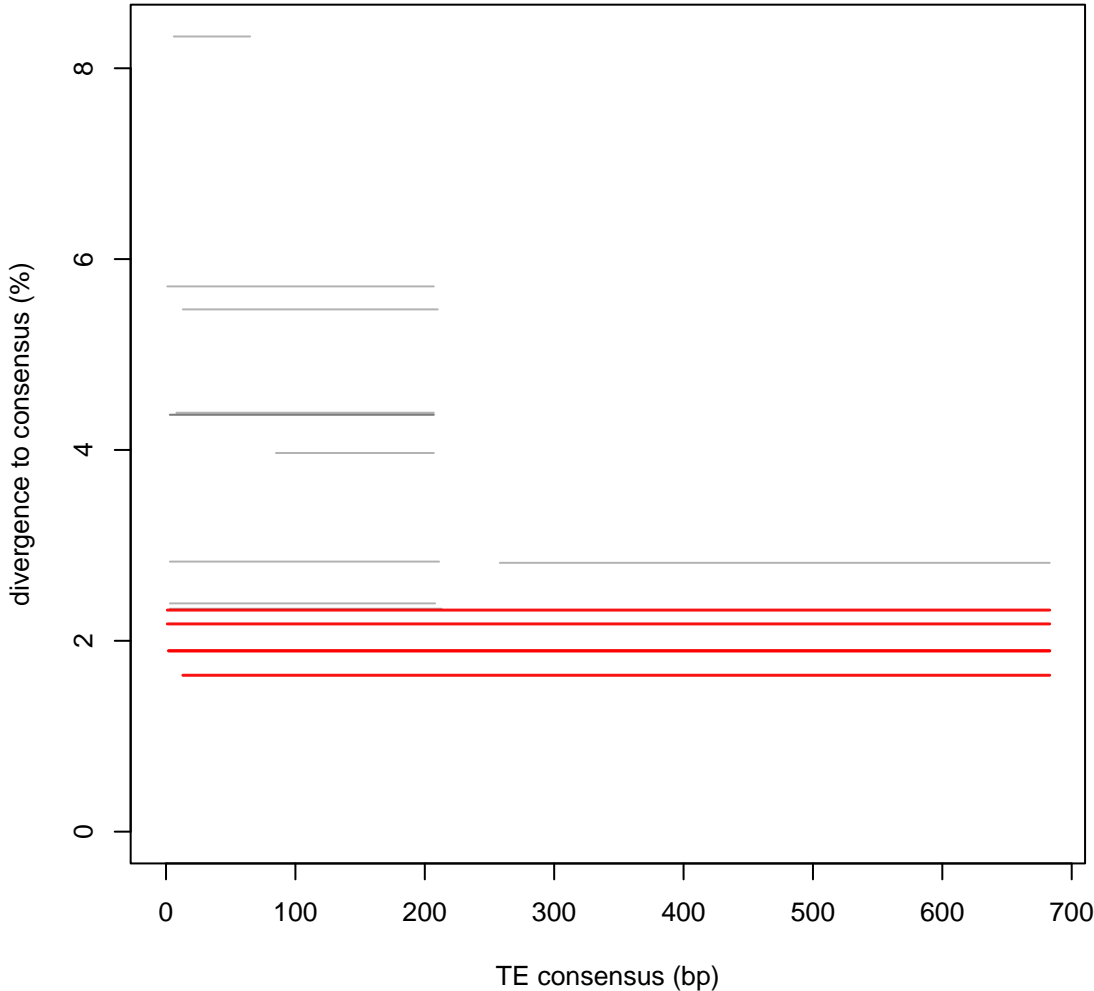
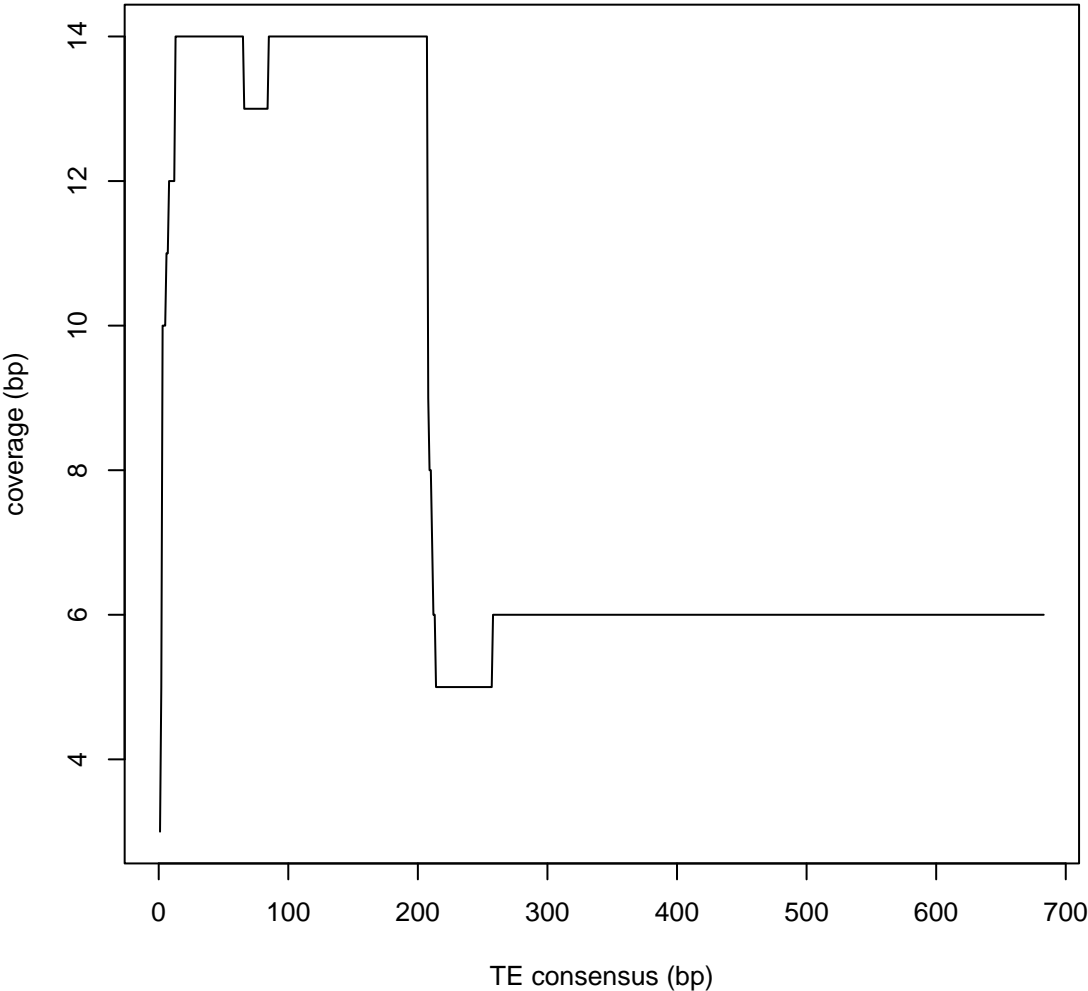


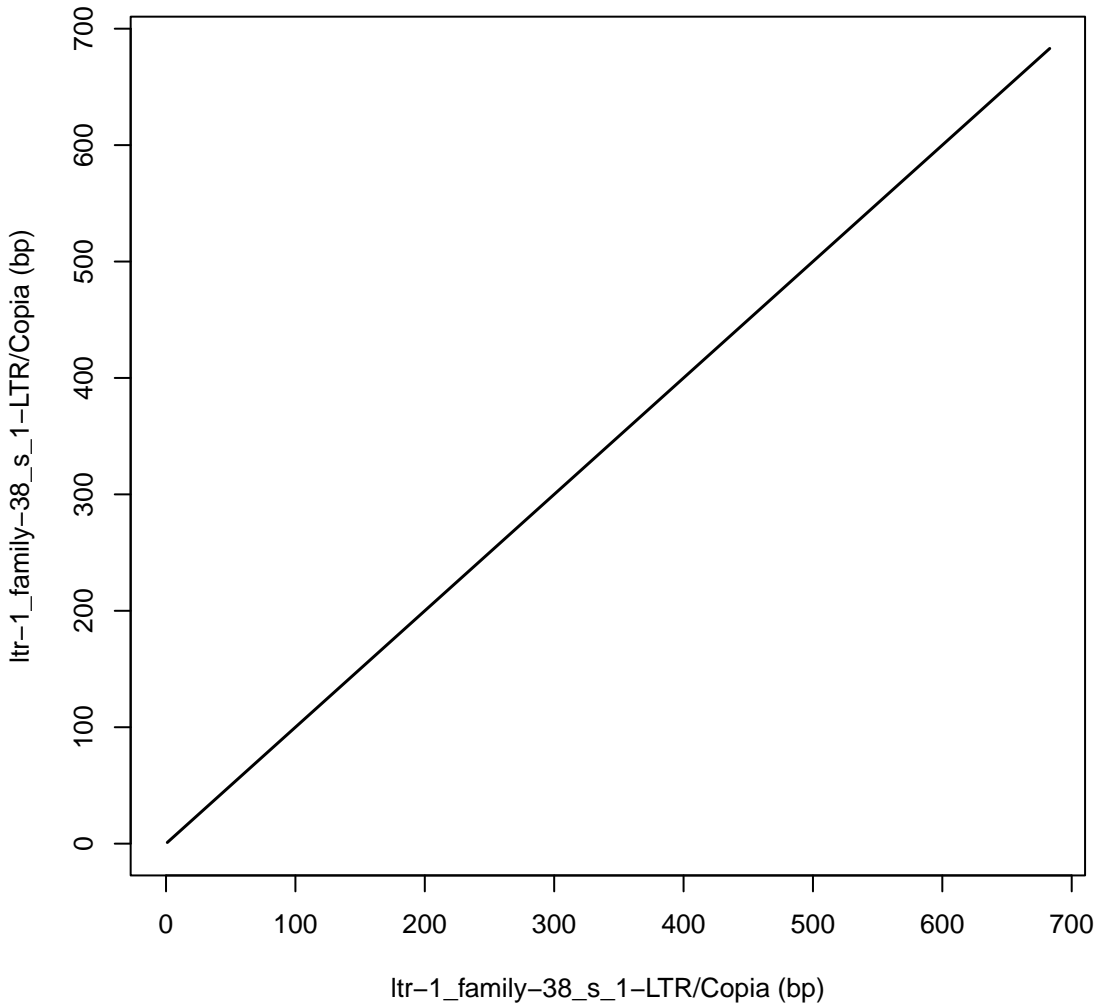
TE: ltr-1_family-38_s_1-LTR/Copia
consensus size: 683bp; fragments: 16; full length: 5 (>=614.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

