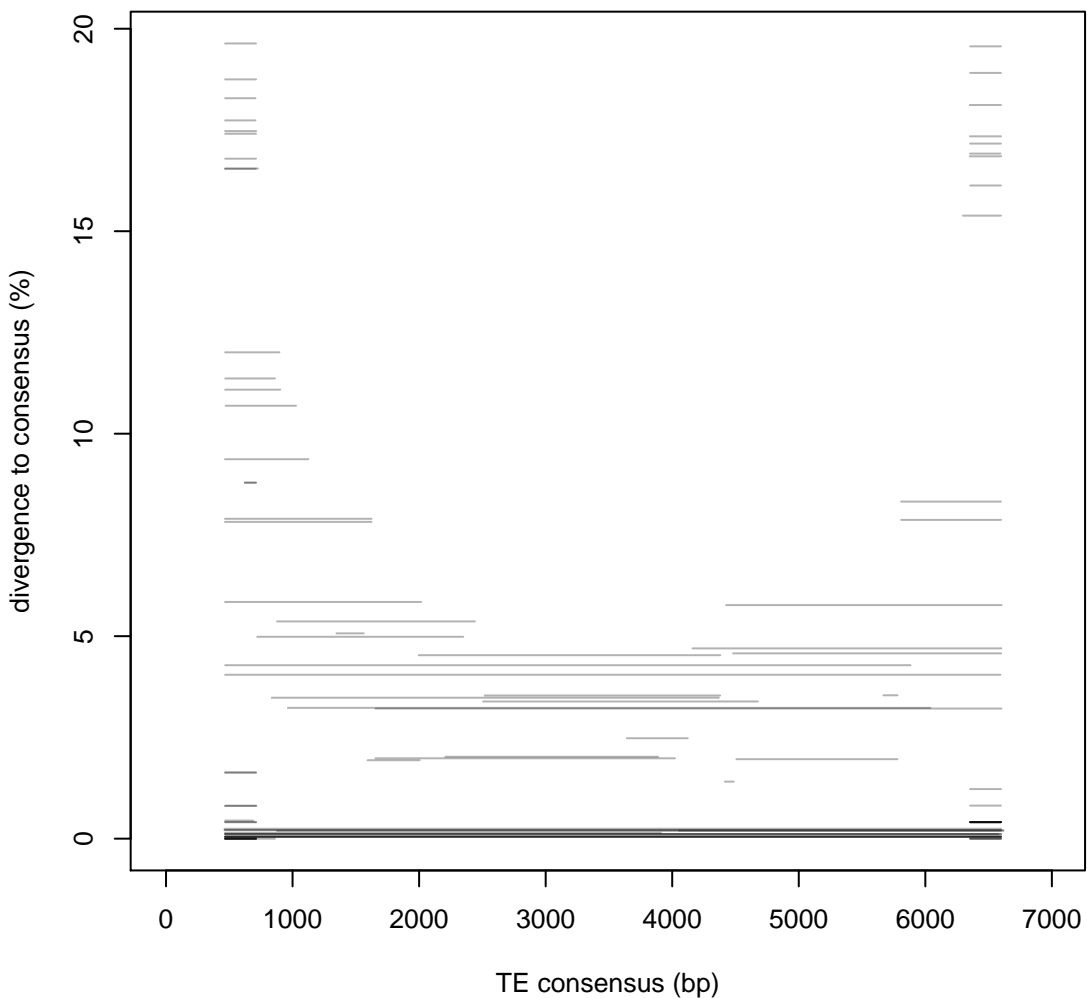
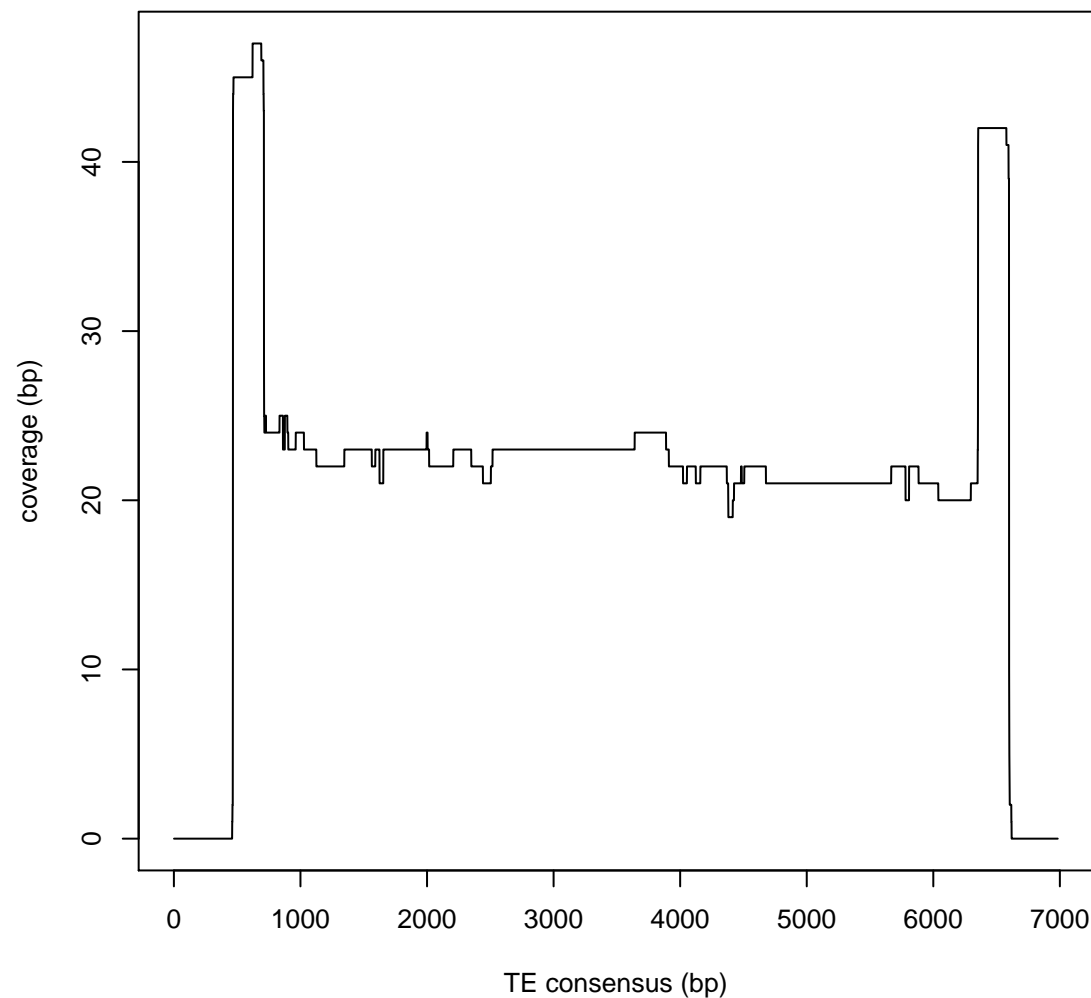


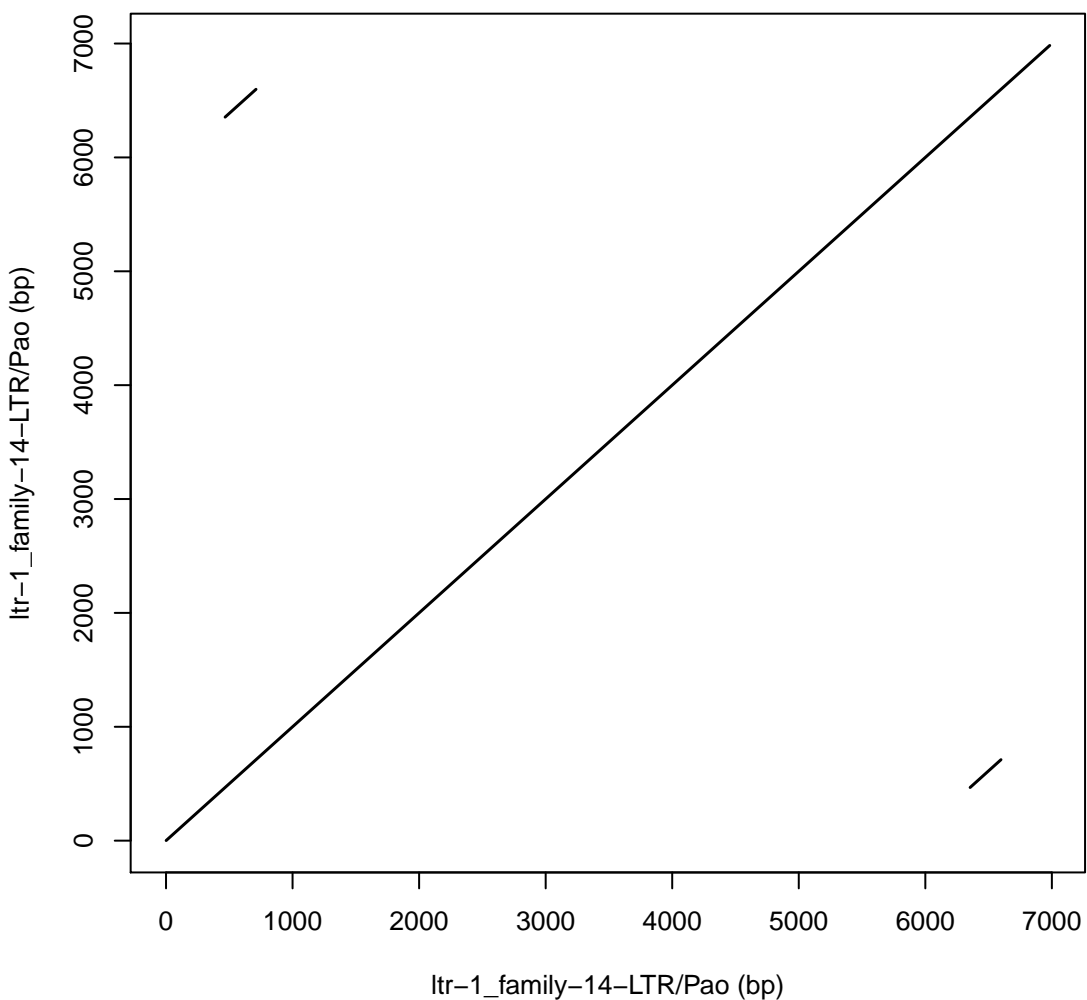
TE: ltr-1_family-14-LTR/Pao
consensus size: 6983bp; fragments: 92; full length: 0 (>=6284.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

