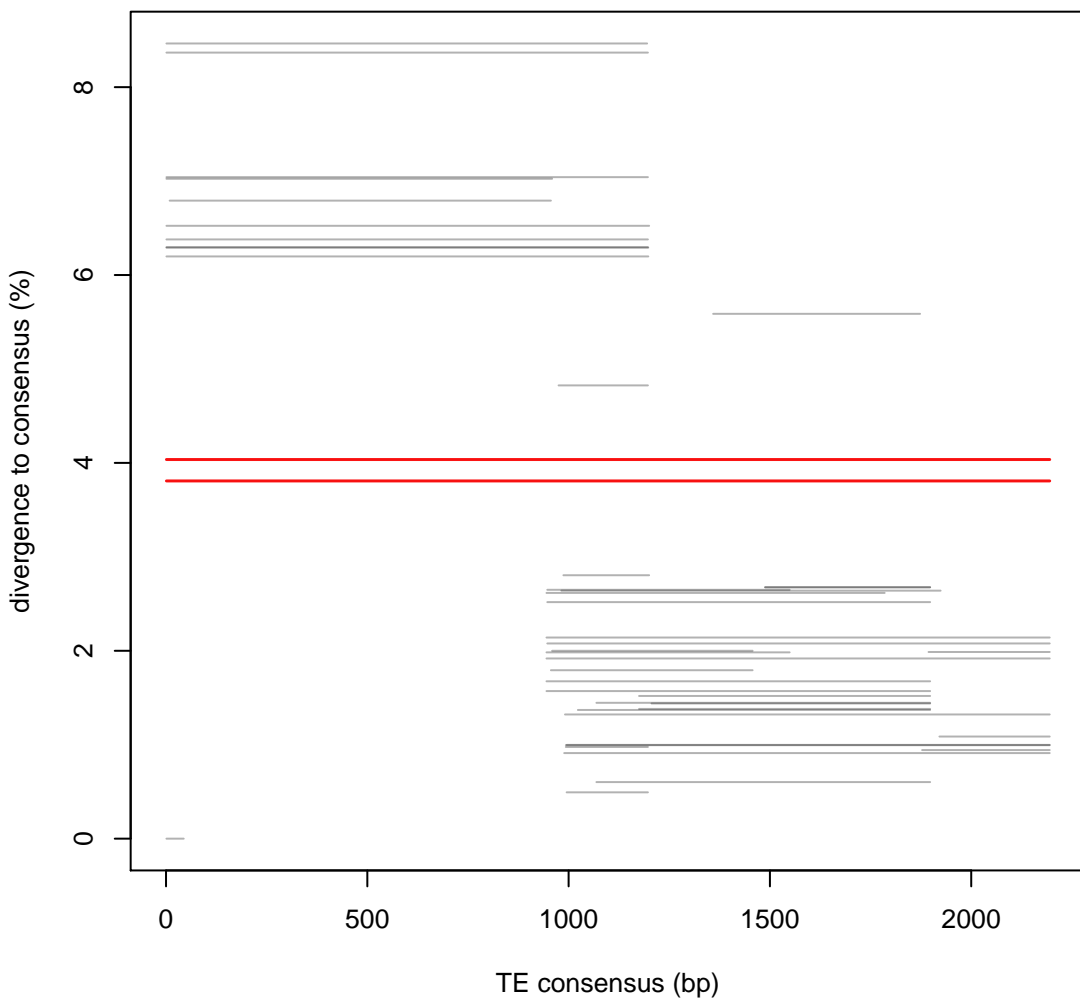
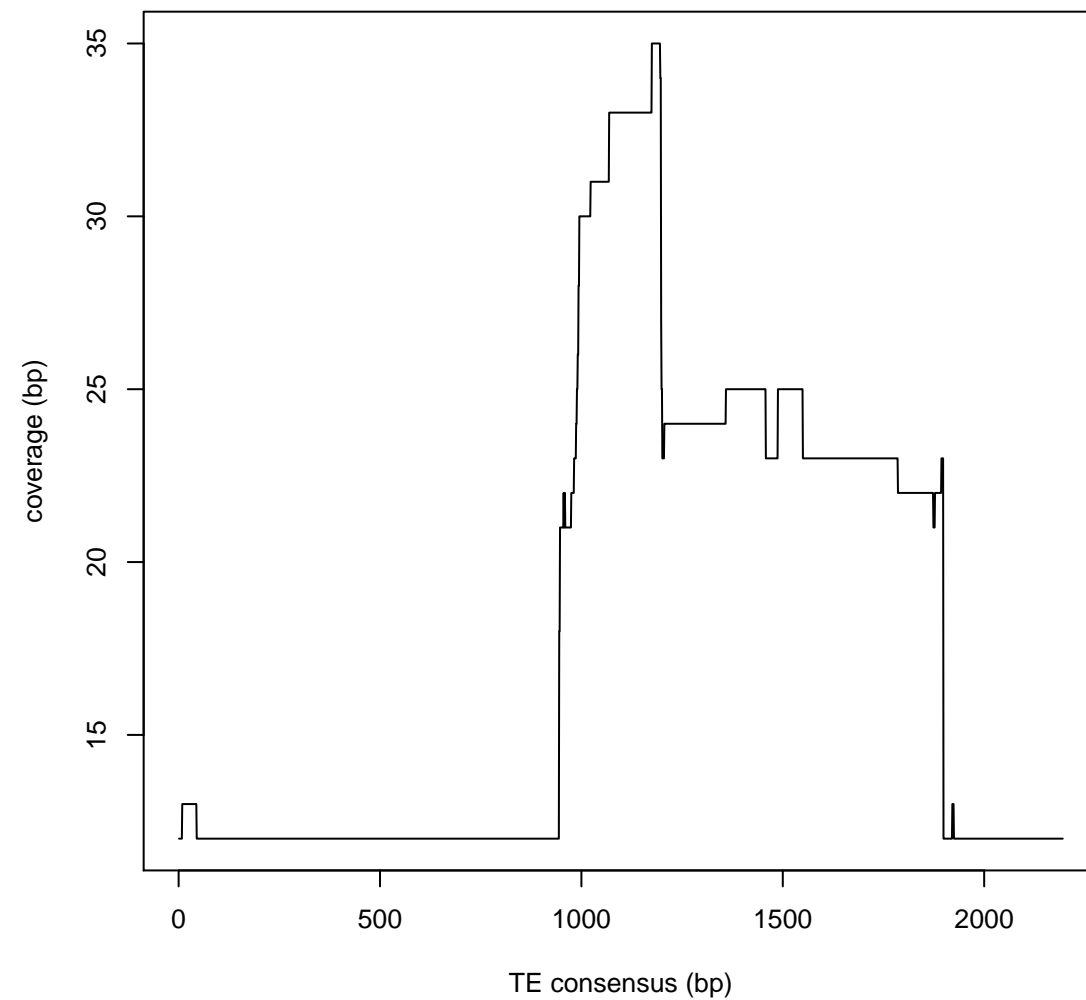


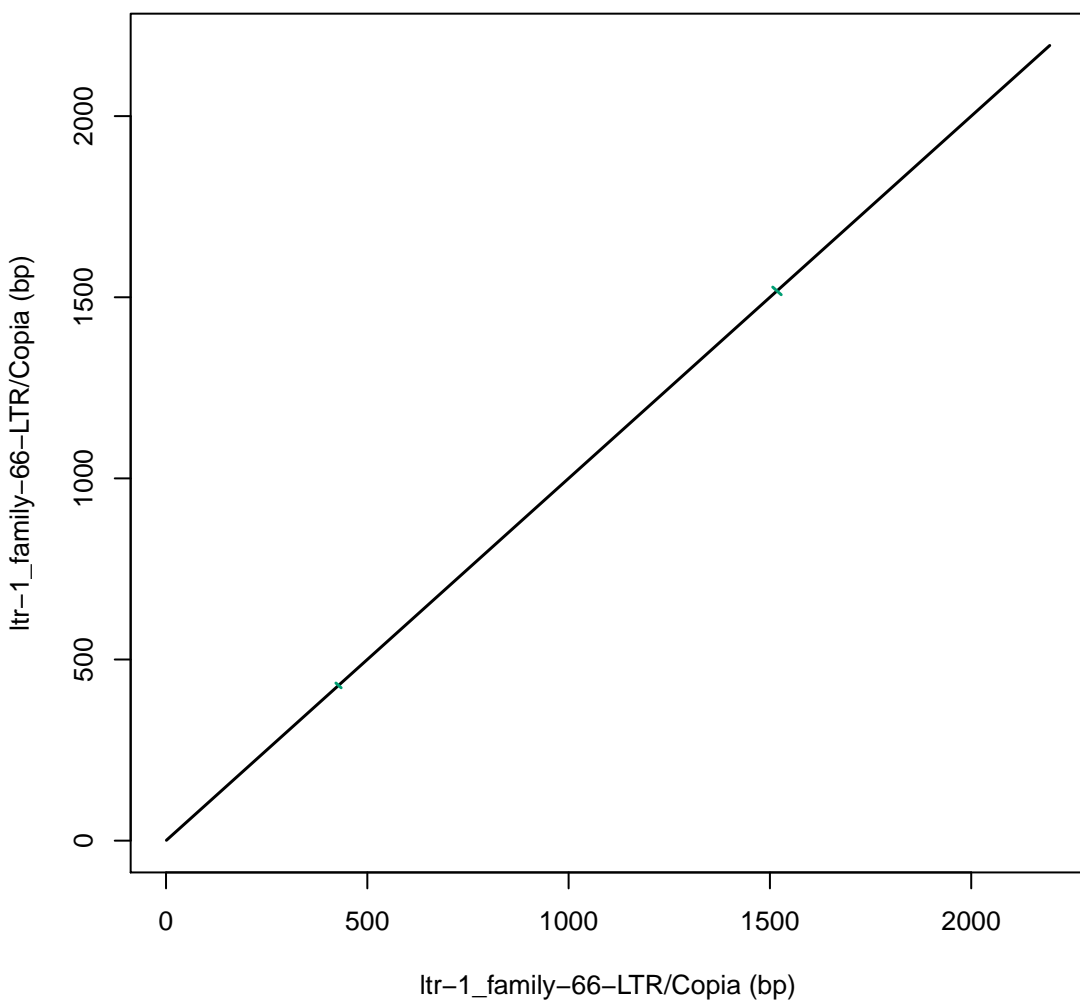
TE: ltr-1\_family-66-LTR/Copia  
consensus size: 2195bp; fragments: 45; full length: 2 ( $\geq 1975.5\text{bp}$ )



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

