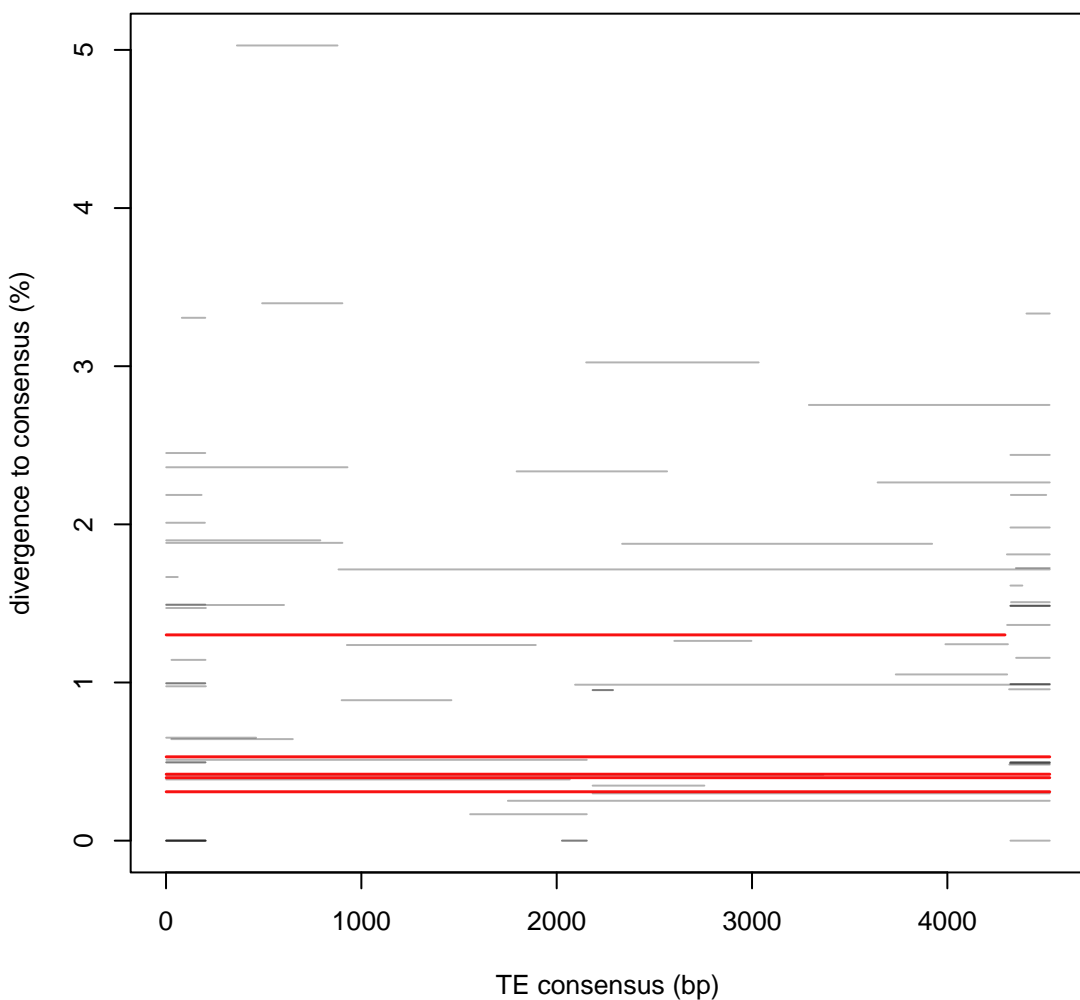
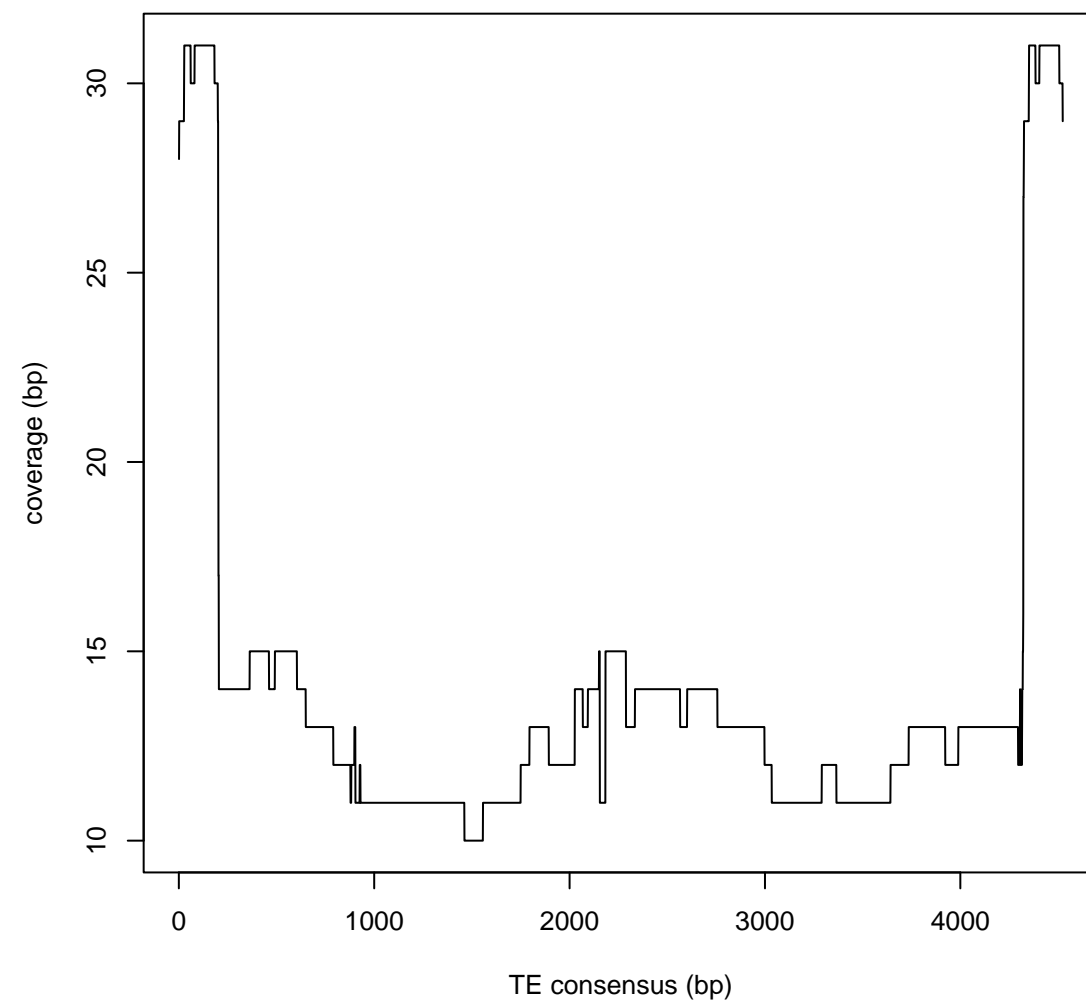


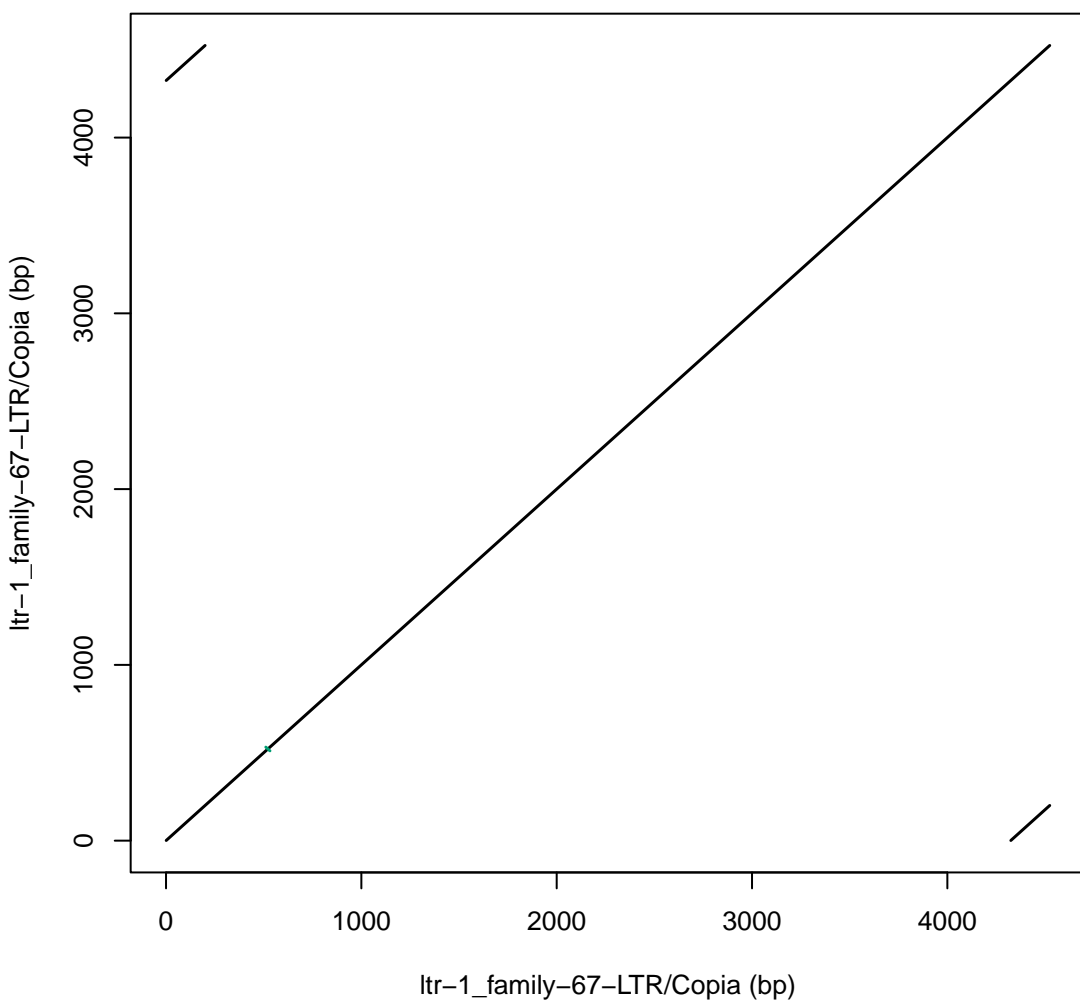
TE: ltr-1\_family-67-LTR/Copia  
consensus size: 4524bp; fragments: 76; full length: 5 ( $\geq 4071.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

