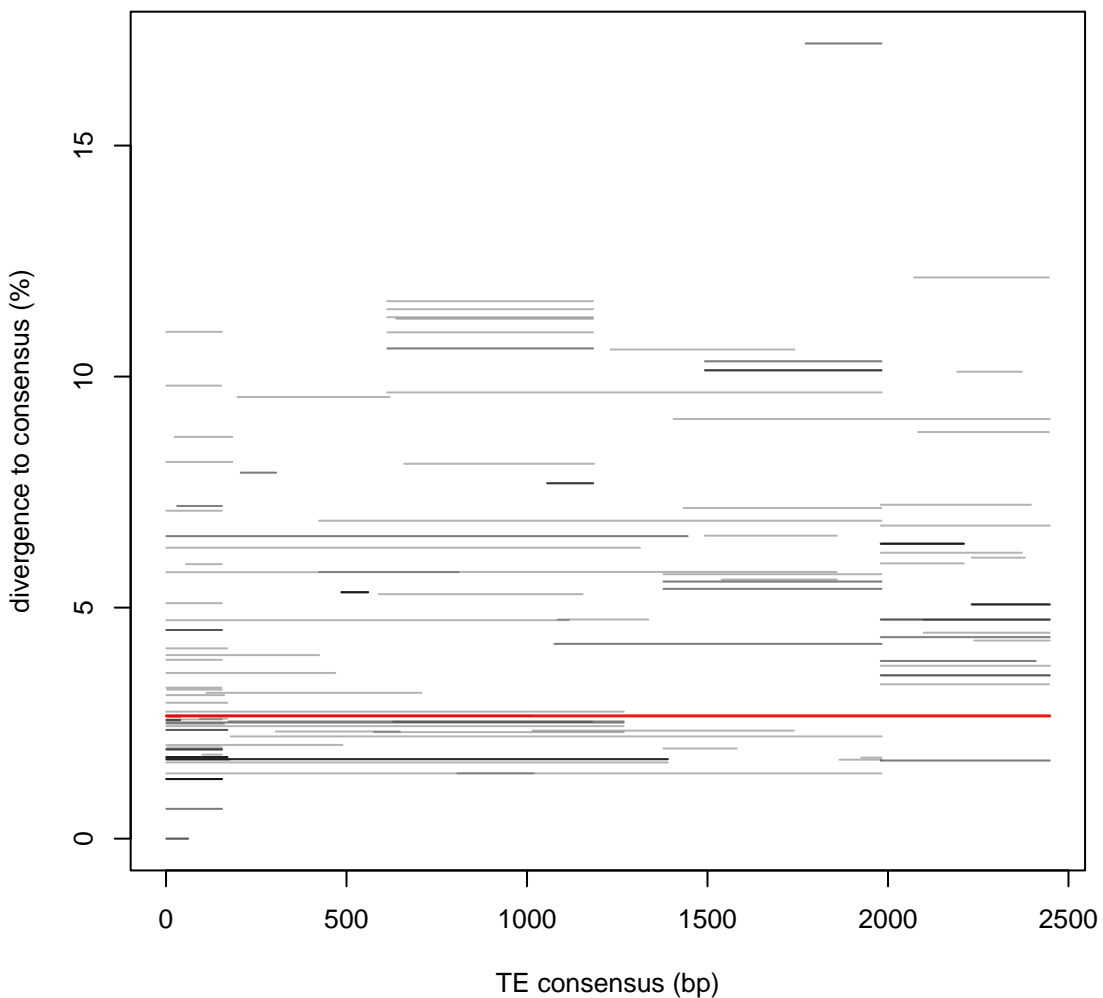
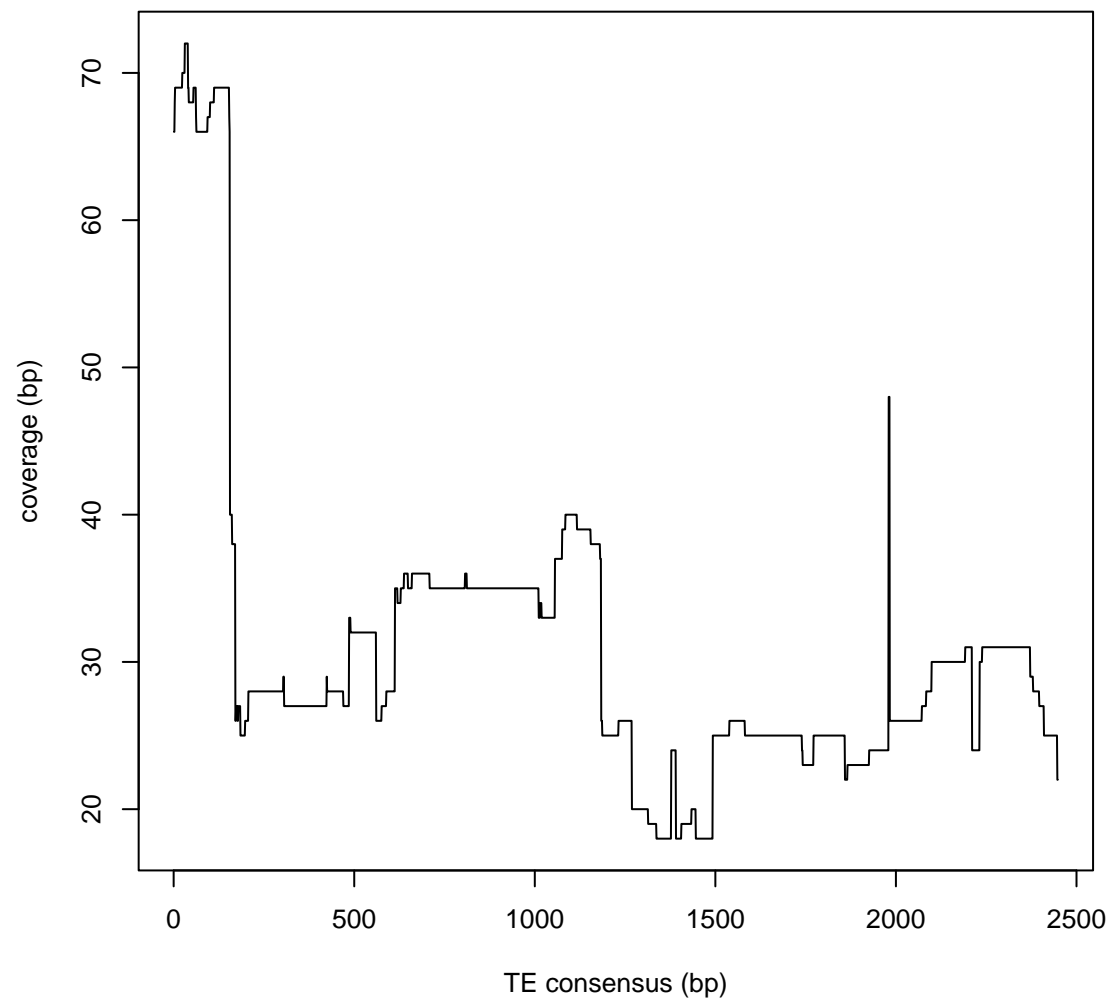


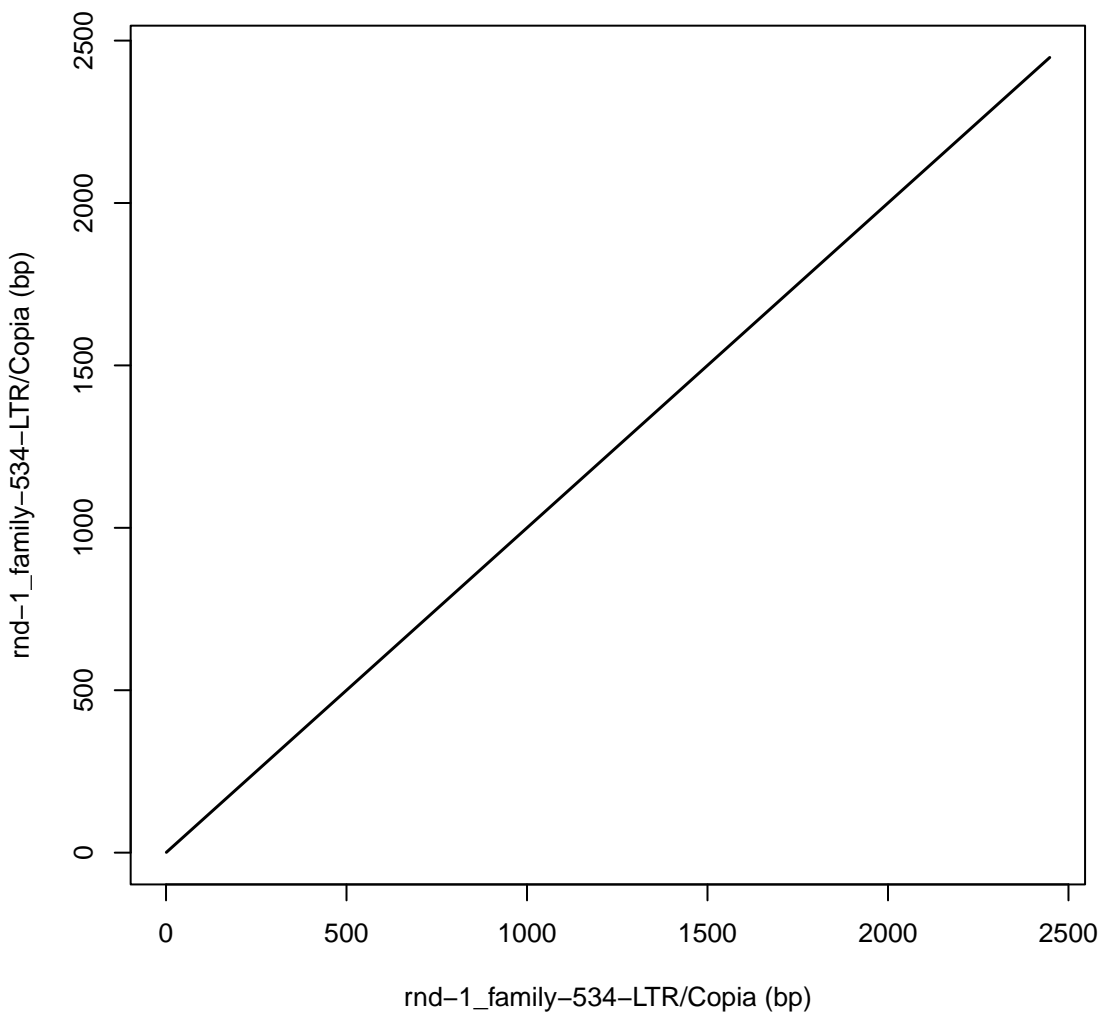
TE: rnd-1_family-534-LTR/Copia
consensus size: 2448bp; fragments: 168; full length: 1 (>=2203.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

