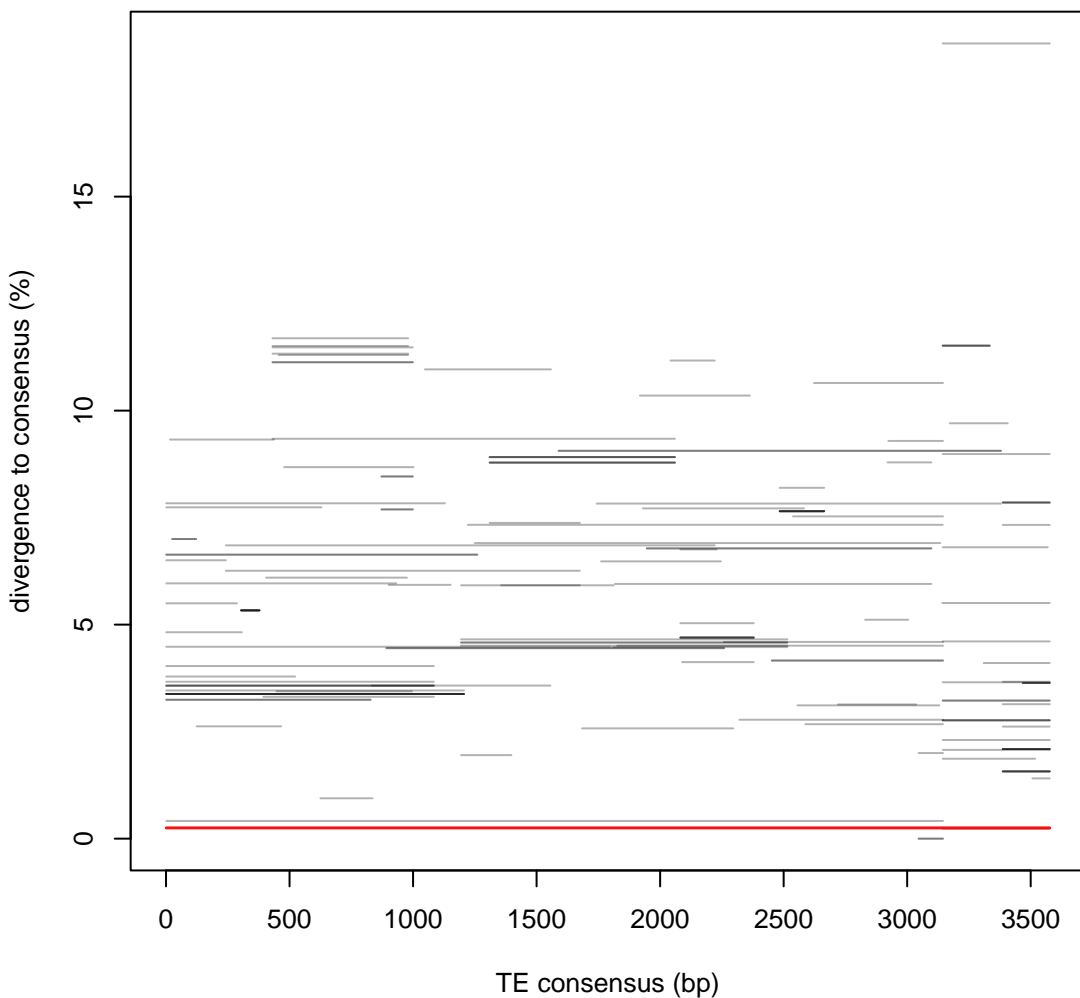
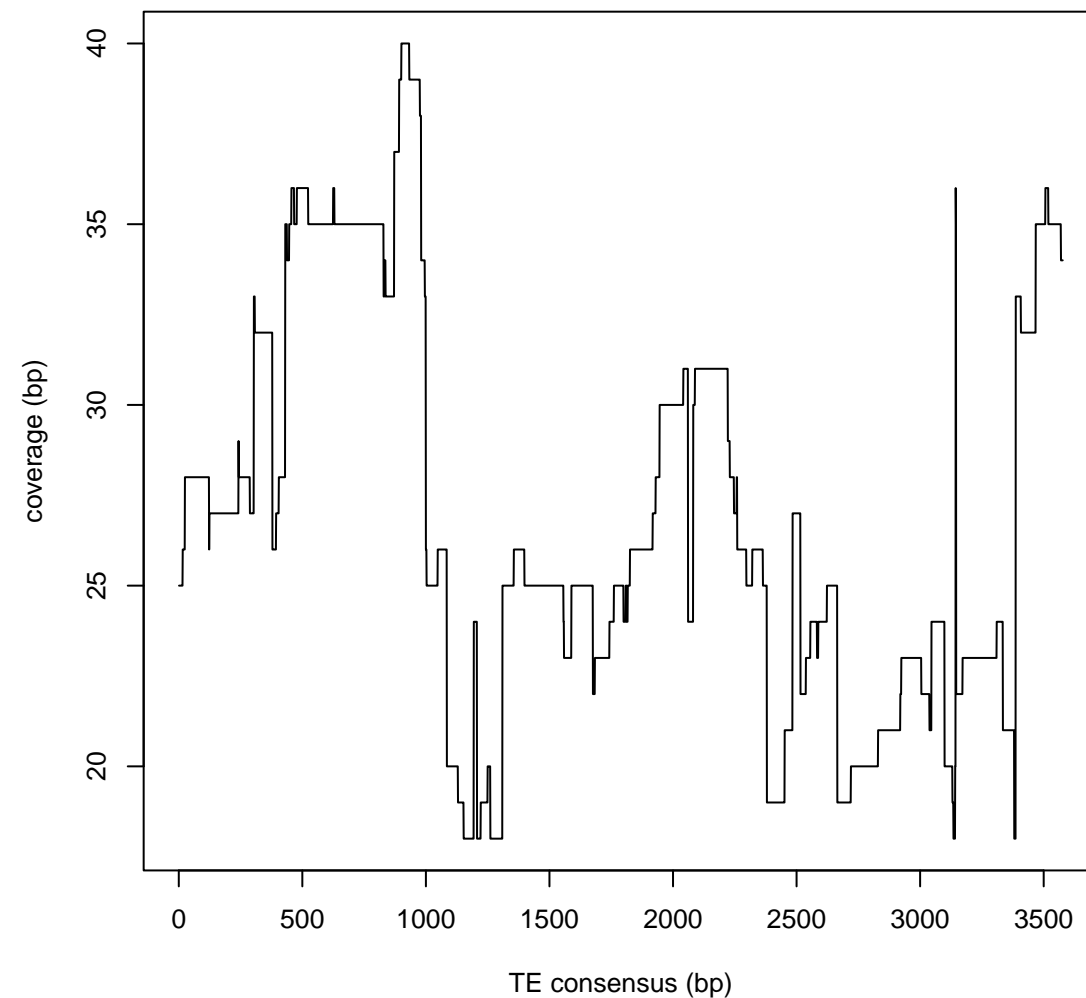


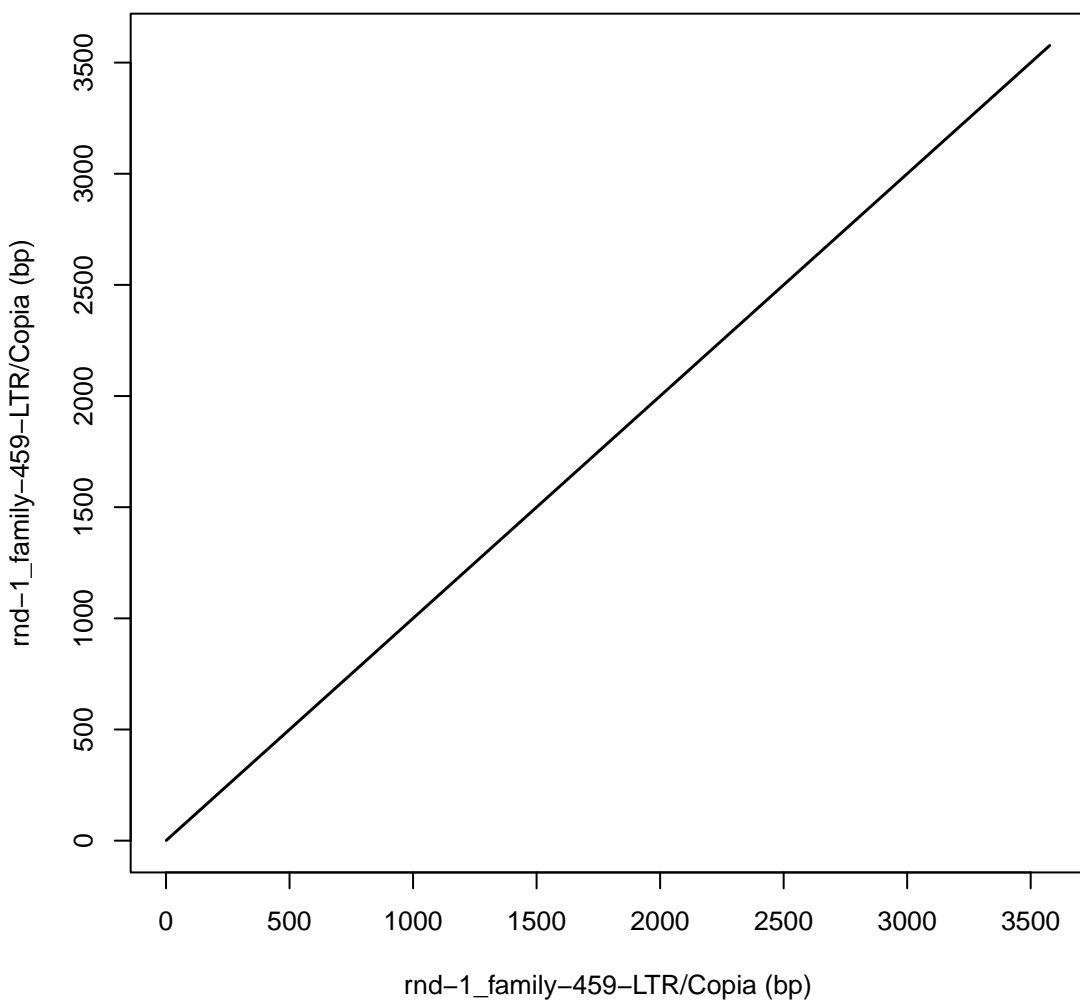
TE: rnd-1_family-459-LTR/Copia
consensus size: 3577bp; fragments: 155; full length: 1 (>=3219.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

