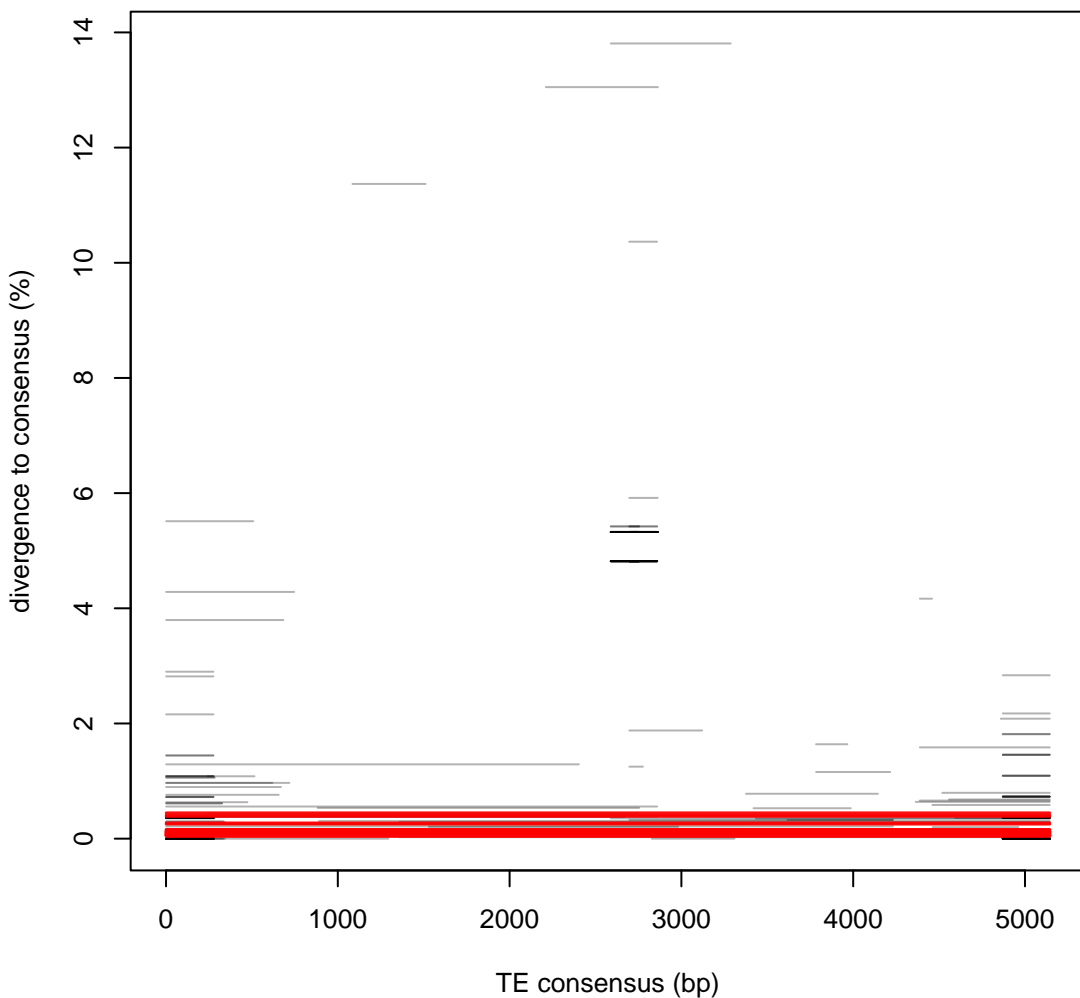
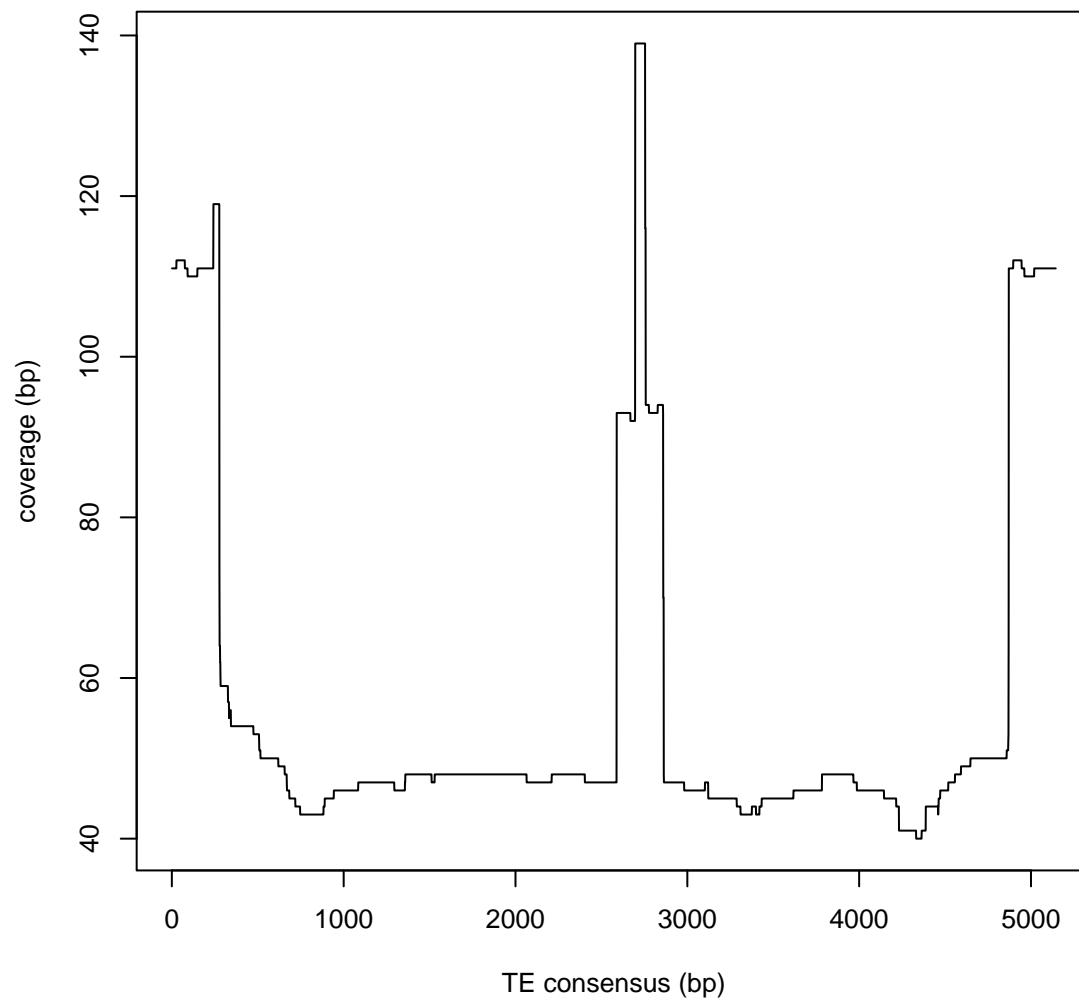


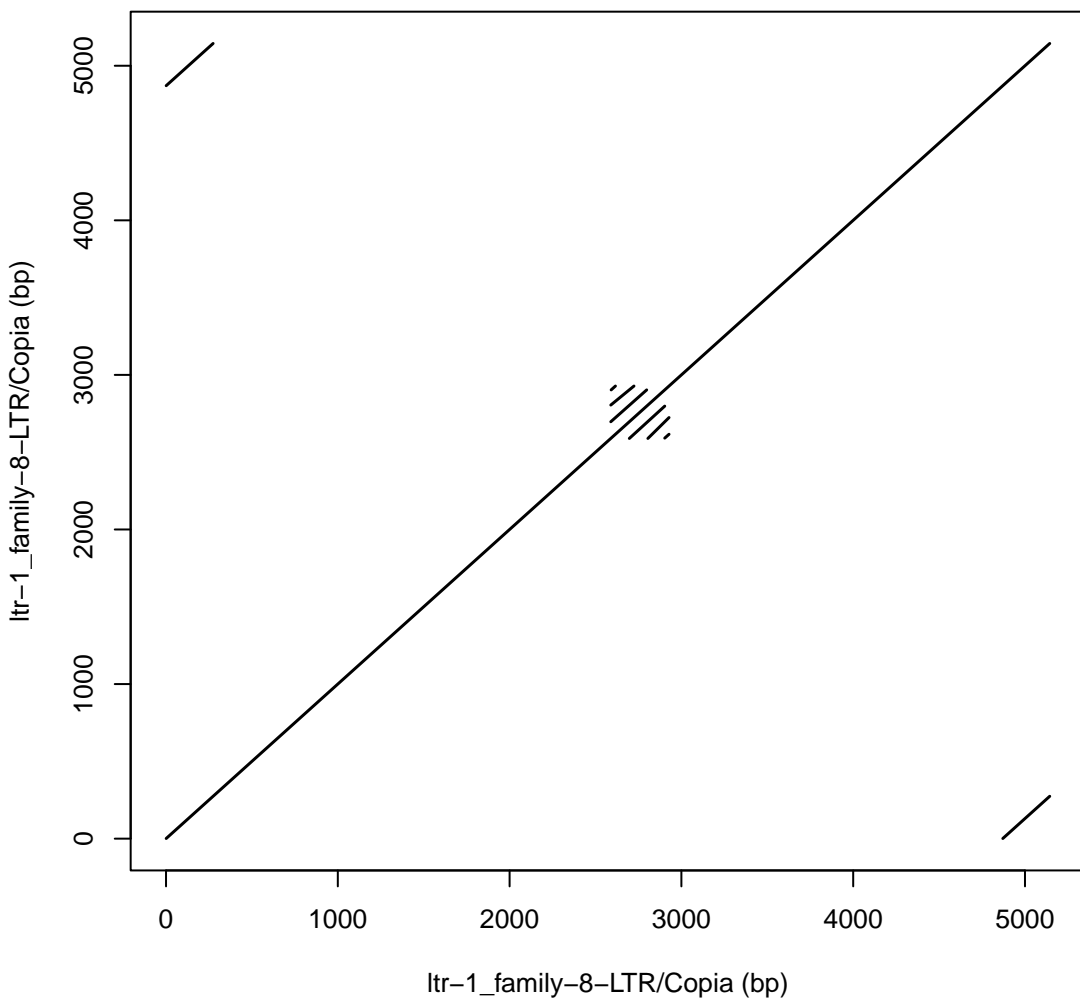
TE: ltr-1\_family-8-LTR/Copia  
consensus size: 5144bp; fragments: 305; full length: 32 (>=4629.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

