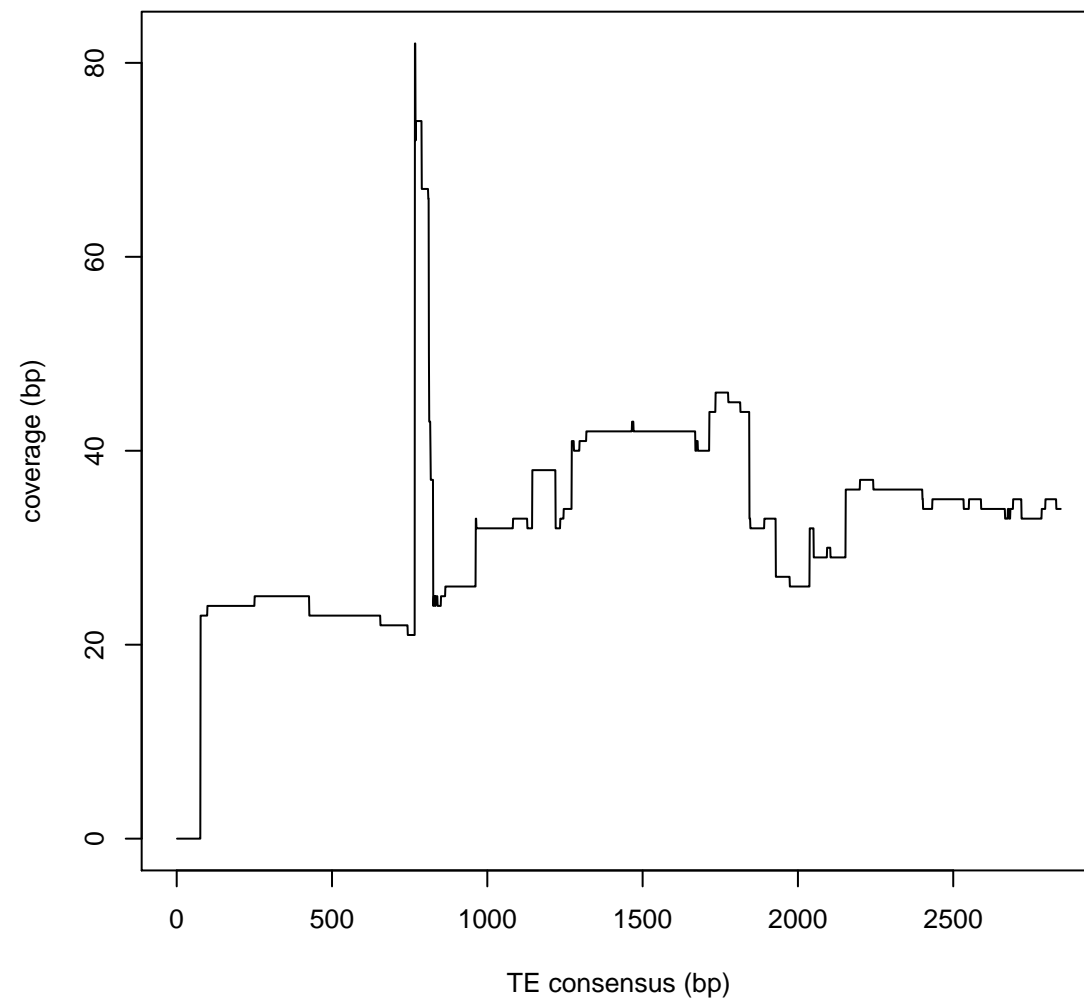


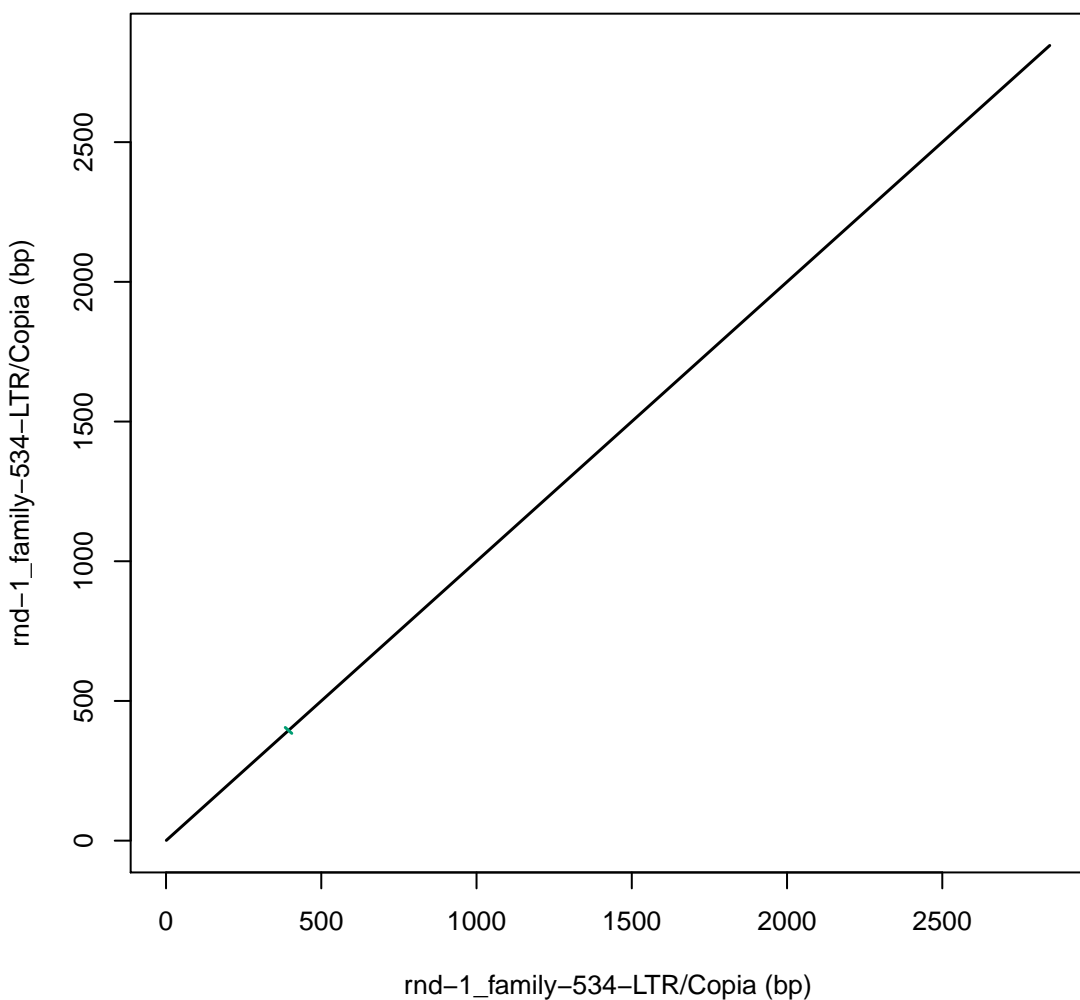
TE: rnd-1_family-534-LTR/Copia
consensus size: 2846bp; fragments: 148; full length: 4 (≥ 2561.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

