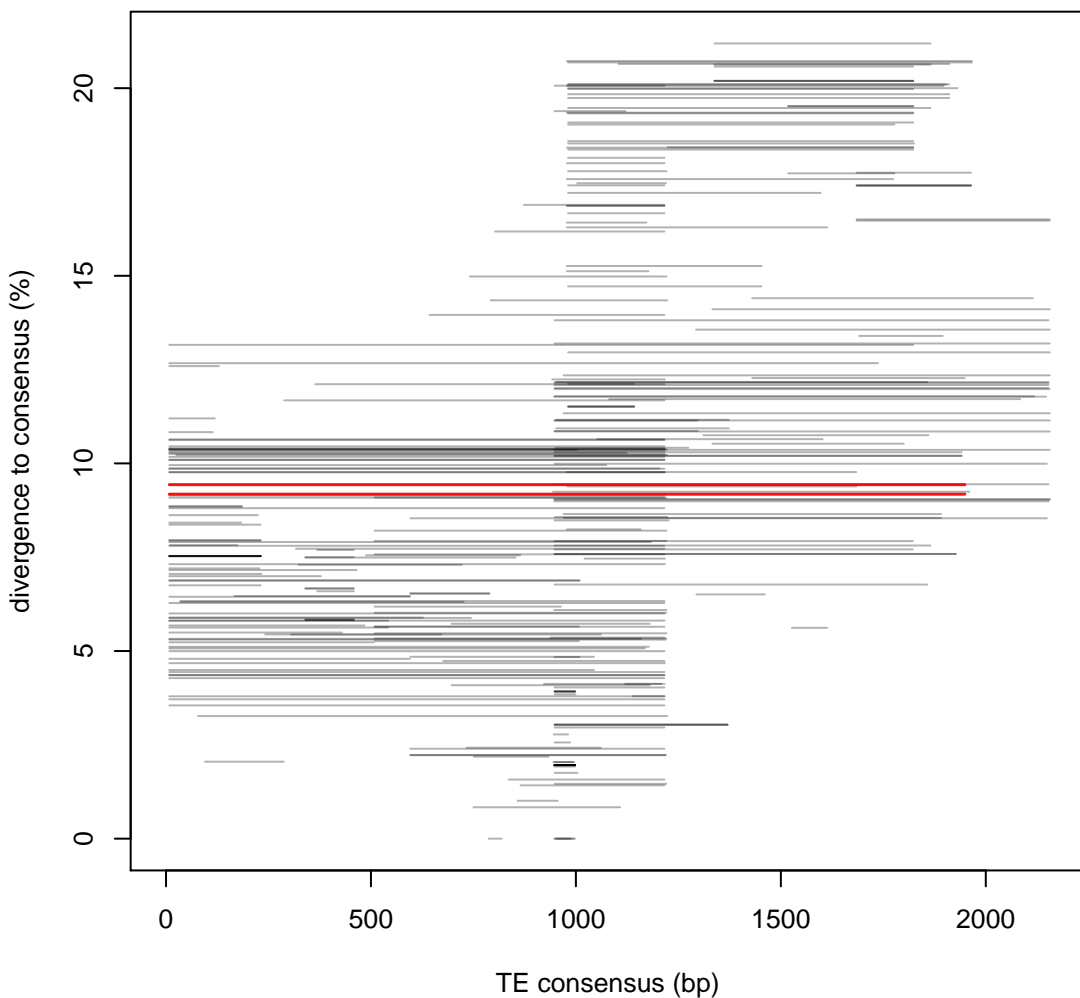
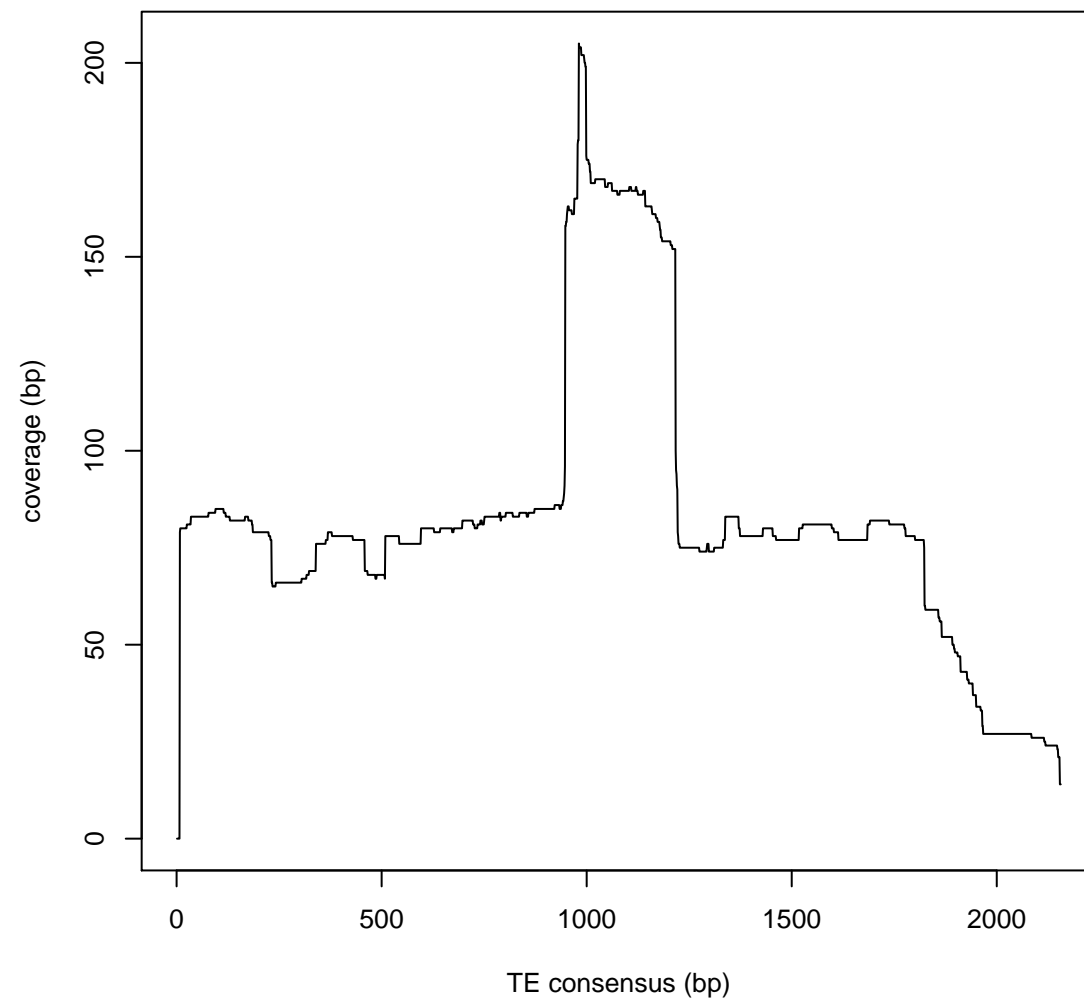


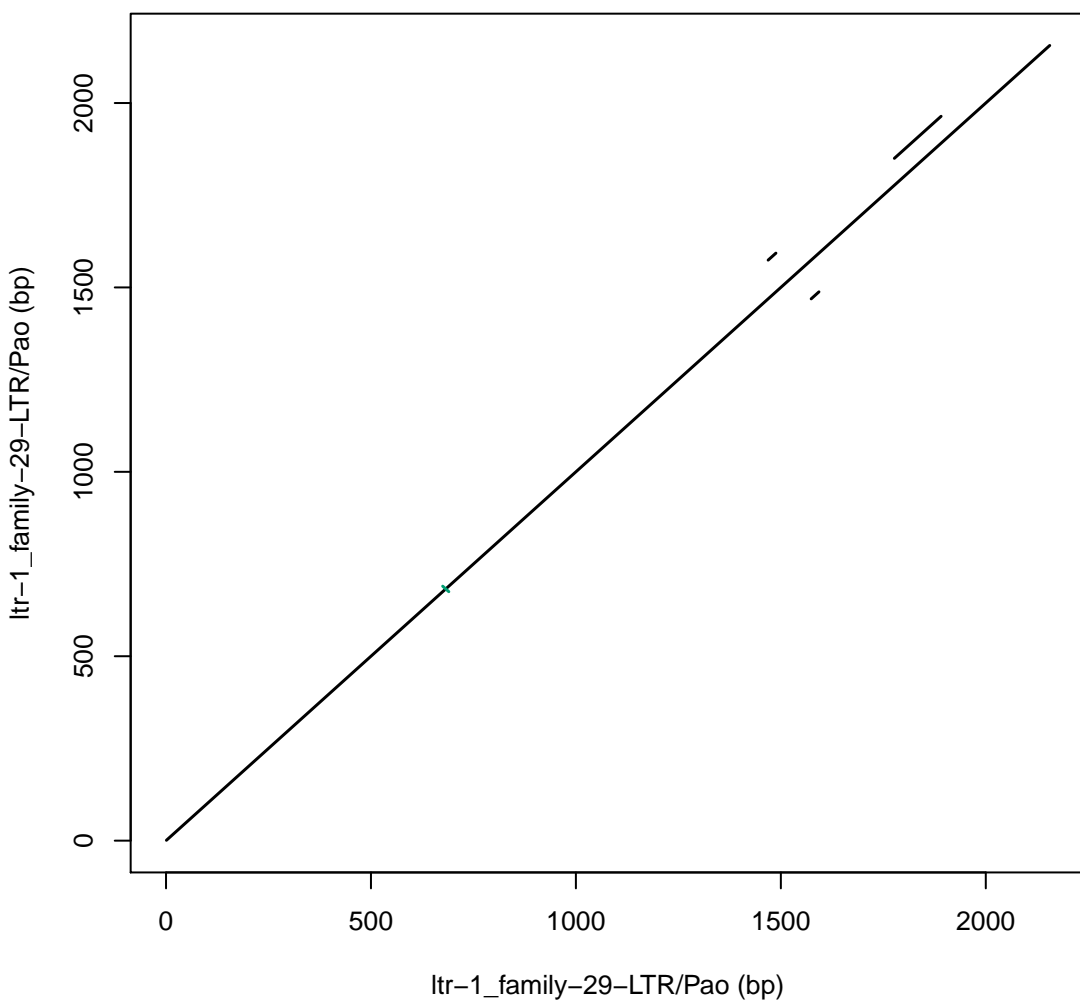
TE: ltr-1\_family-29-LTR/Pao  
consensus size: 2156bp; fragments: 290; full length: 2 ( $\geq 1940.4$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

