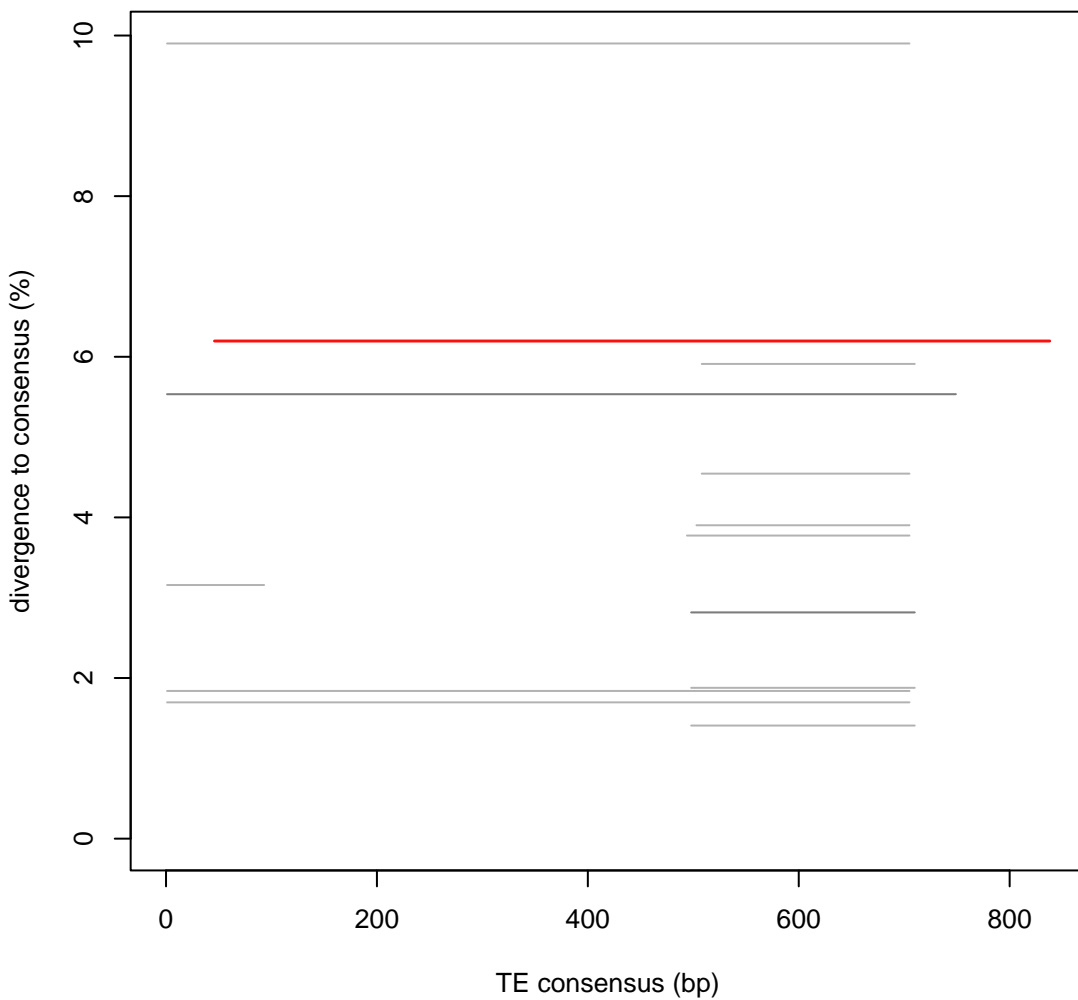
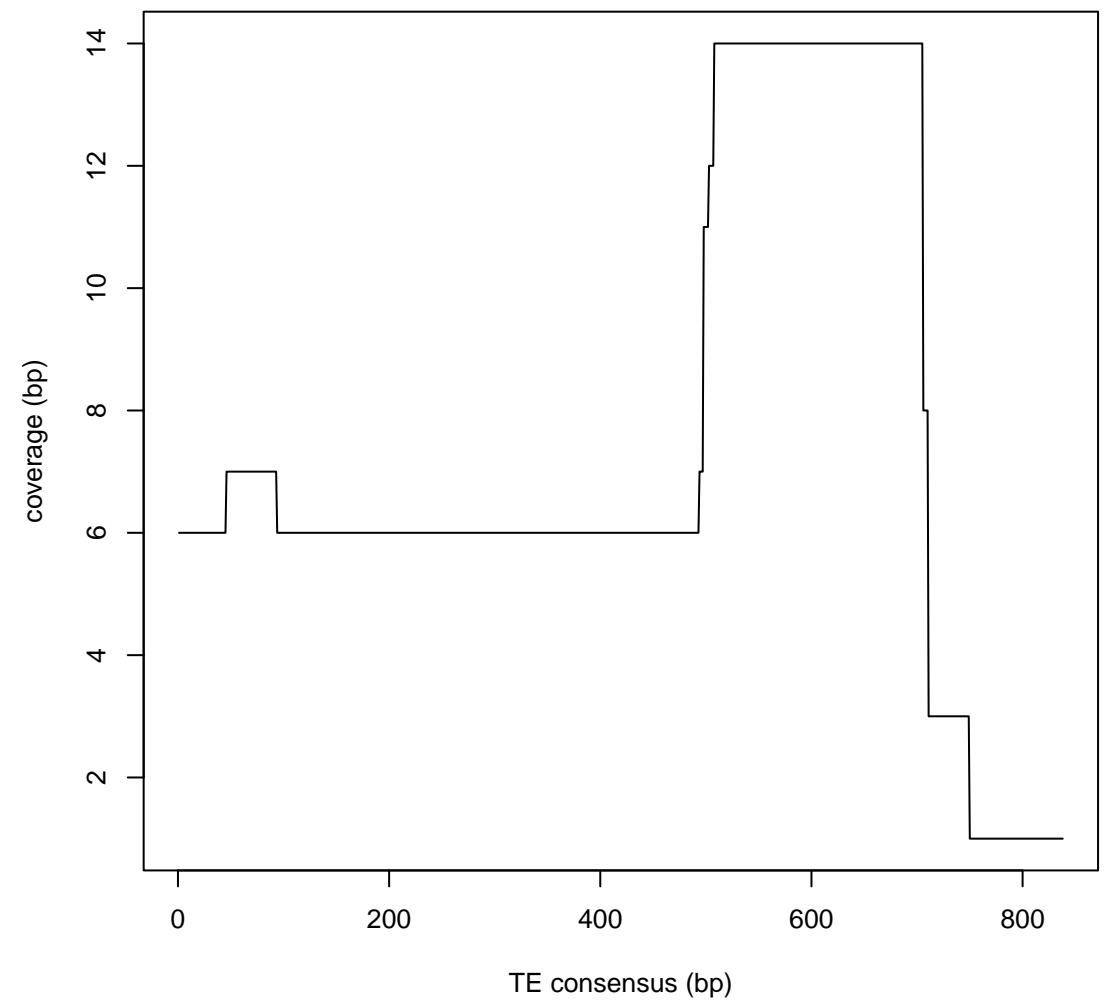


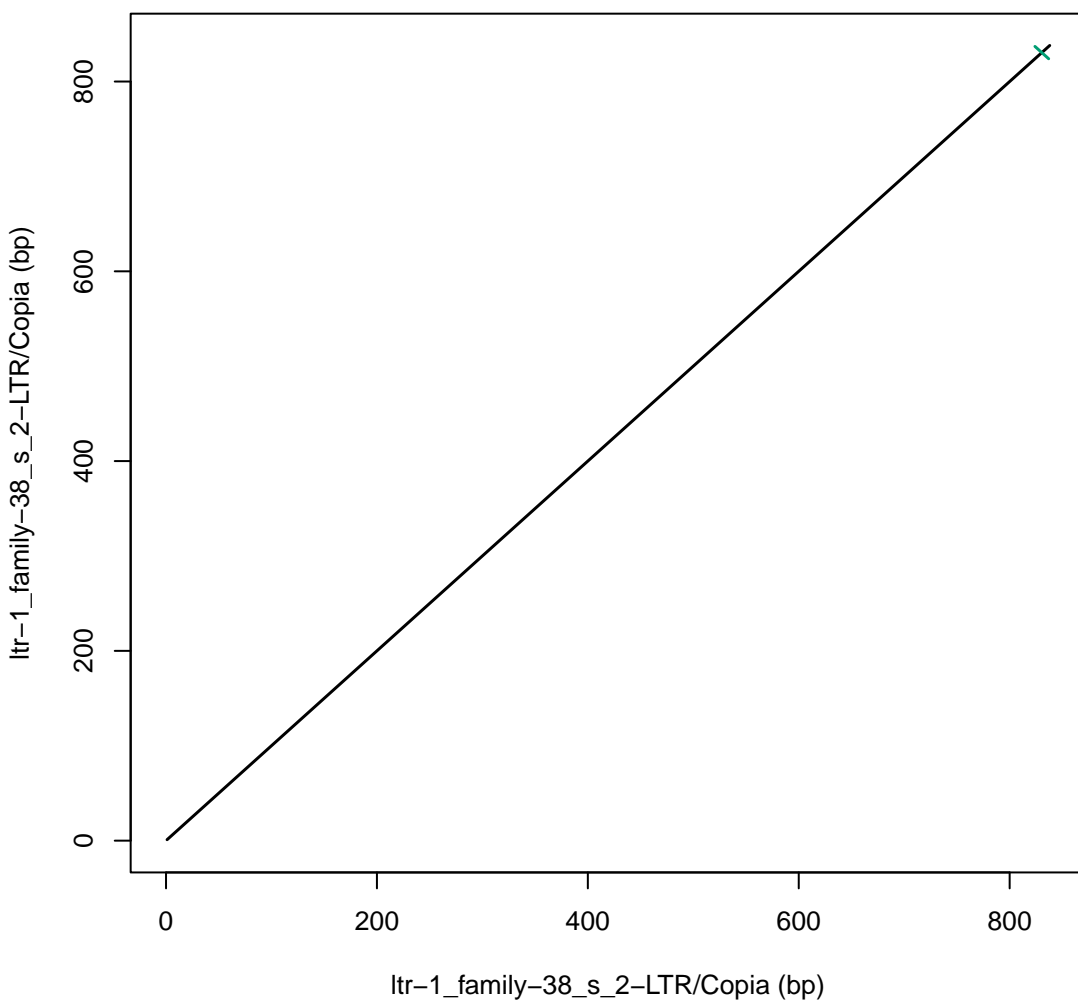
TE: ltr-1_family-38_s_2-LTR/Copia
consensus size: 838bp; fragments: 15; full length: 1 (>=754.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

