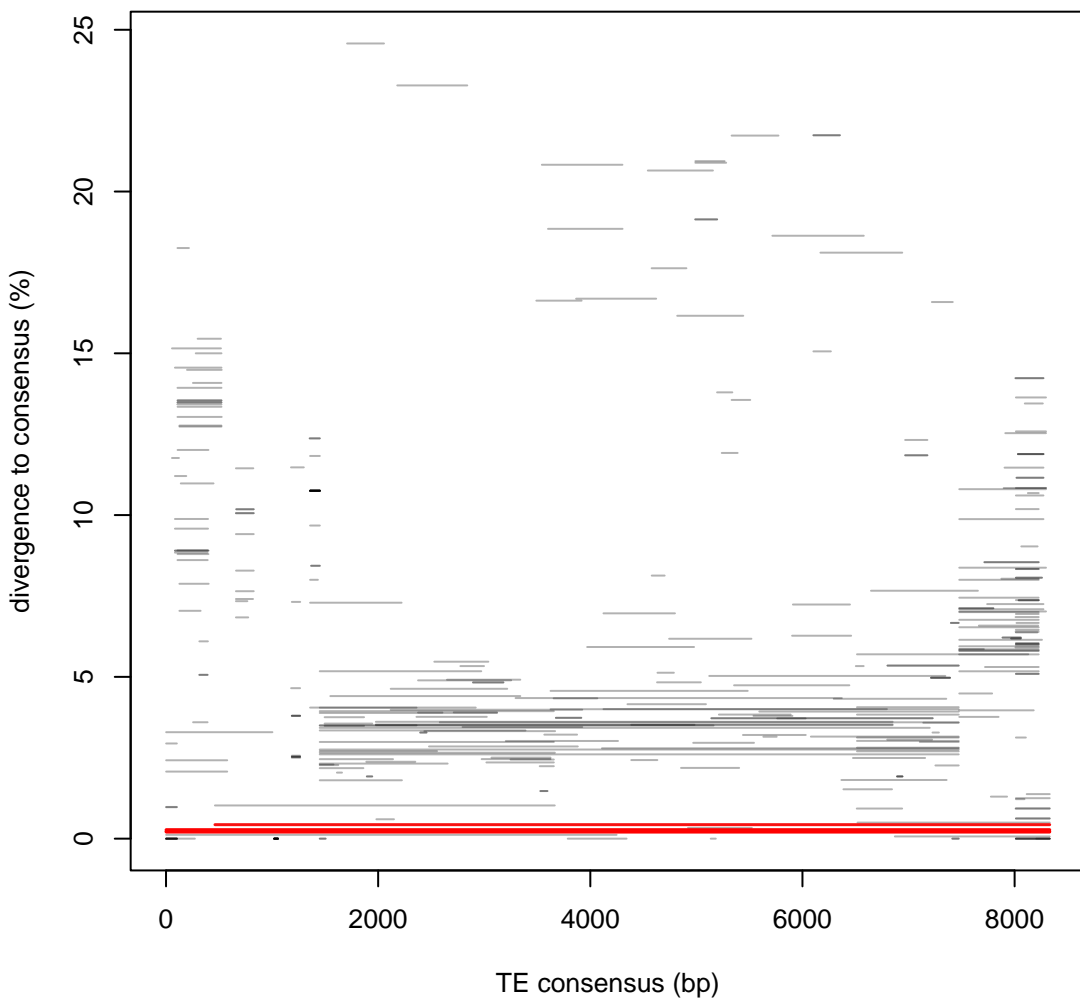
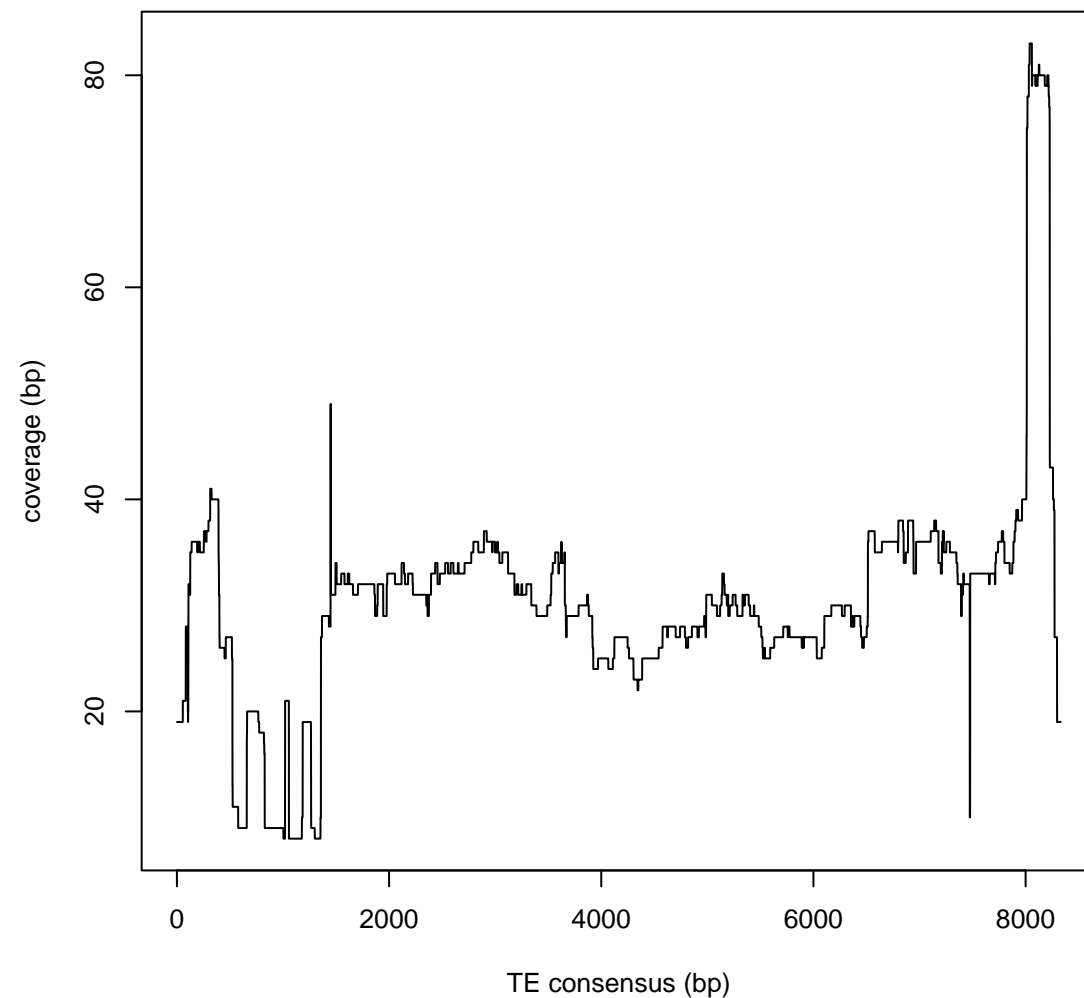


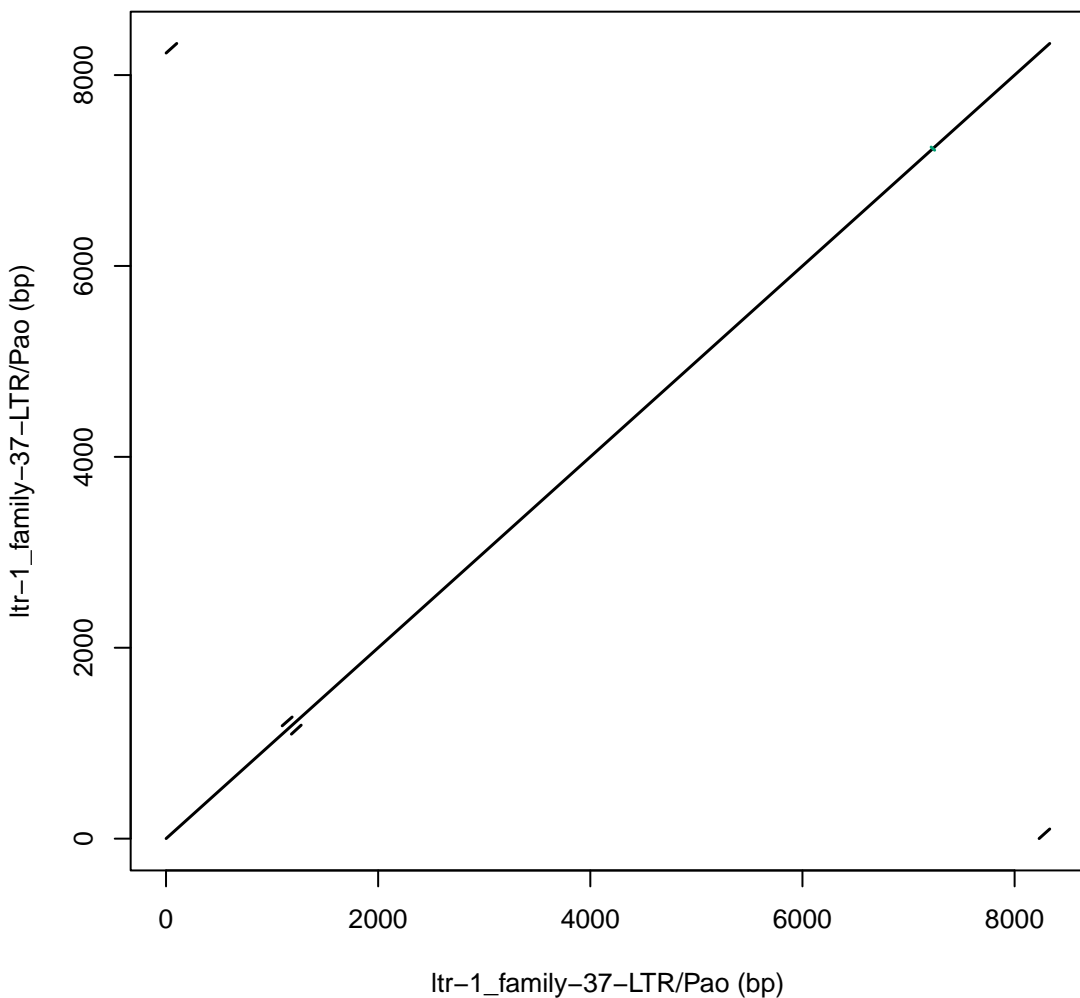
TE: ltr-1\_family-37-LTR/Pao  
consensus size: 8331bp; fragments: 350; full length: 6 ( $\geq 7497.9$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

