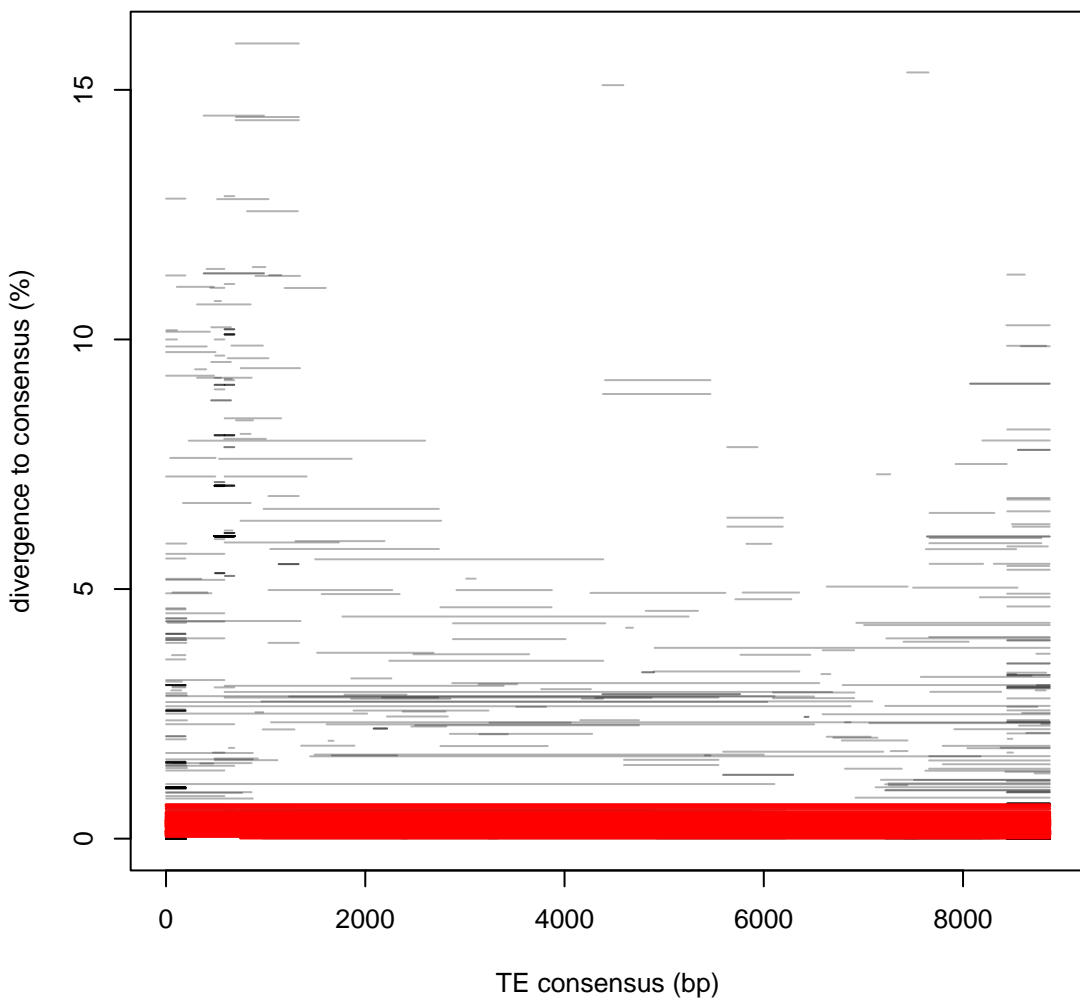
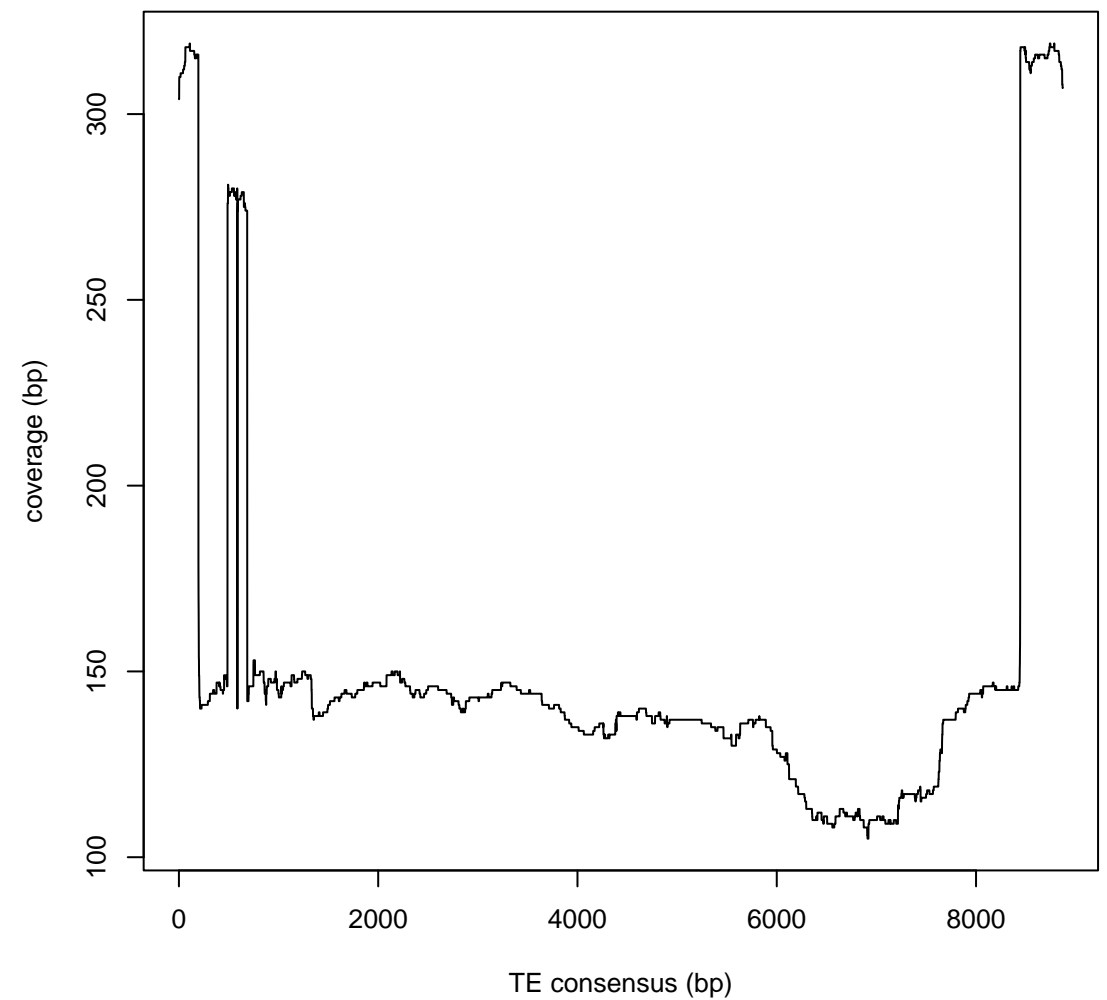


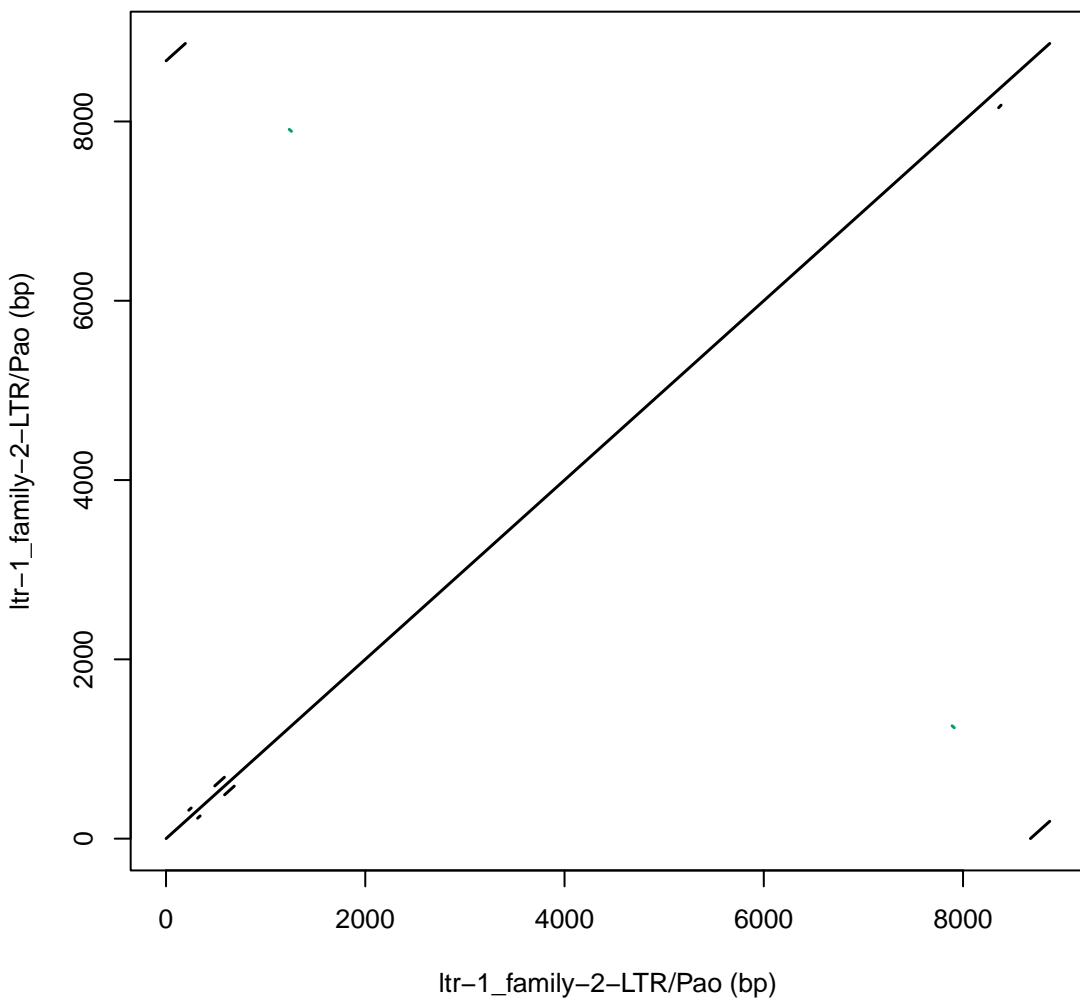
TE: ltr-1_family-2-LTR/Pao
consensus size: 8870bp; fragments: 1009; full length: 72 (>=7983bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

