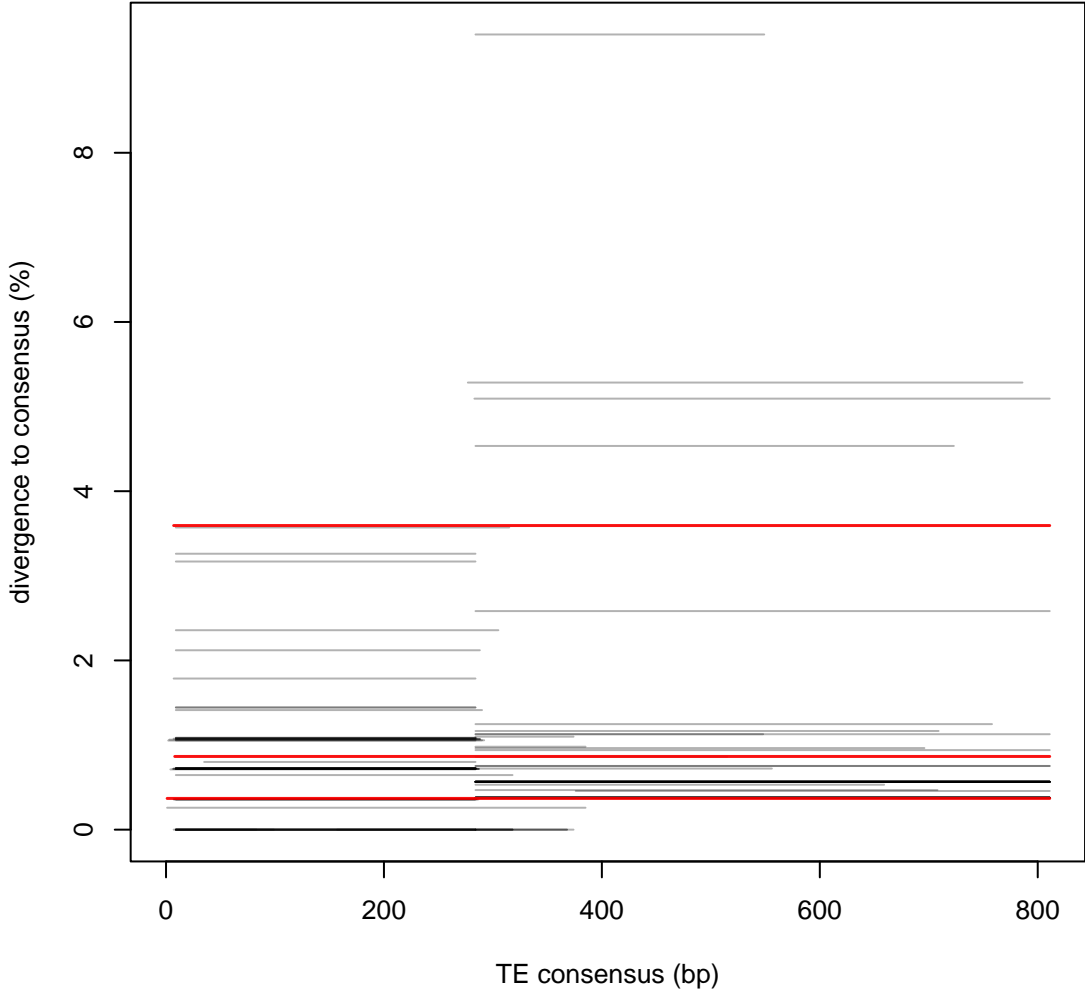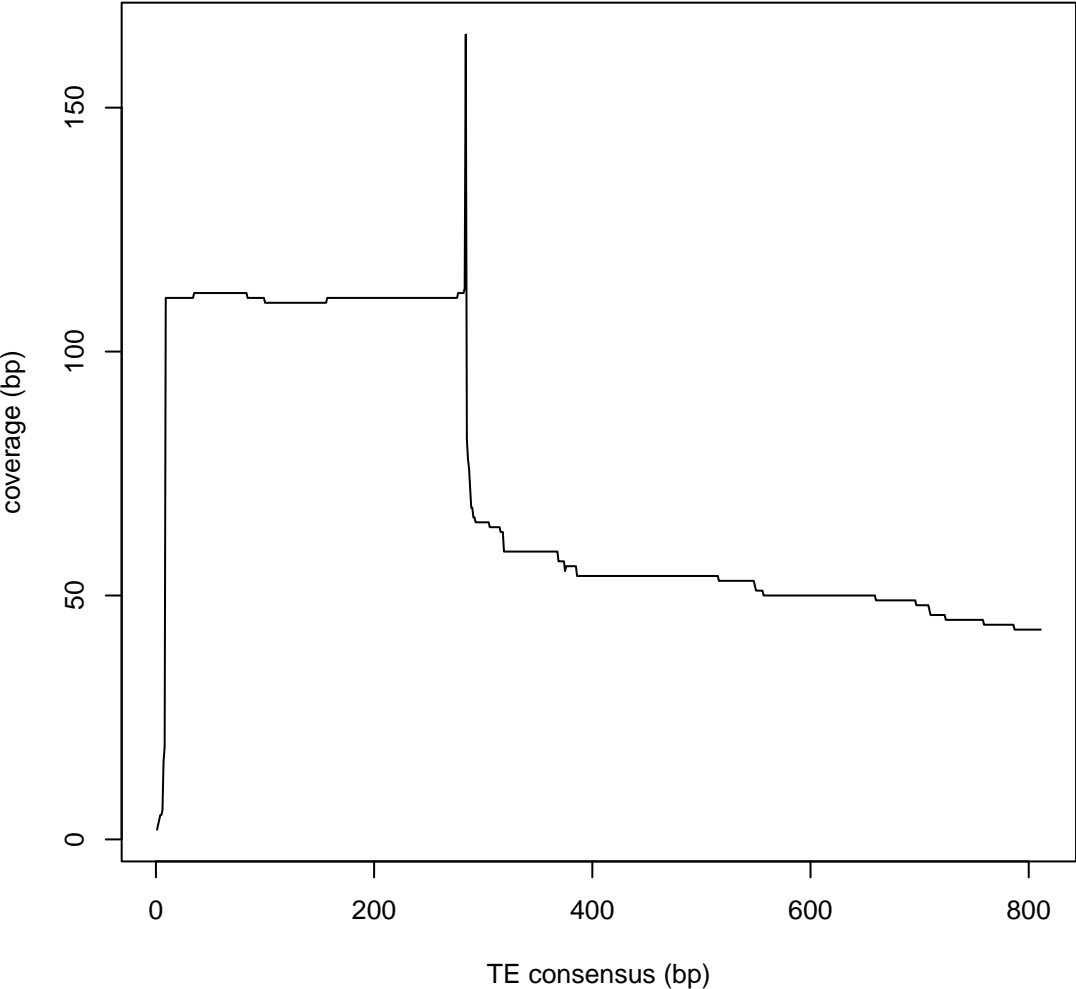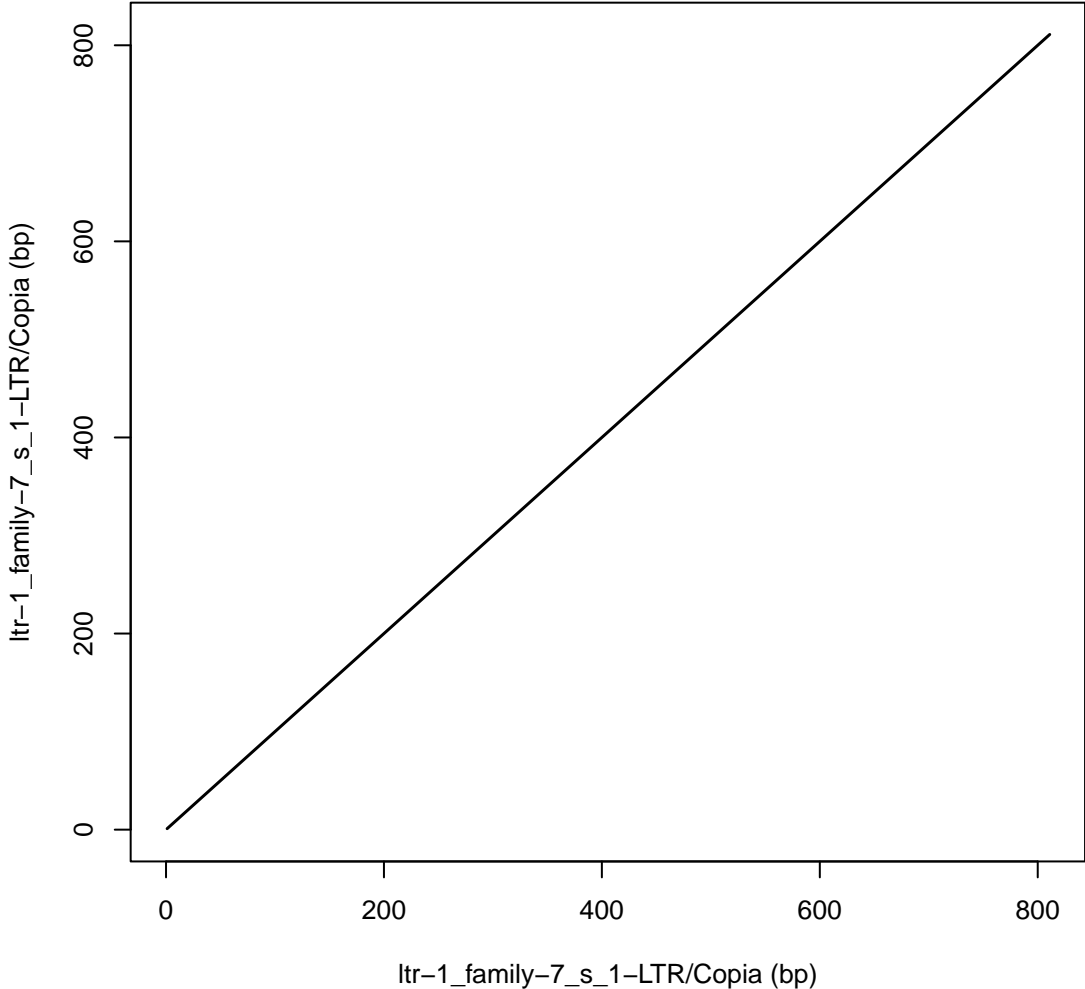**TE: ltr−1_family−7_s_1−LTR/Copia**
consensus size: 811bp; fragments: 168; full length: 3 (>=729.9bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected