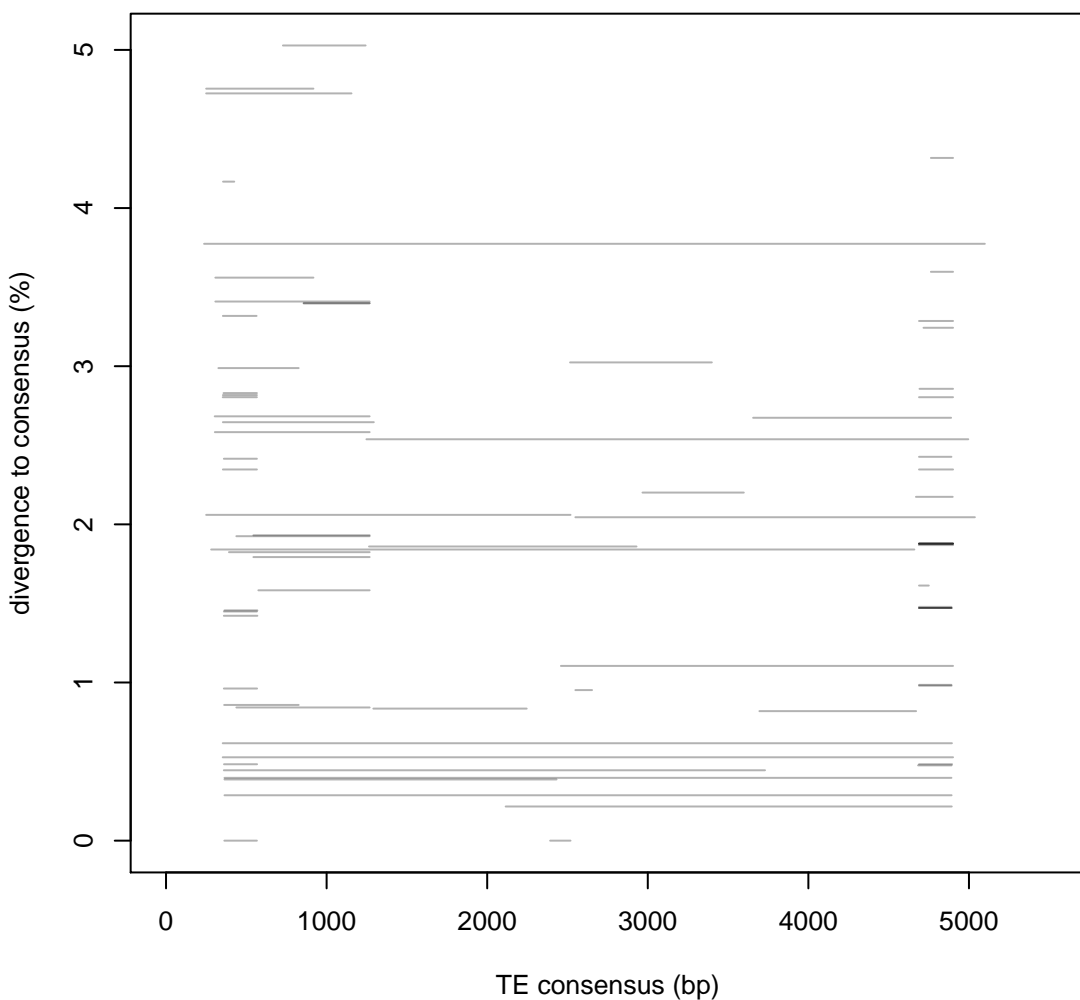
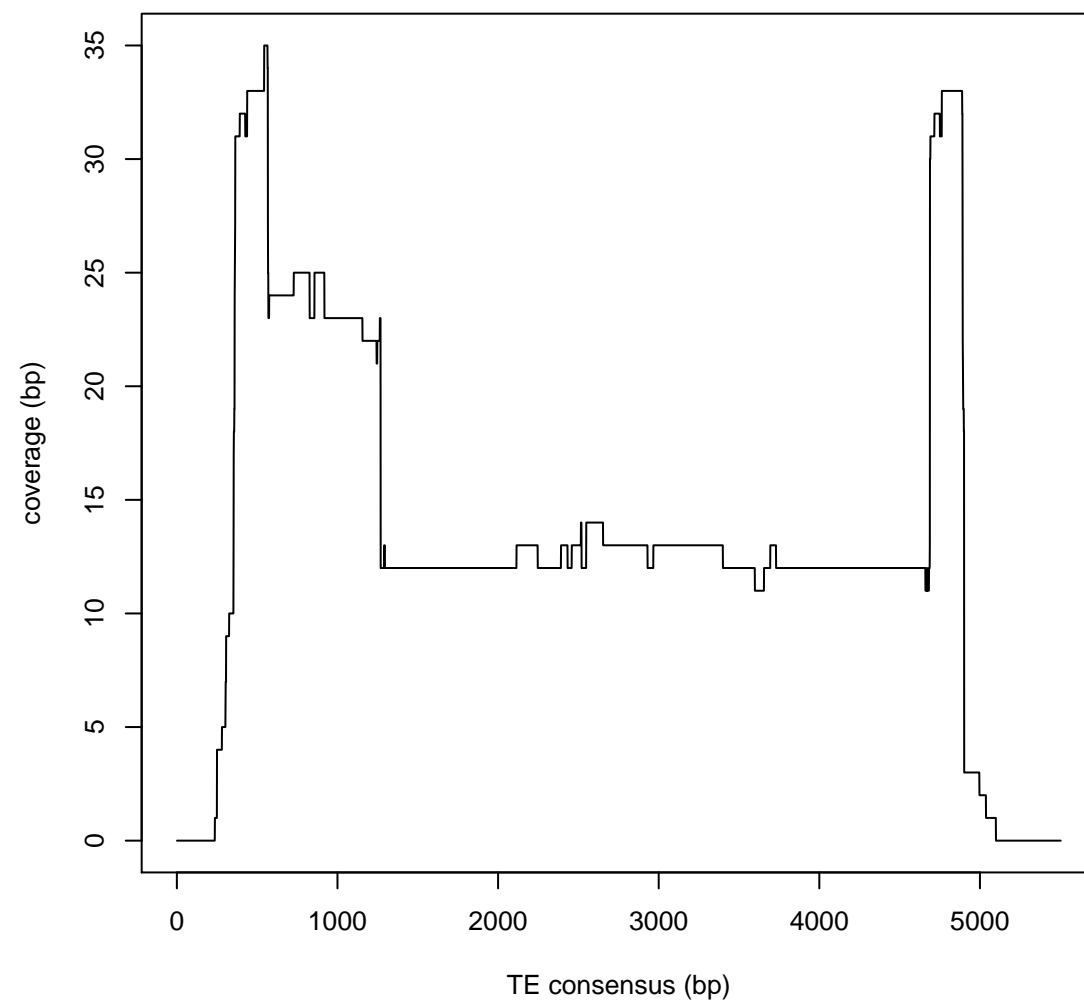


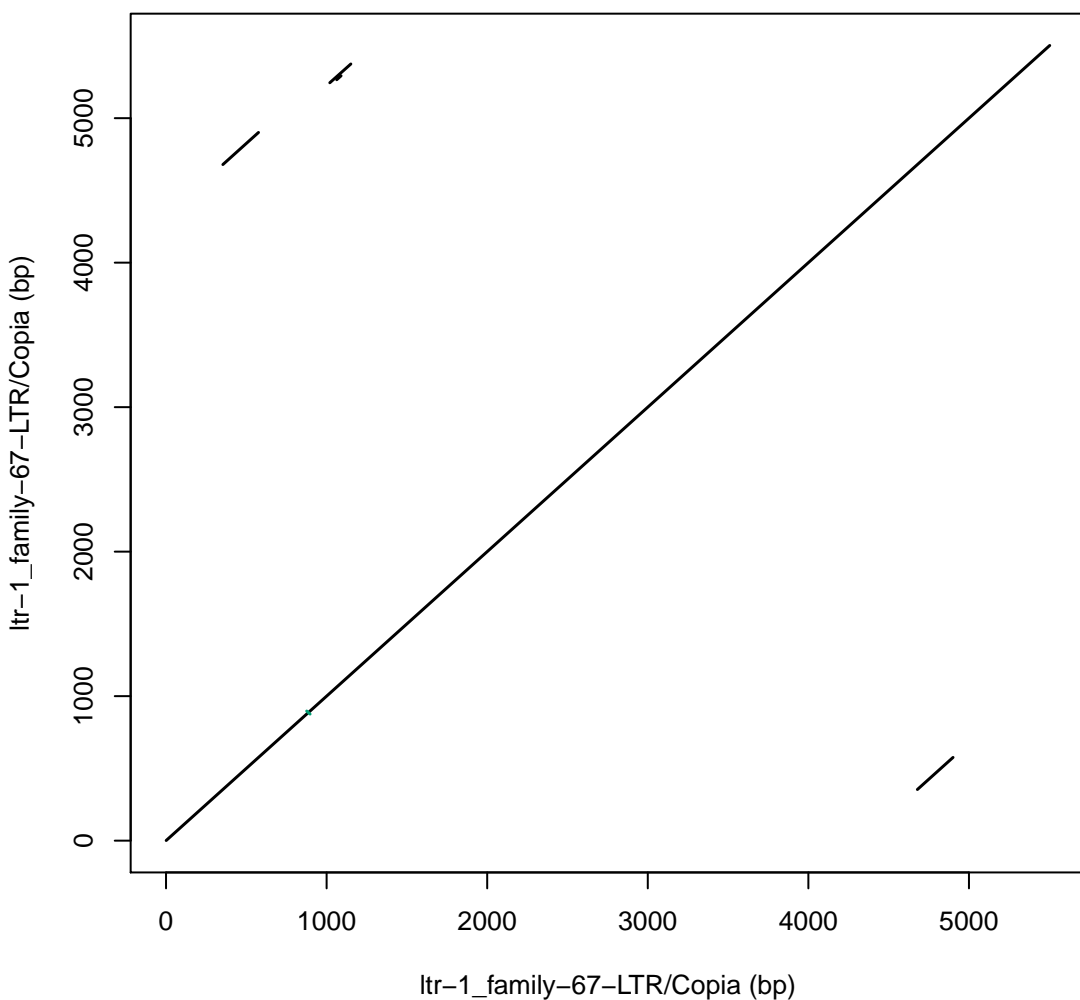
TE: ltr-1\_family-67-LTR/Copia  
consensus size: 5503bp; fragments: 76; full length: 0 ( $\geq 4952.7$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

