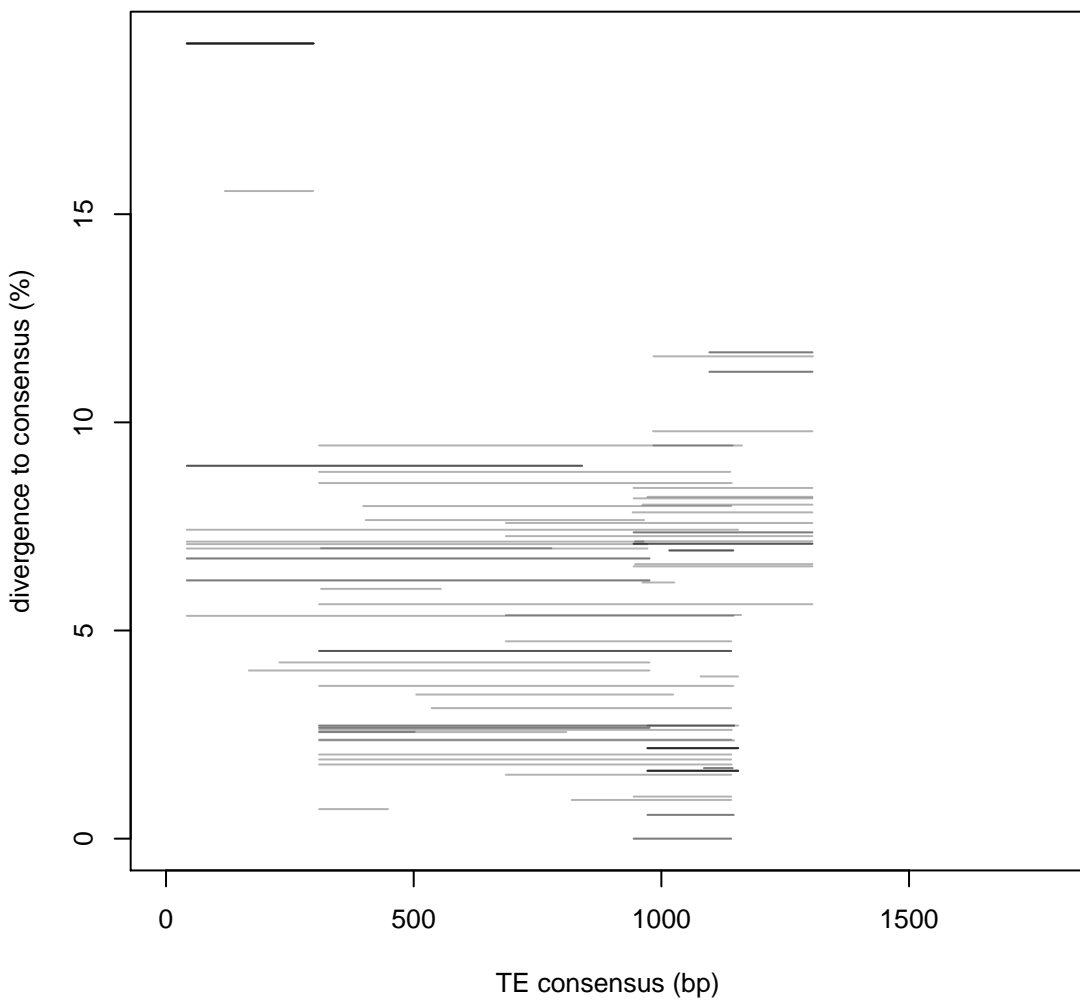
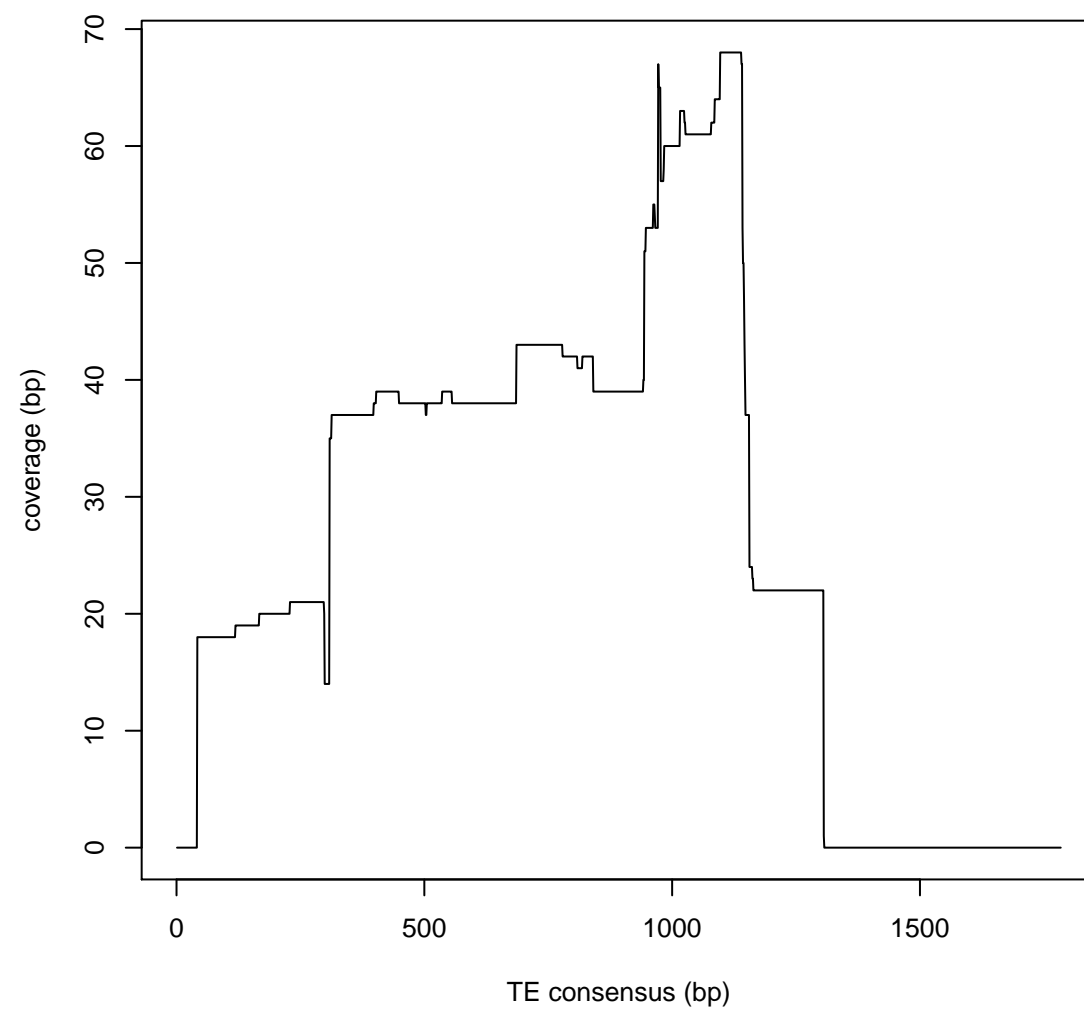


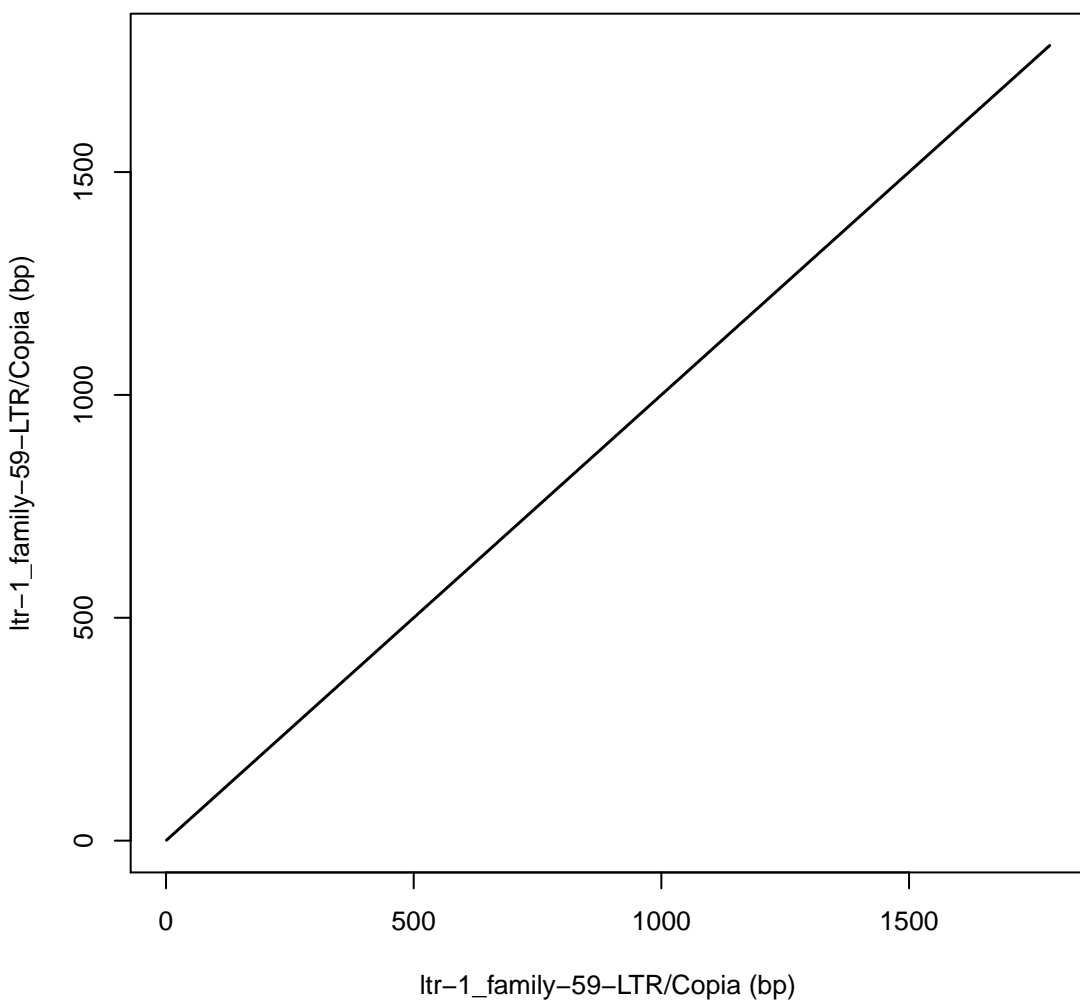
TE: ltr-1\_family-59-LTR/Copia  
consensus size: 1784bp; fragments: 97; full length: 0 (>=1605.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

