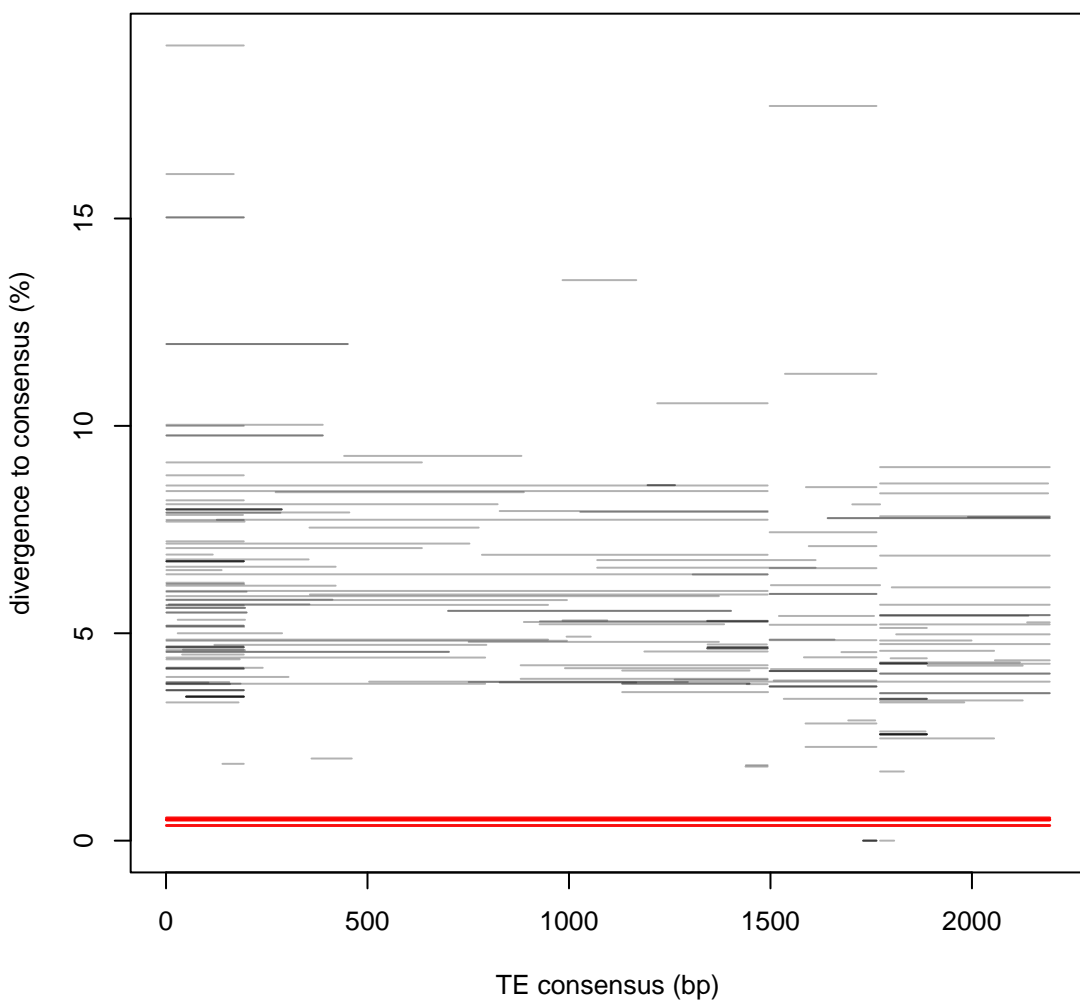
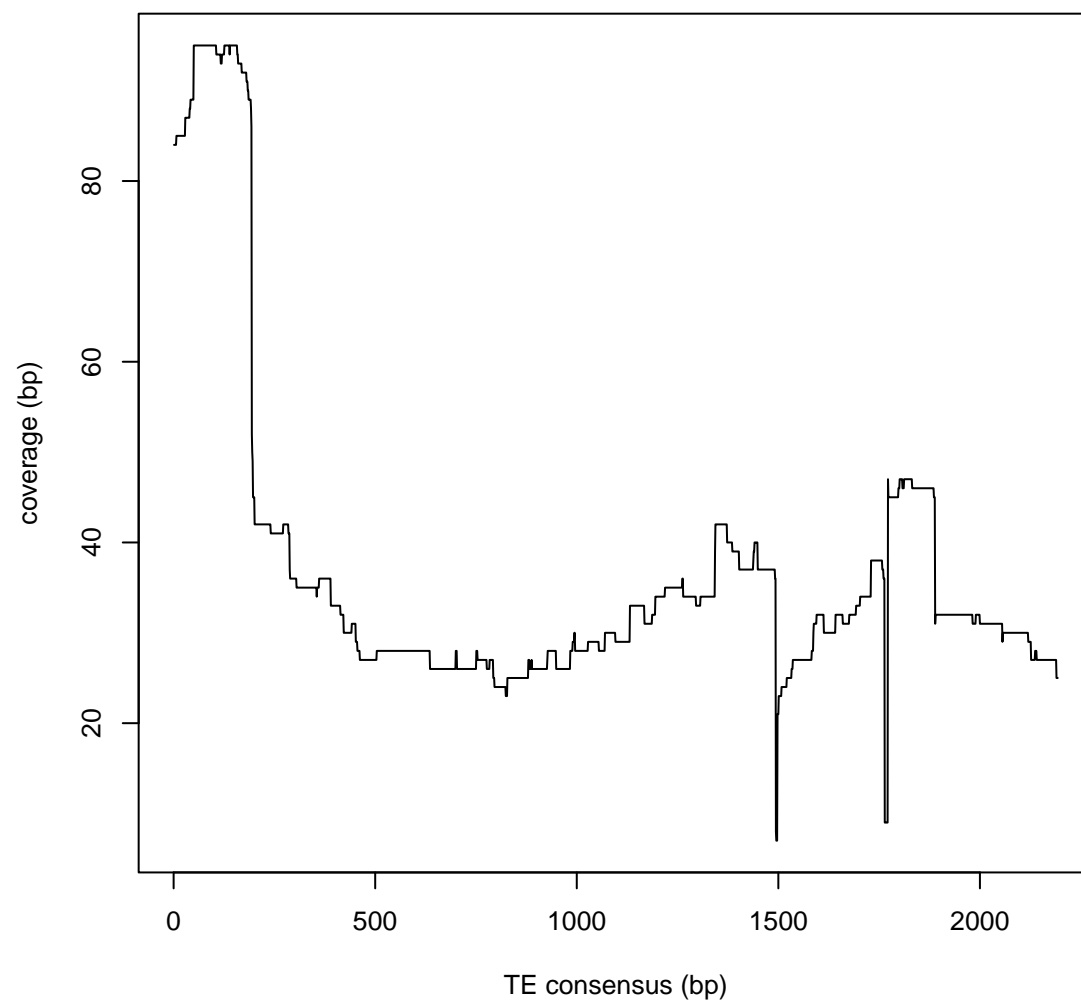


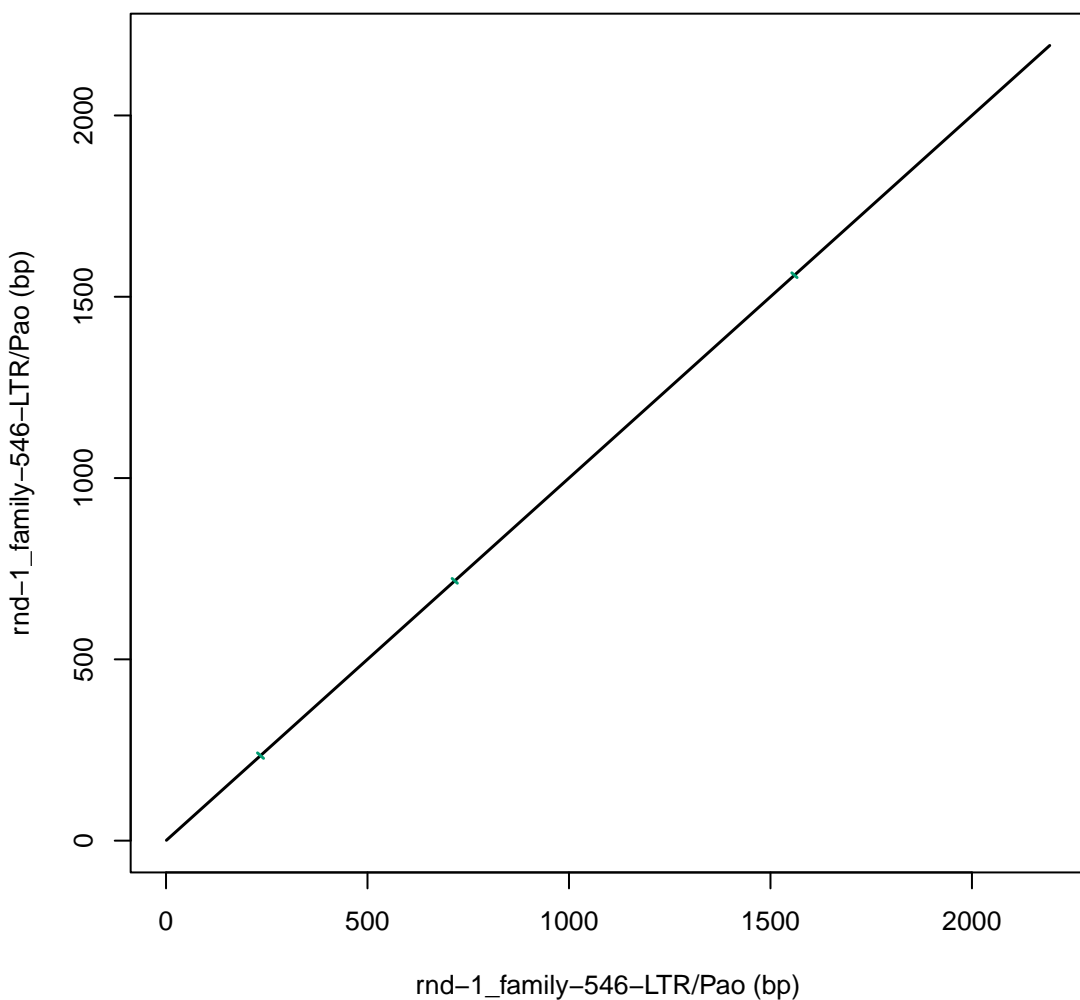
TE: rnd-1\_family-546-LTR/Pao  
consensus size: 2193bp; fragments: 223; full length: 4 ( $\geq 1973.7$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

