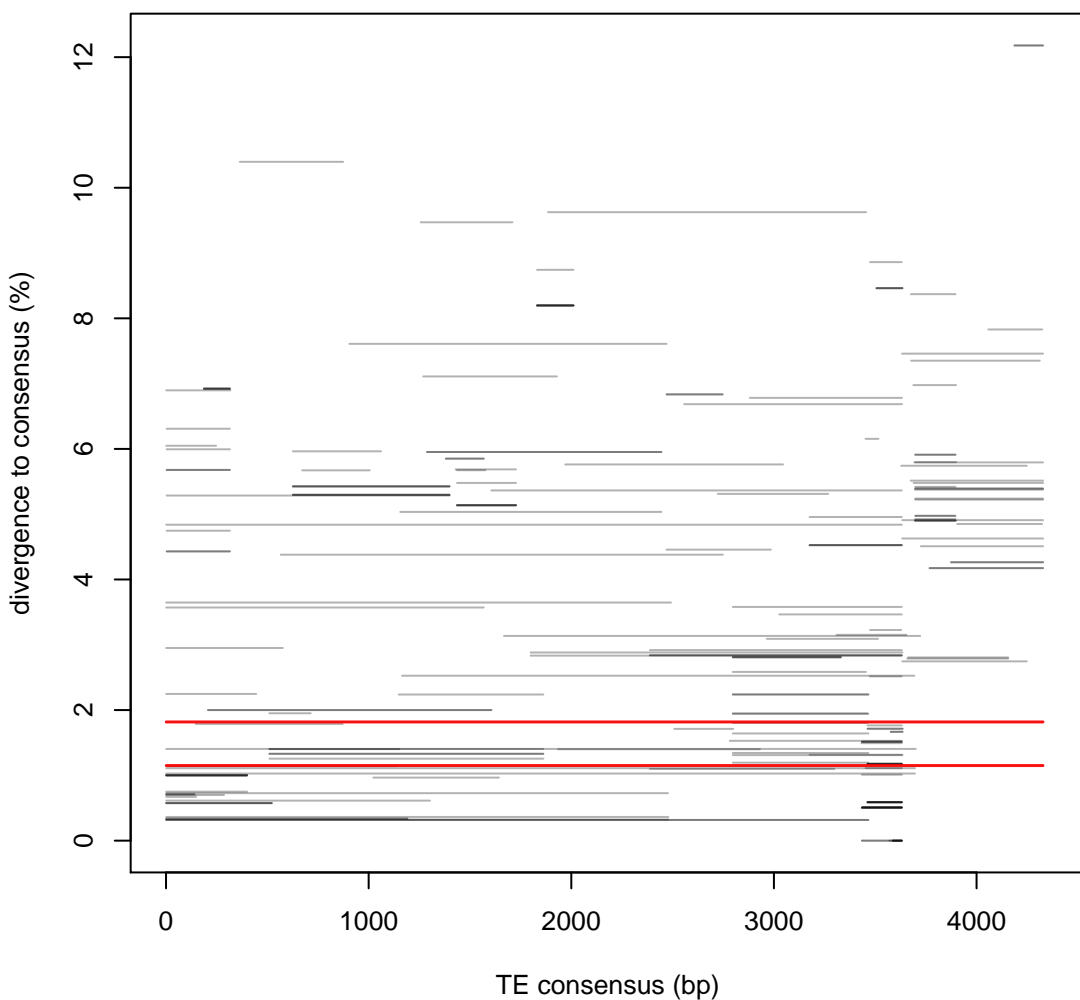
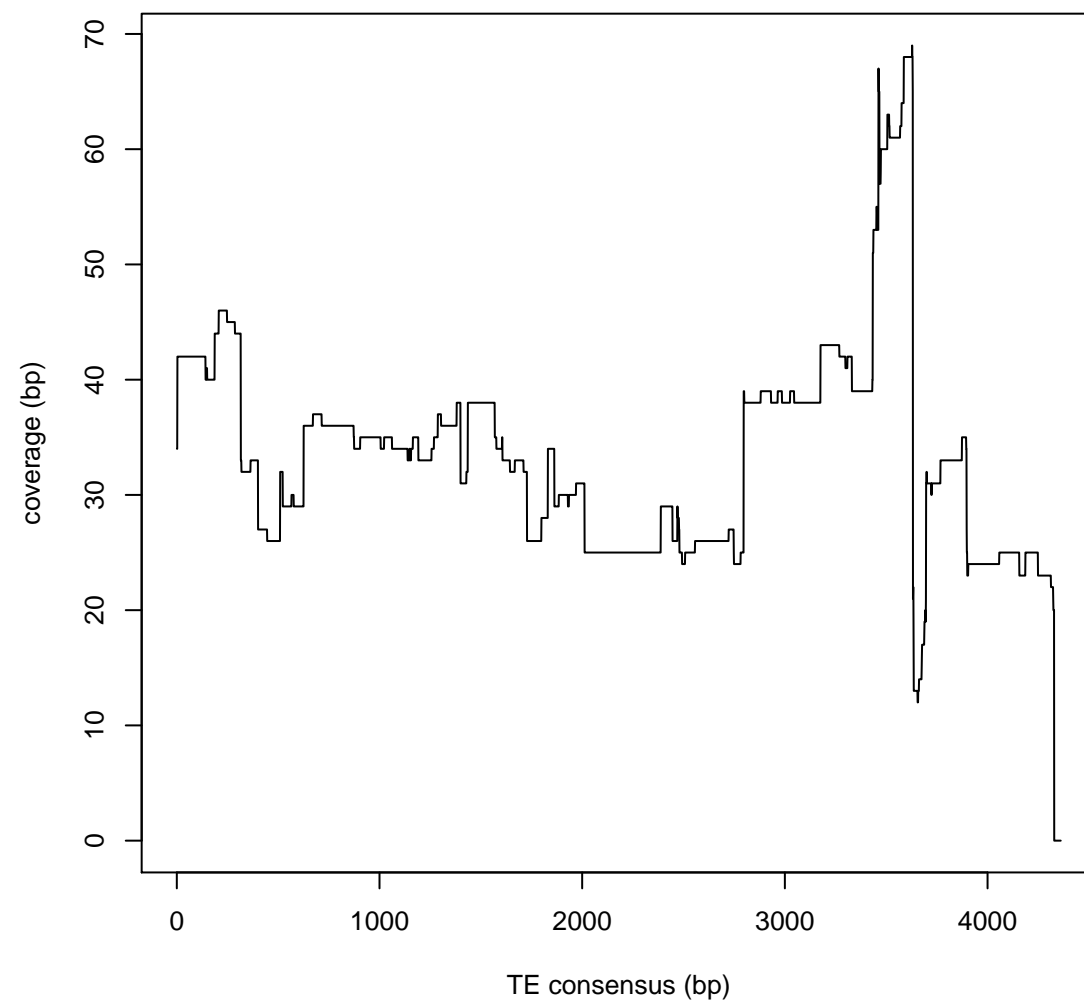


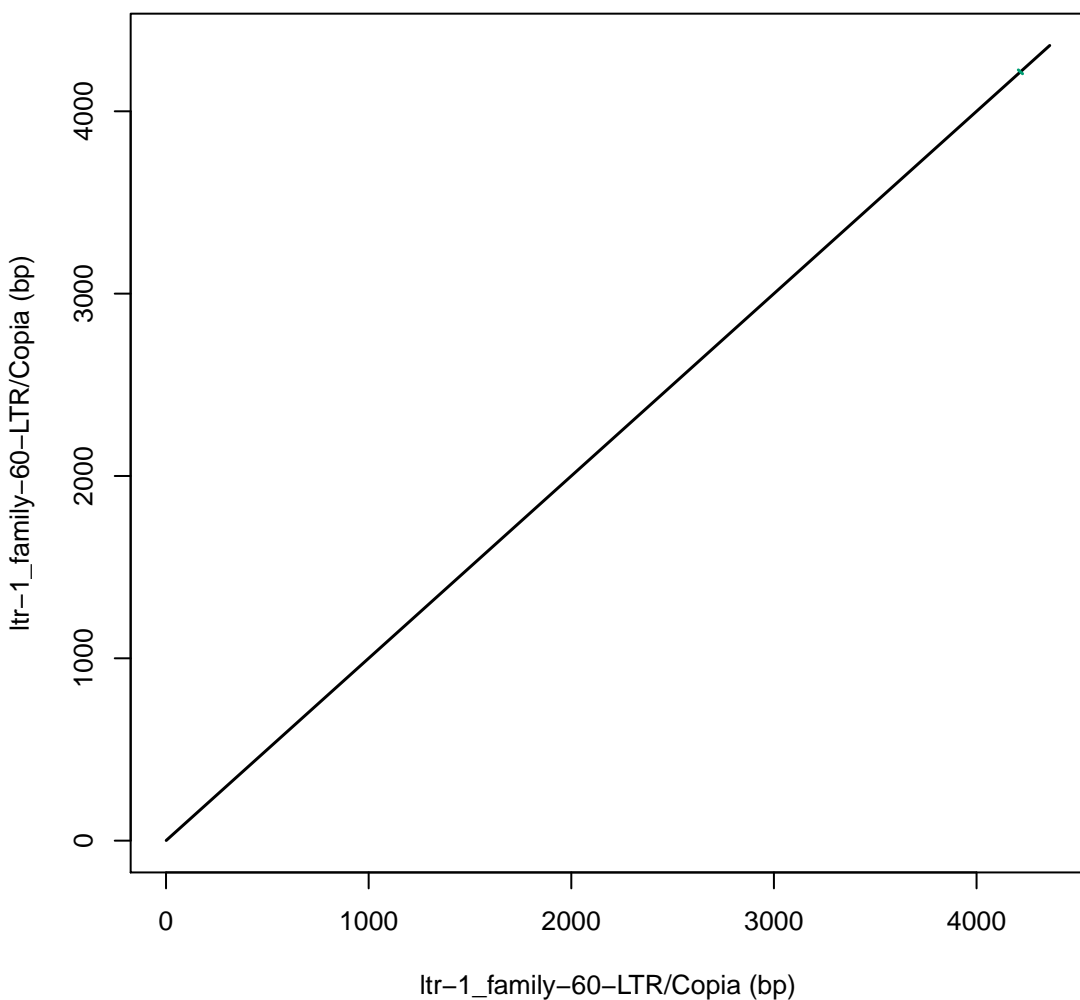
TE: ltr-1_family-60-LTR/Copia
consensus size: 4361bp; fragments: 210; full length: 2 (≥ 3924.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

