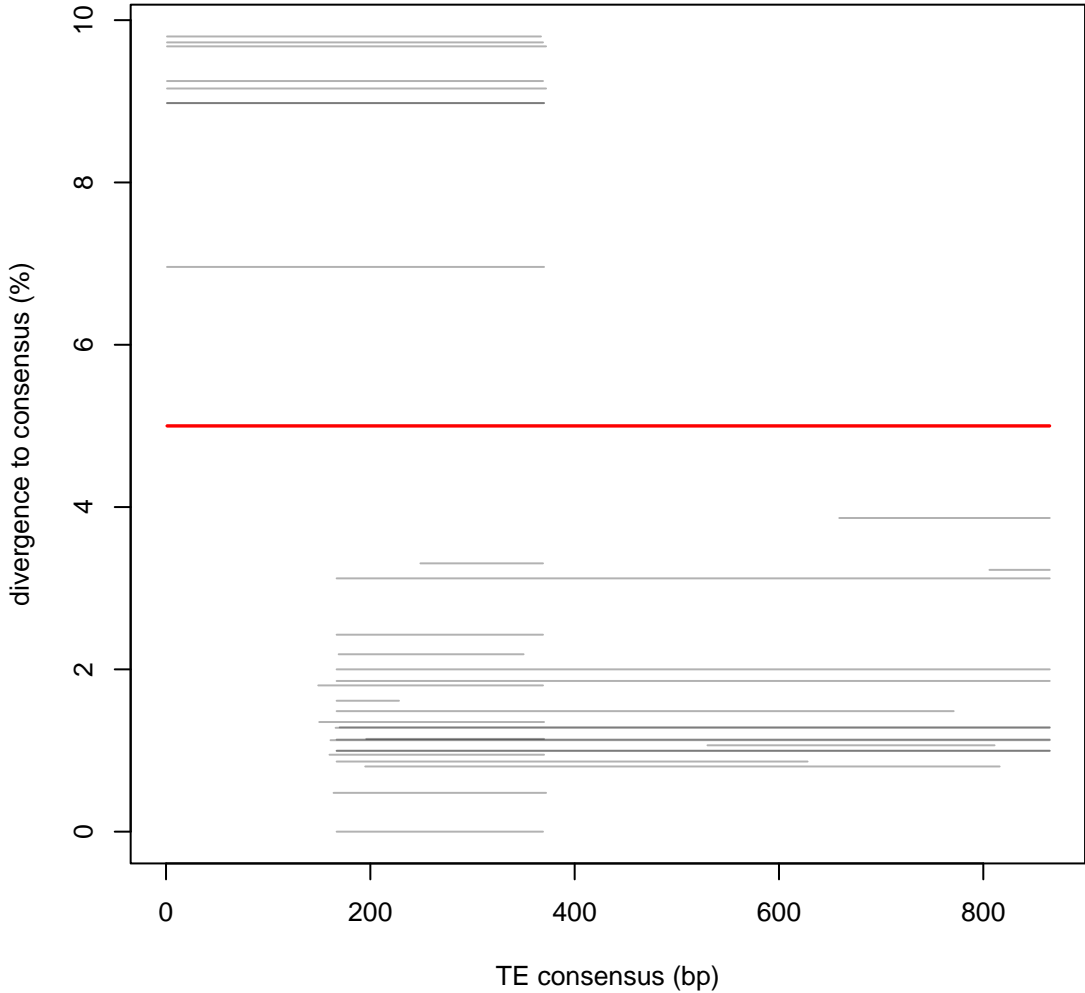
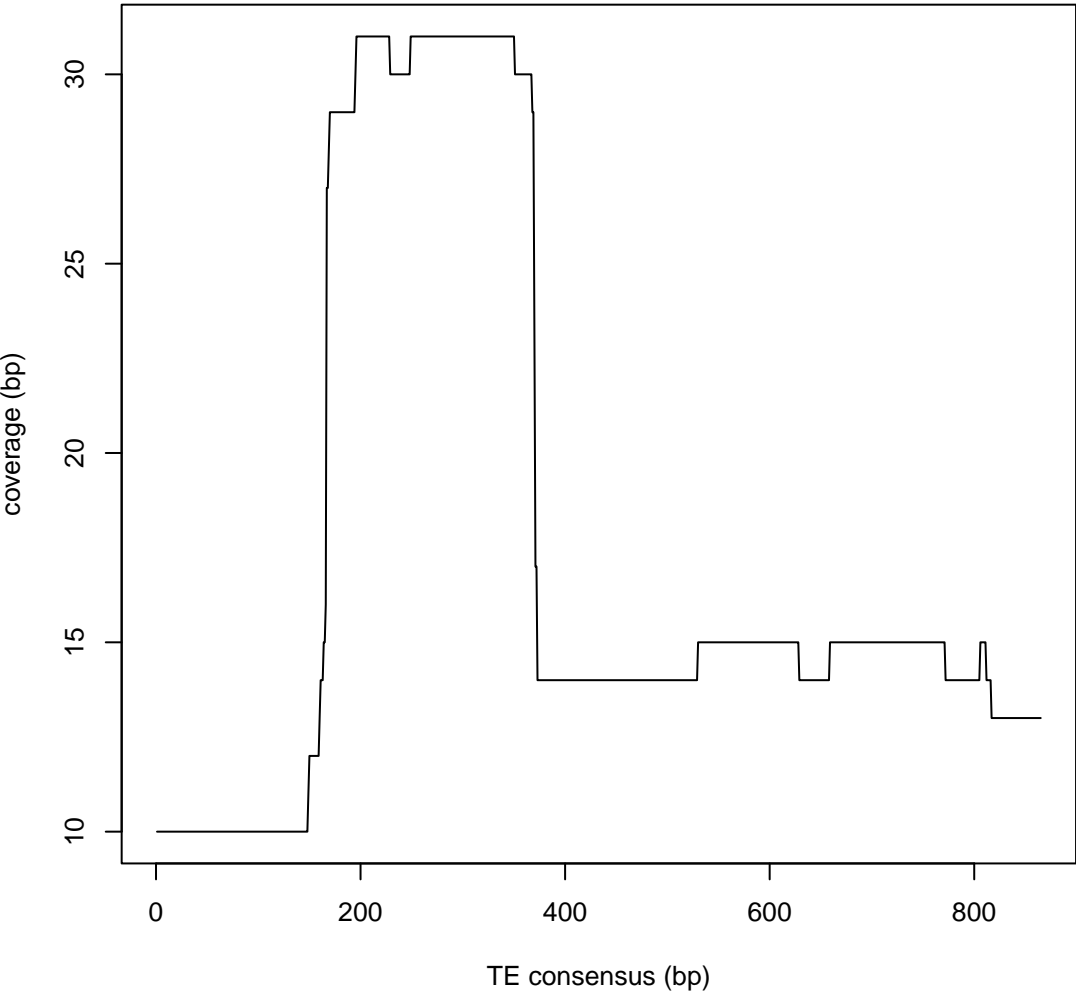


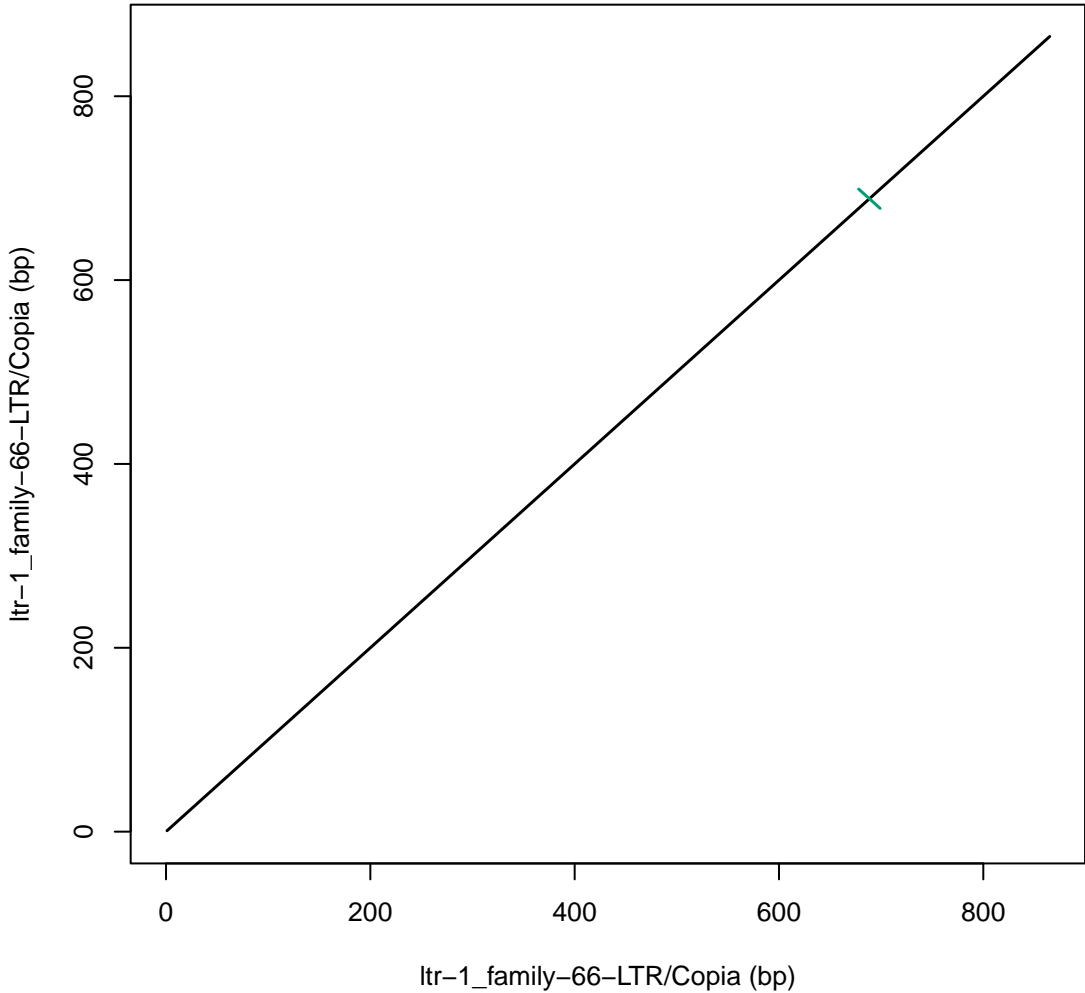
TE: ltr-1_family-66-LTR/Copia
consensus size: 865bp; fragments: 35; full length: 2 (>=778.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

