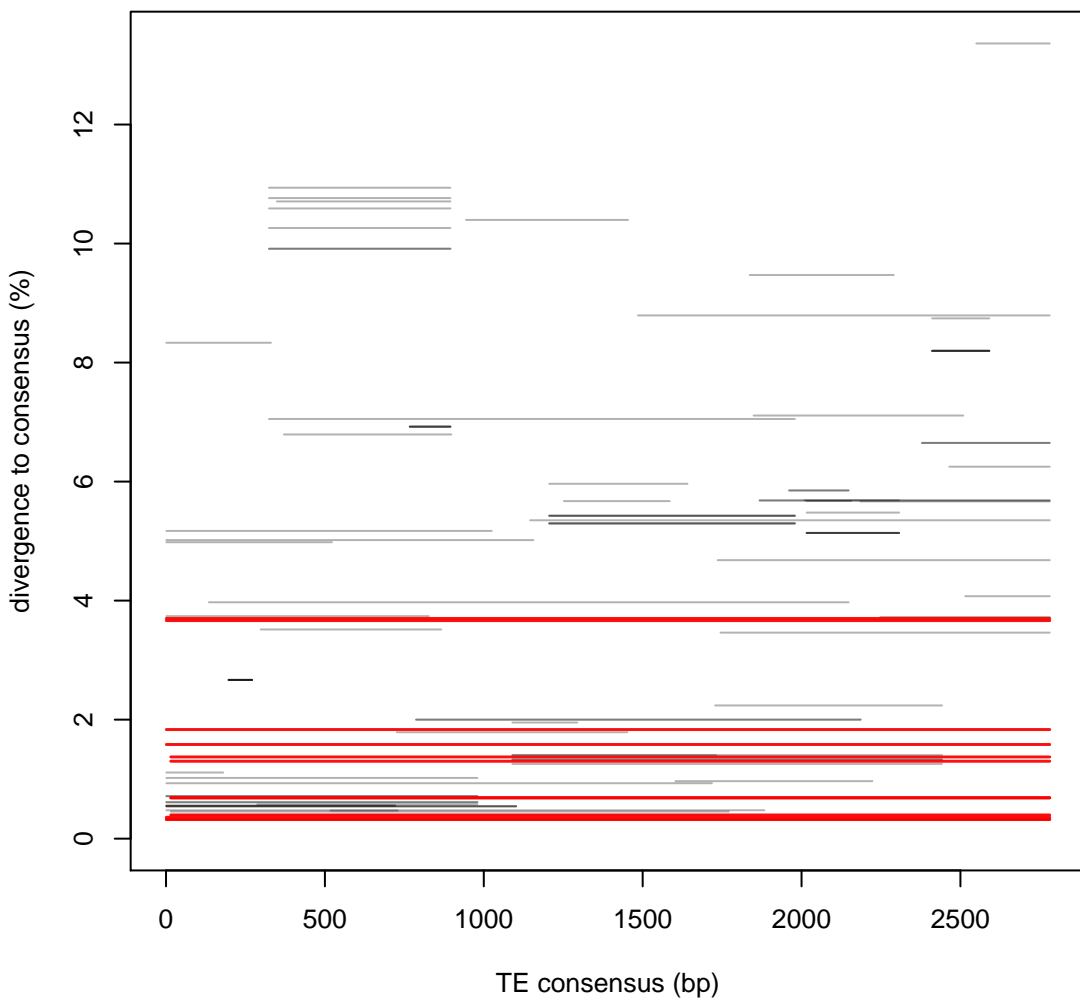
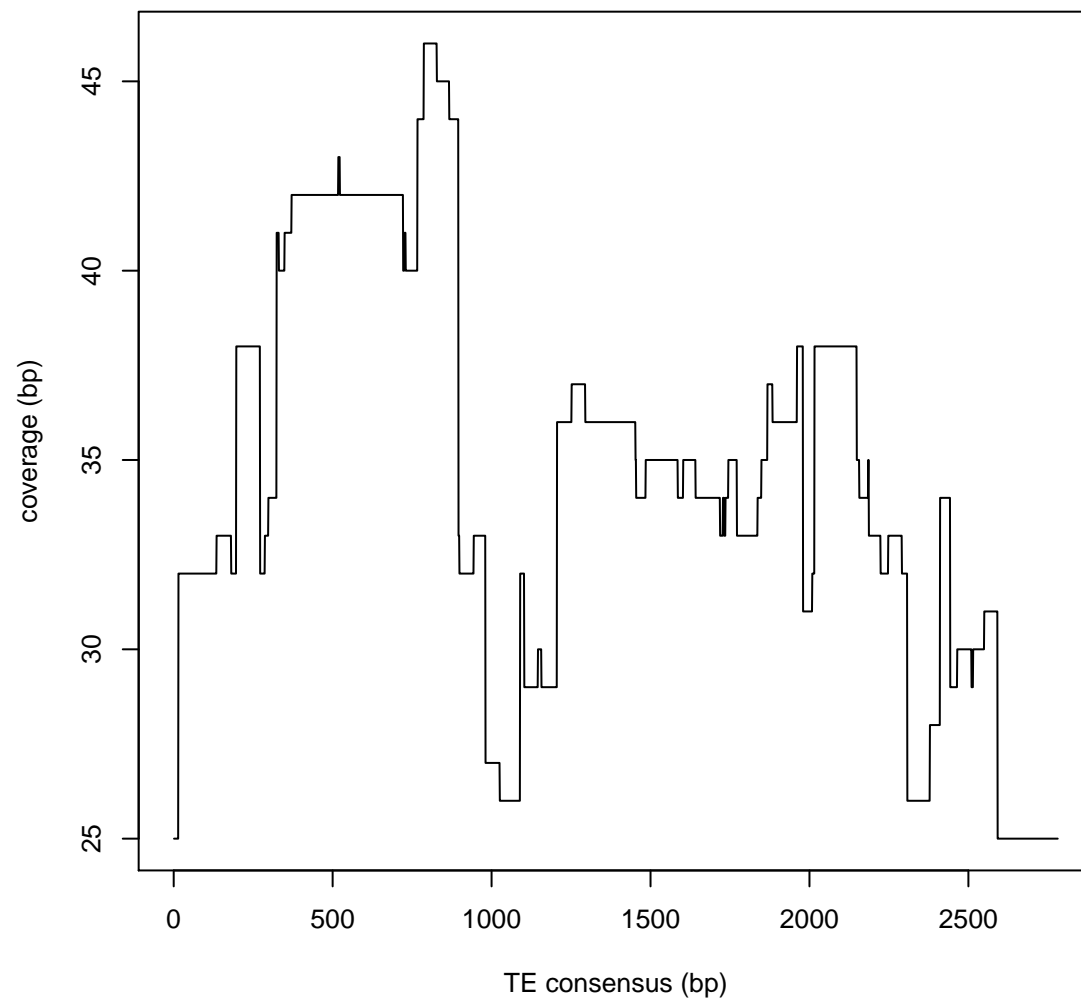


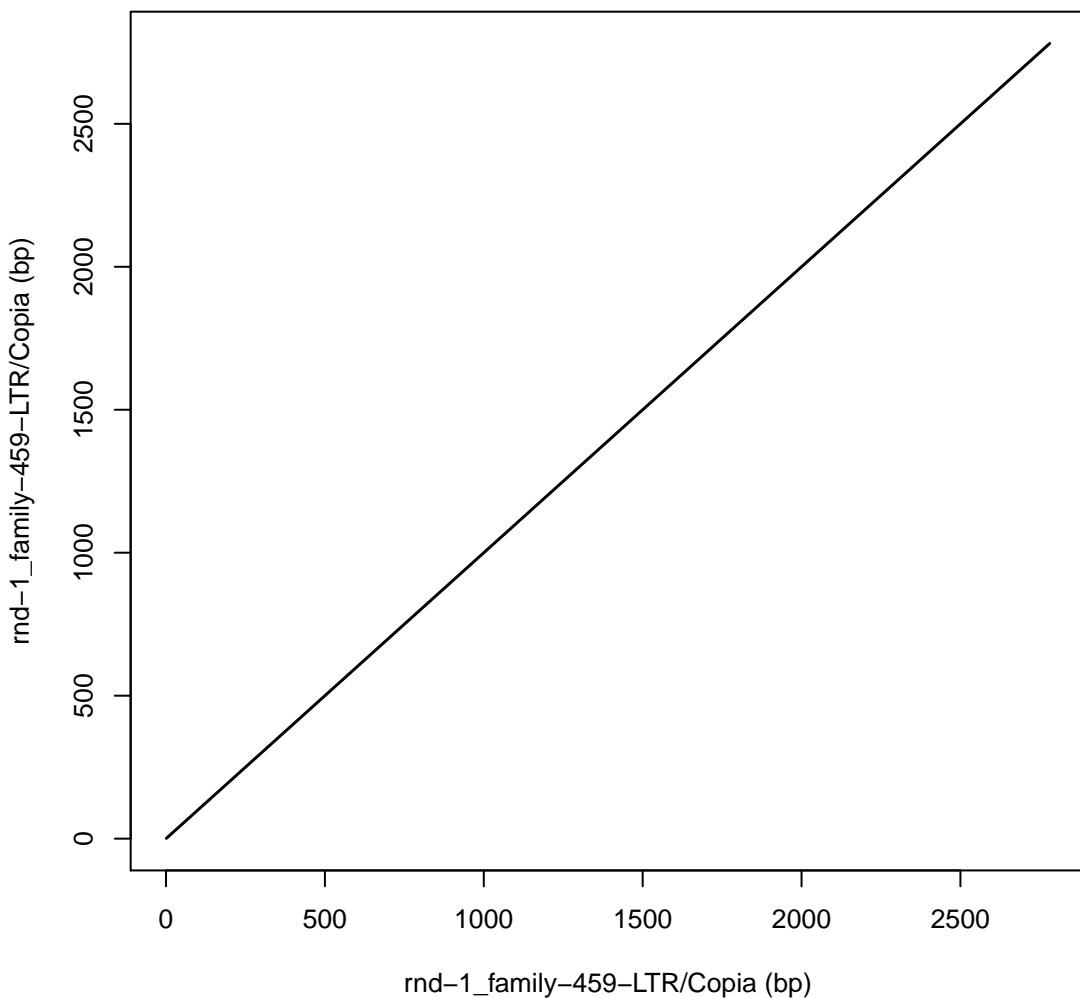
TE: rnd-1_family-459-LTR/Copia
consensus size: 2781bp; fragments: 105; full length: 12 (>=2502.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

