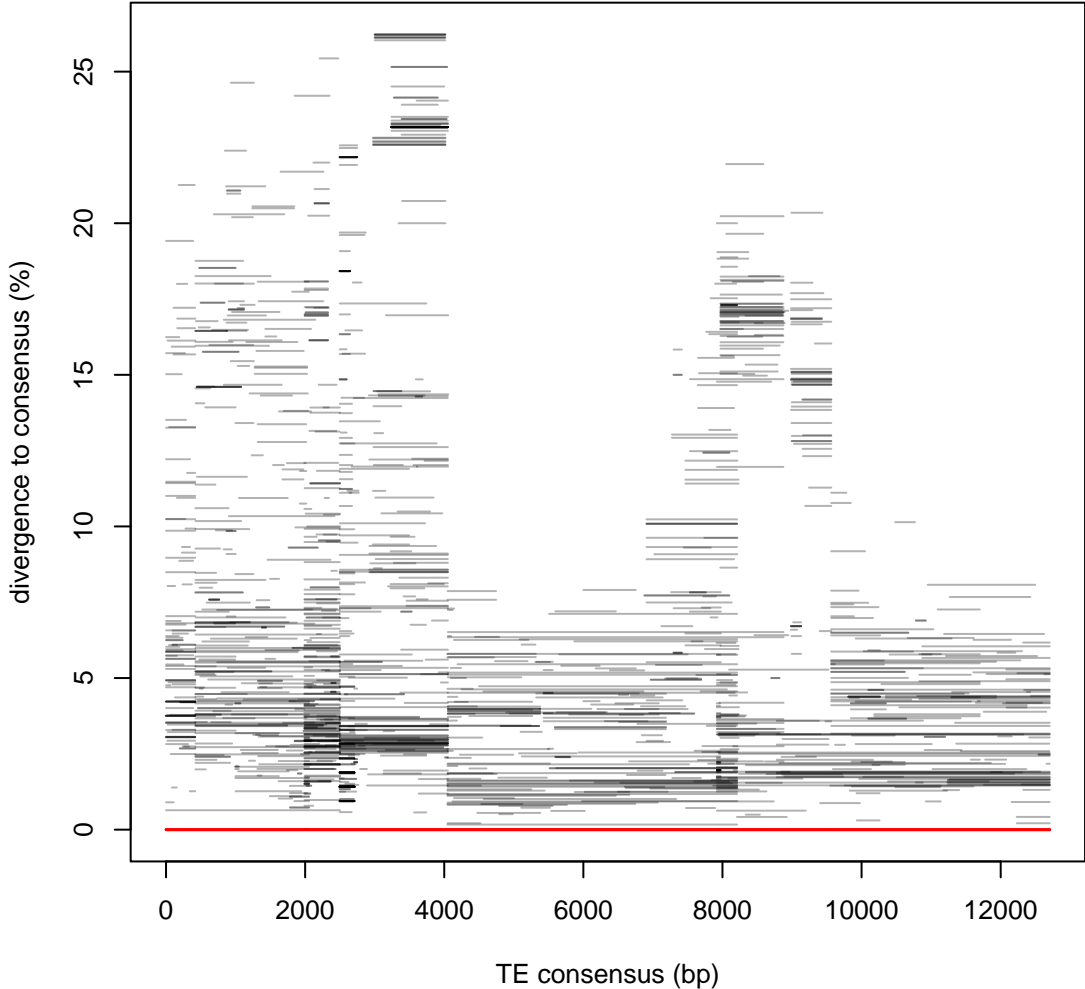
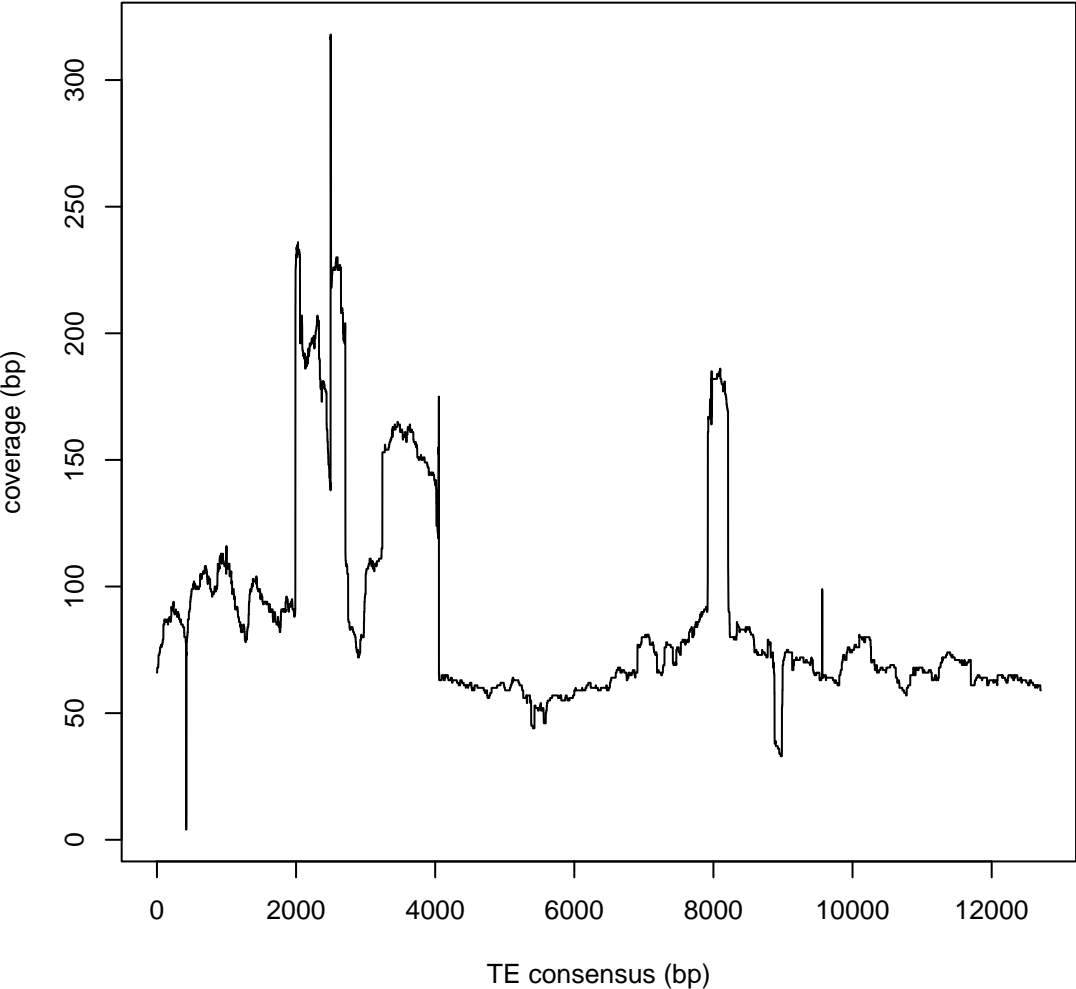


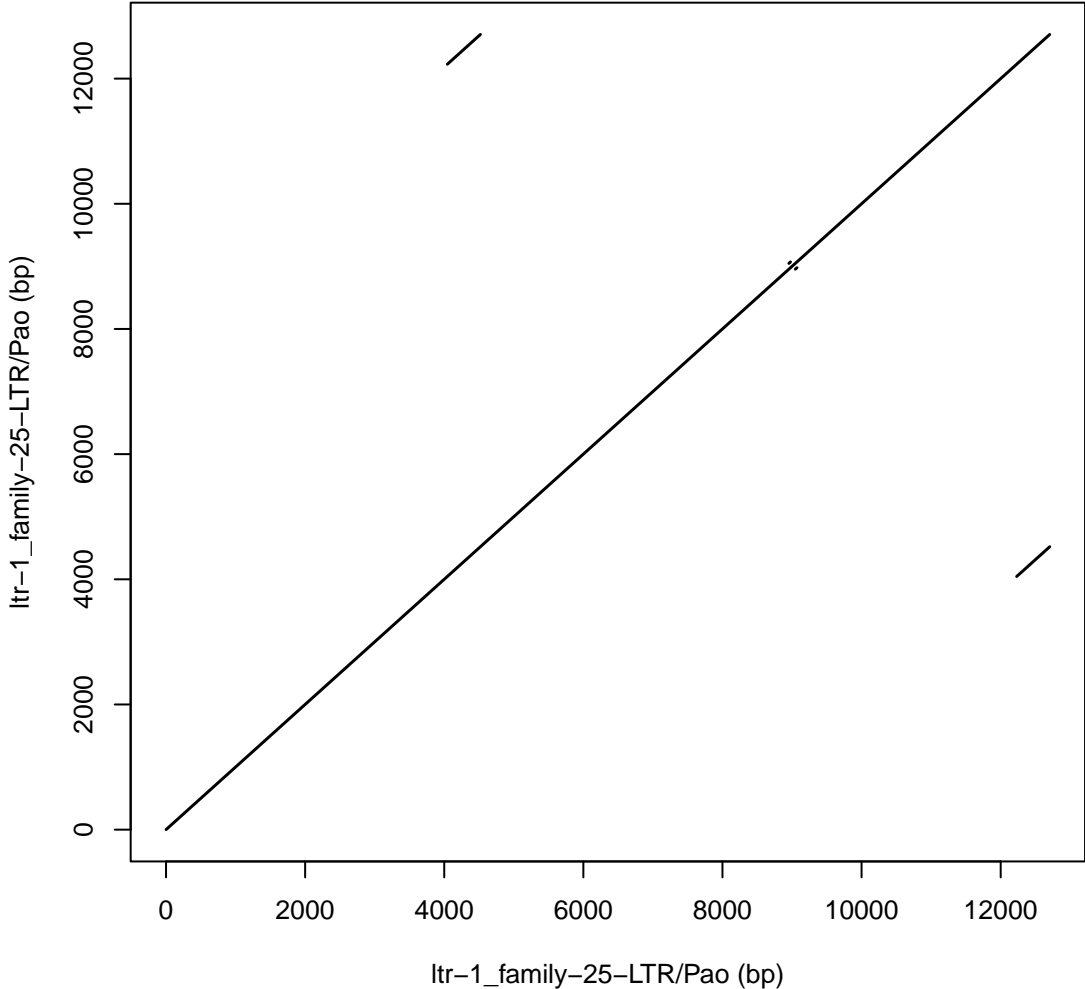
TE: ltr-1_family-25-LTR/Pao
consensus size: 12705bp; fragments: 1590; full length: 2 (>=11434.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

