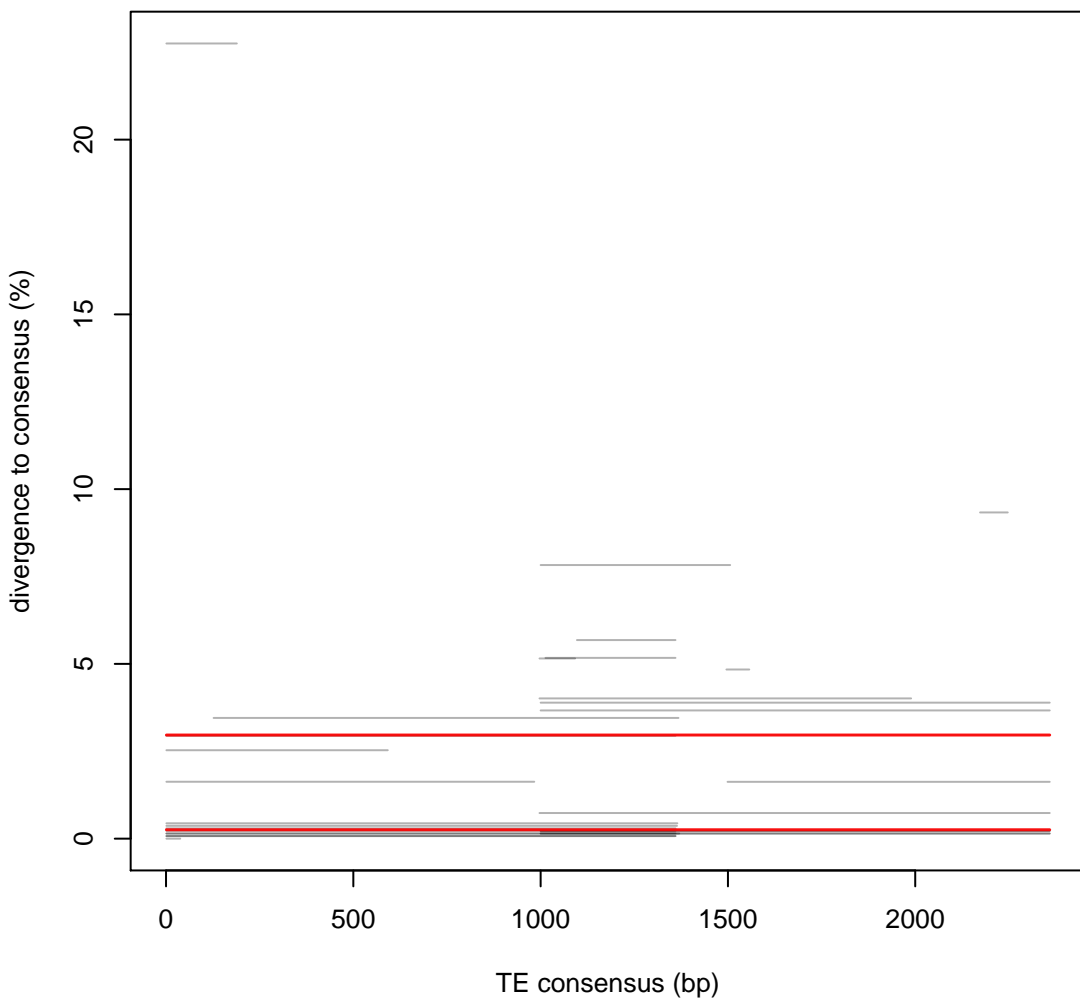
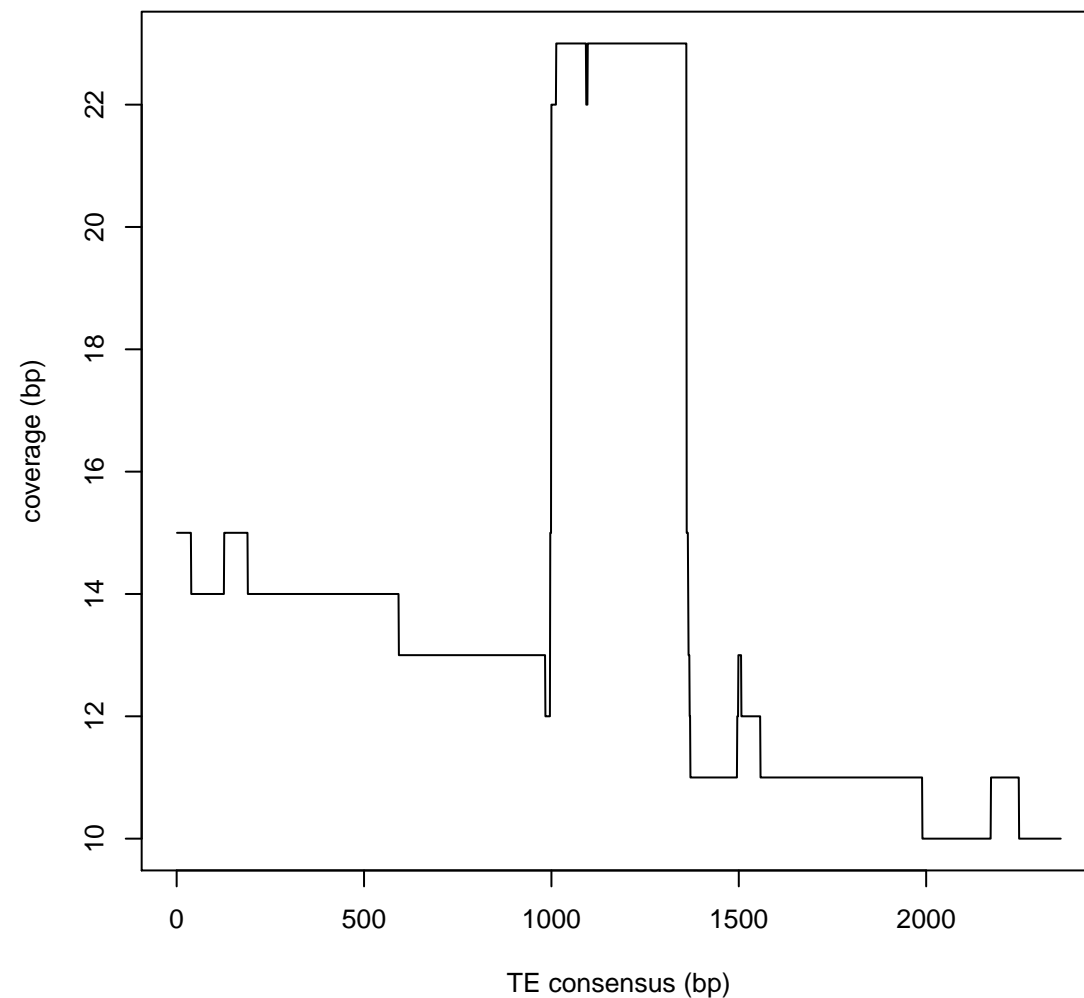


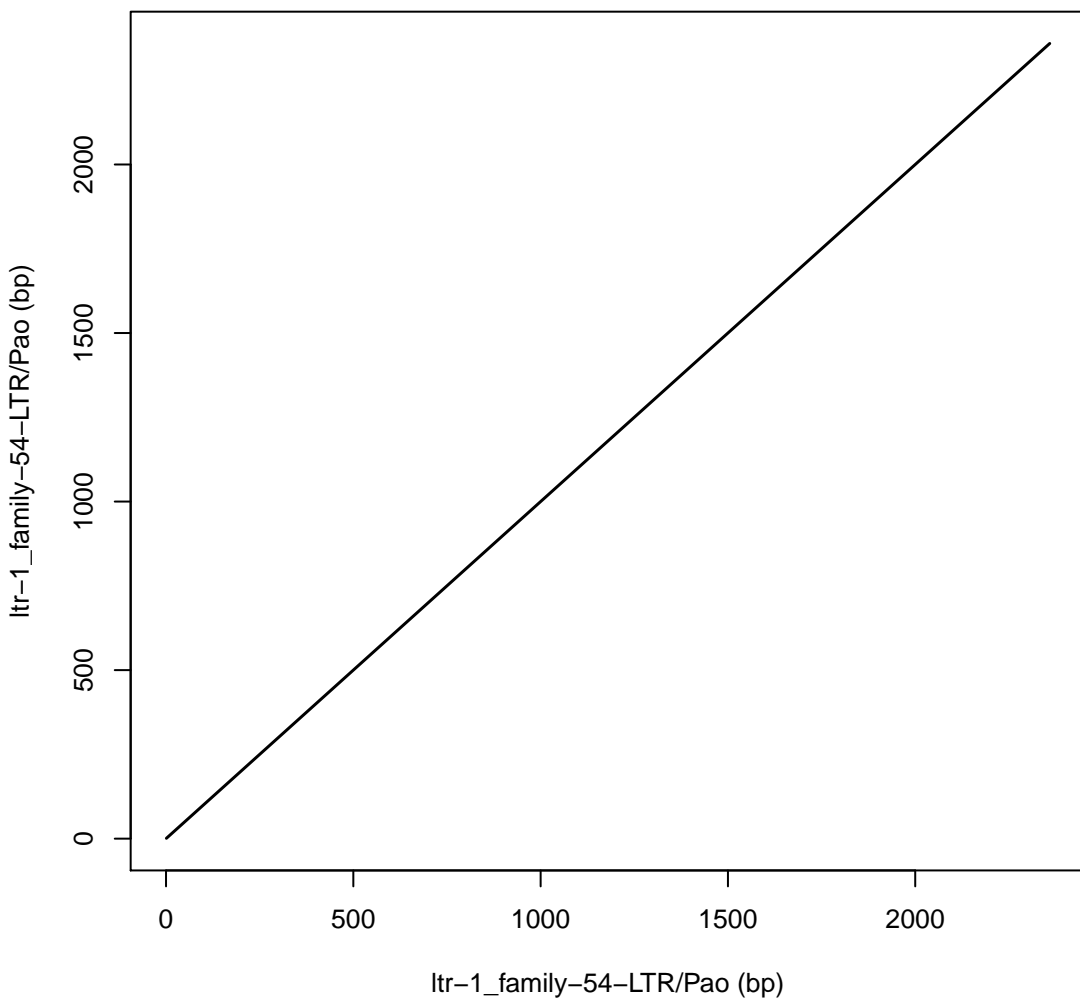
TE: ltr-1\_family-54-LTR/Pao  
consensus size: 2359bp; fragments: 31; full length: 2 ( $\geq 2123.1$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

