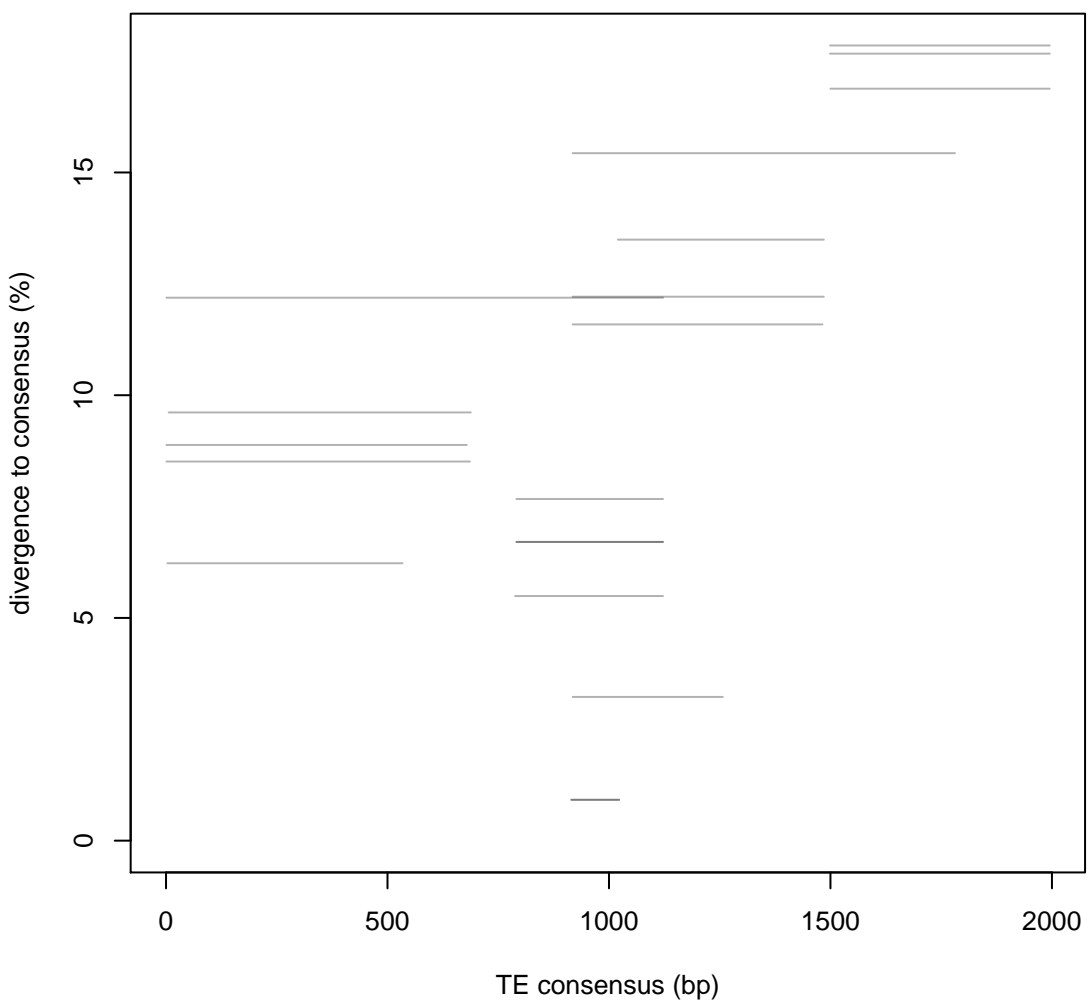
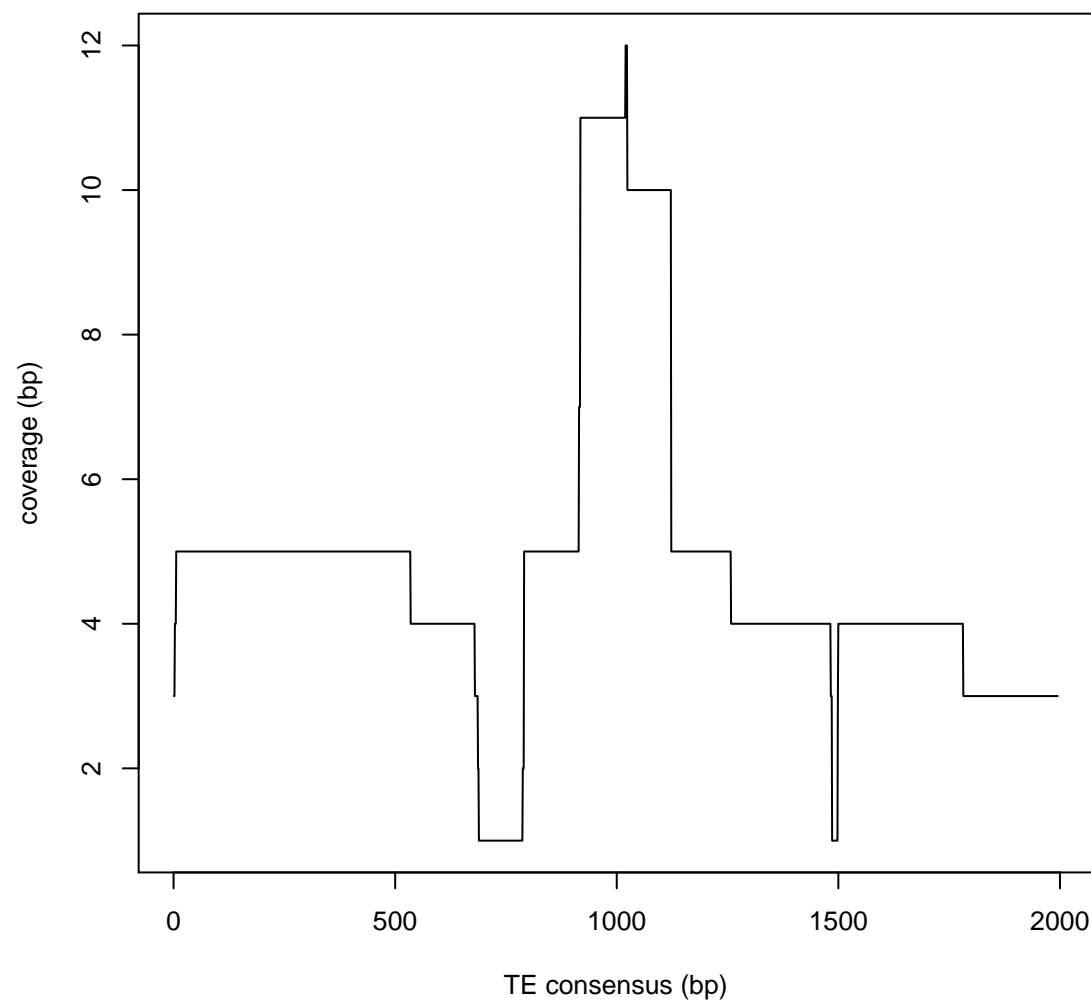


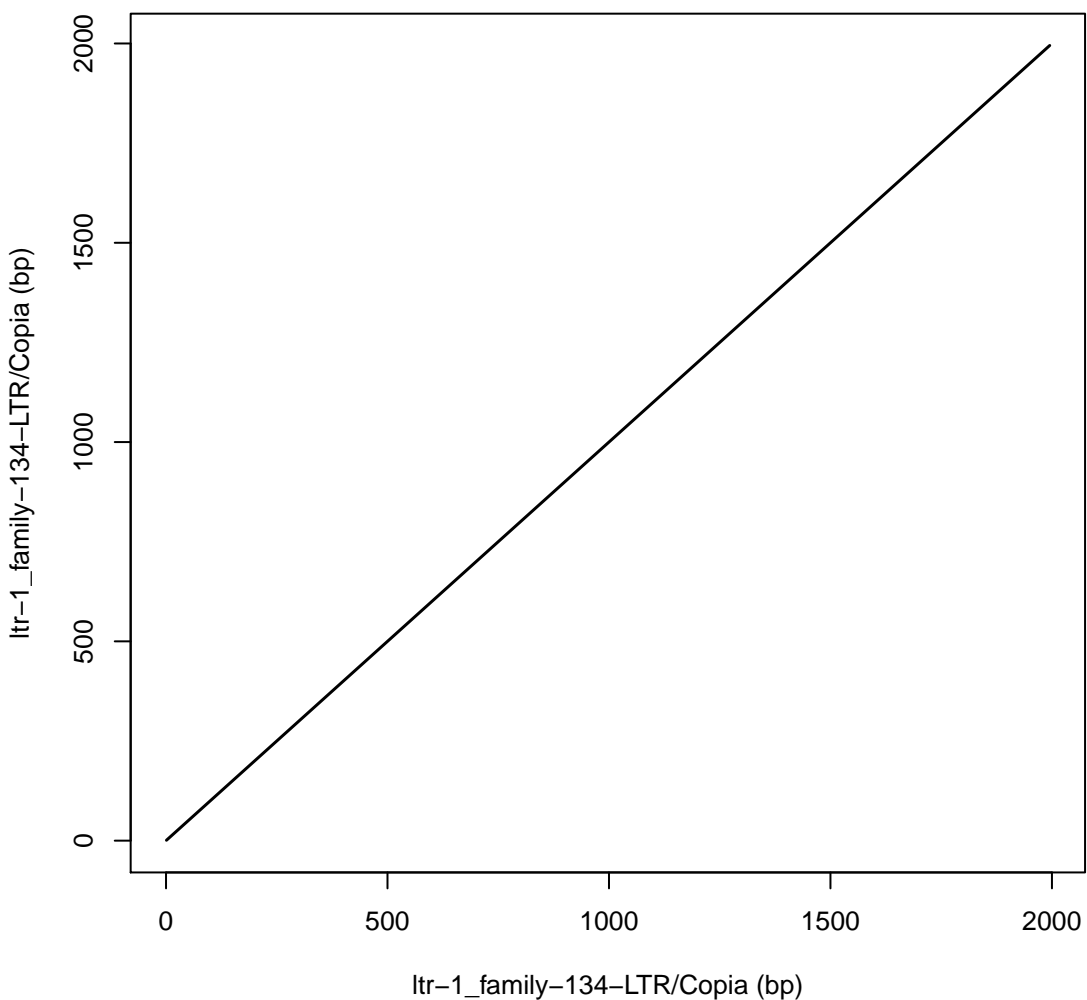
TE: ltr-1_family-134-LTR/Copia
consensus size: 1995bp; fragments: 19; full length: 0 ($\geq 1795.5\text{bp}$)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

