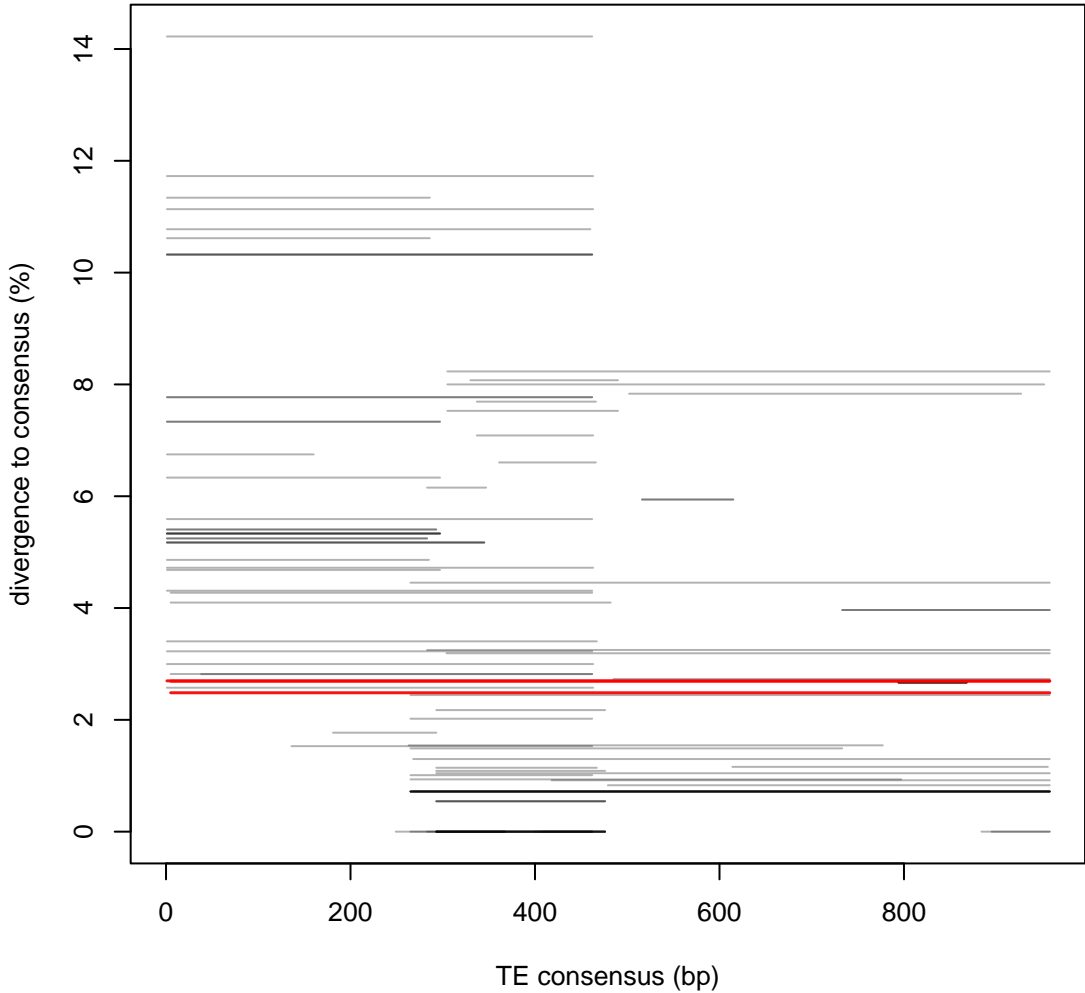
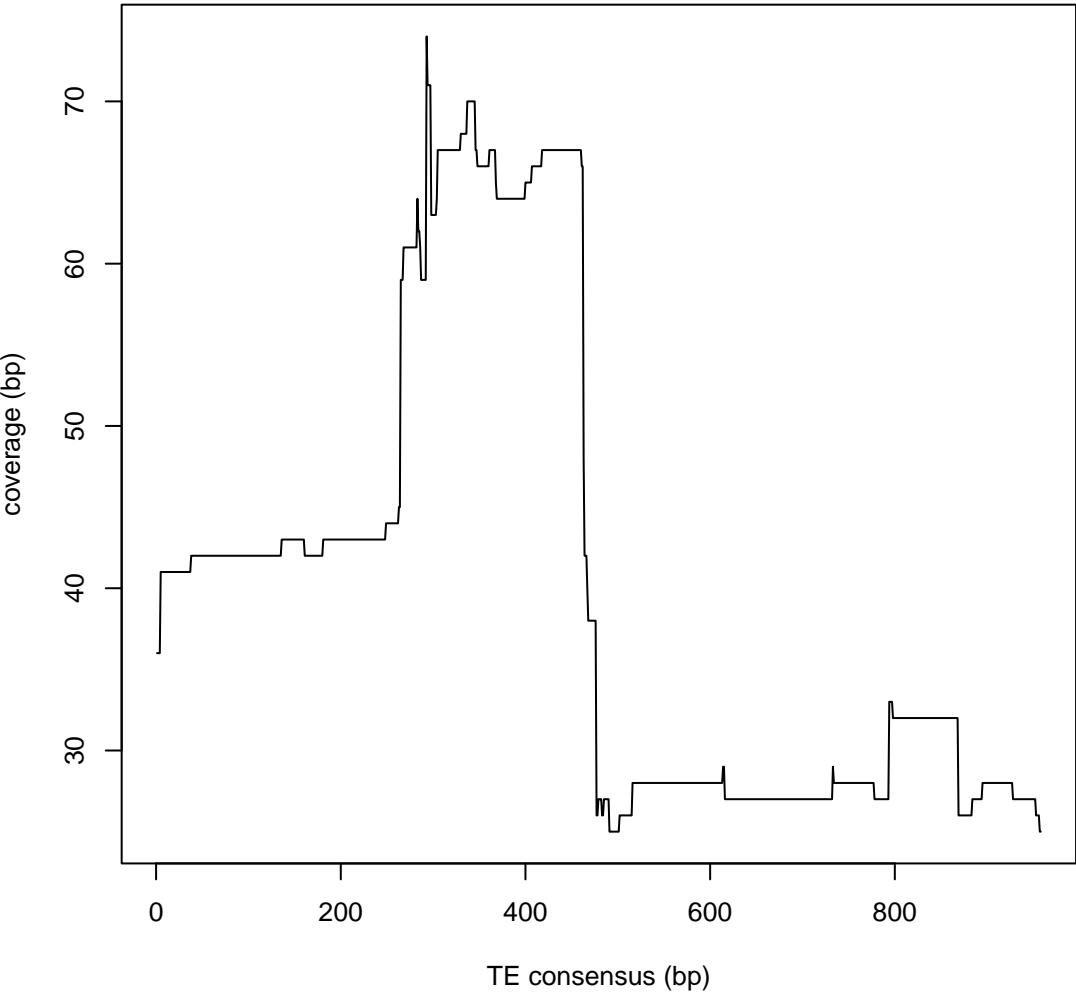


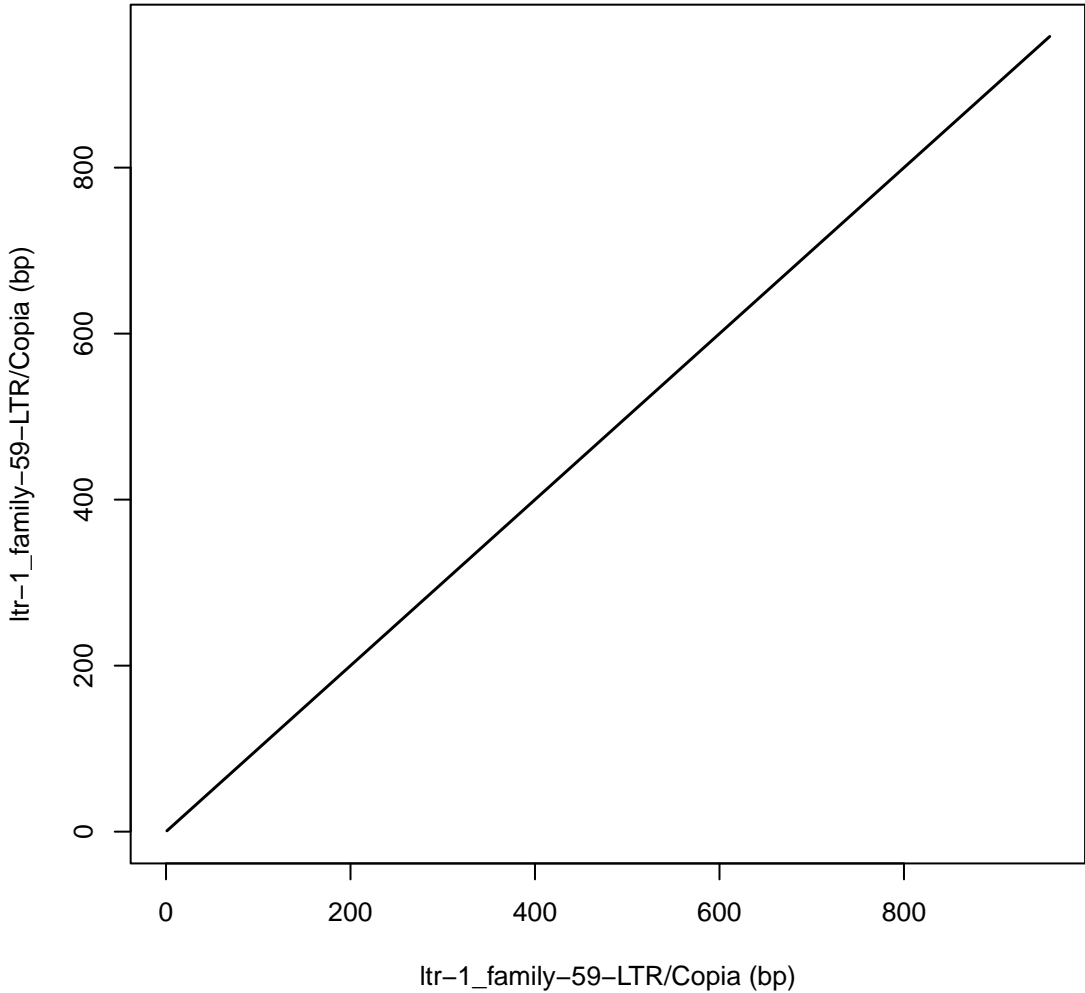
TE: ltr-1_family-59-LTR/Copia
consensus size: 958bp; fragments: 107; full length: 3 (>=862.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

