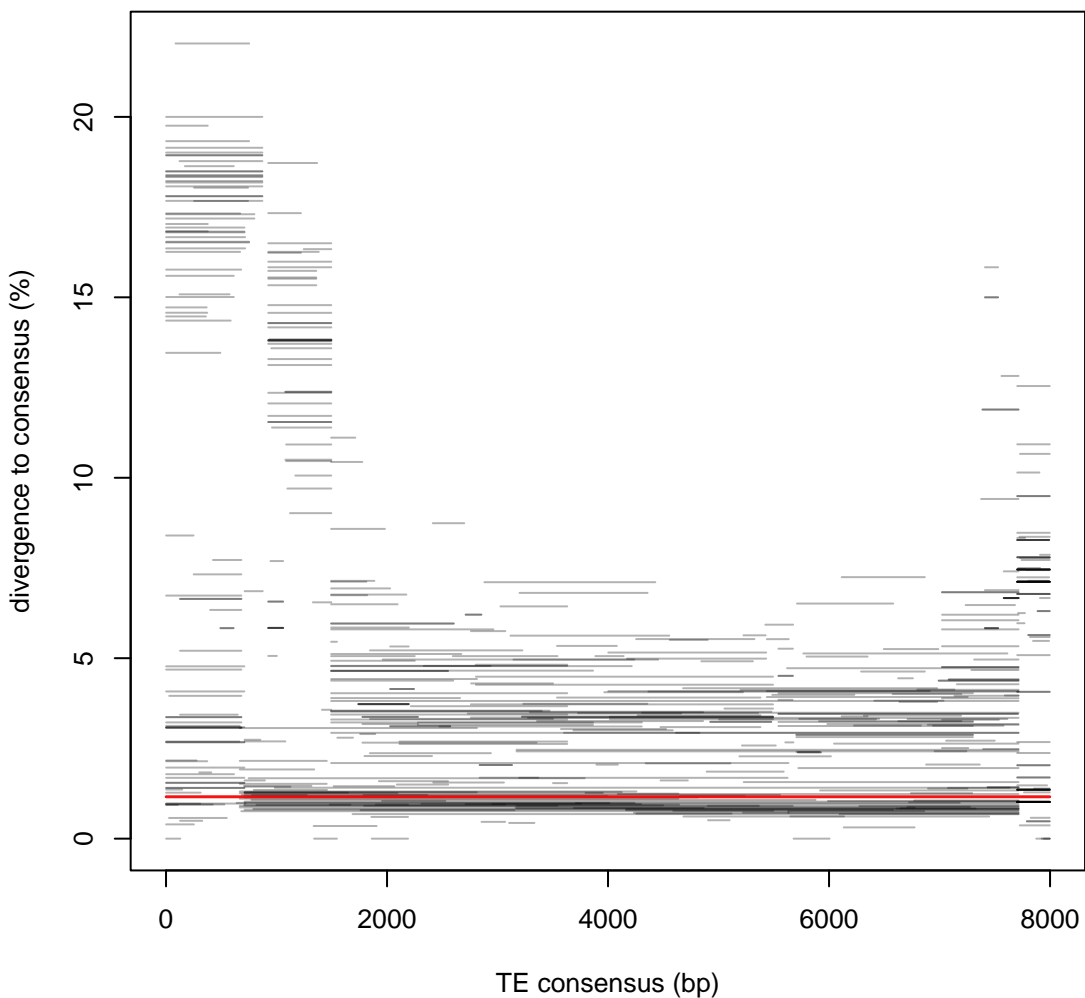
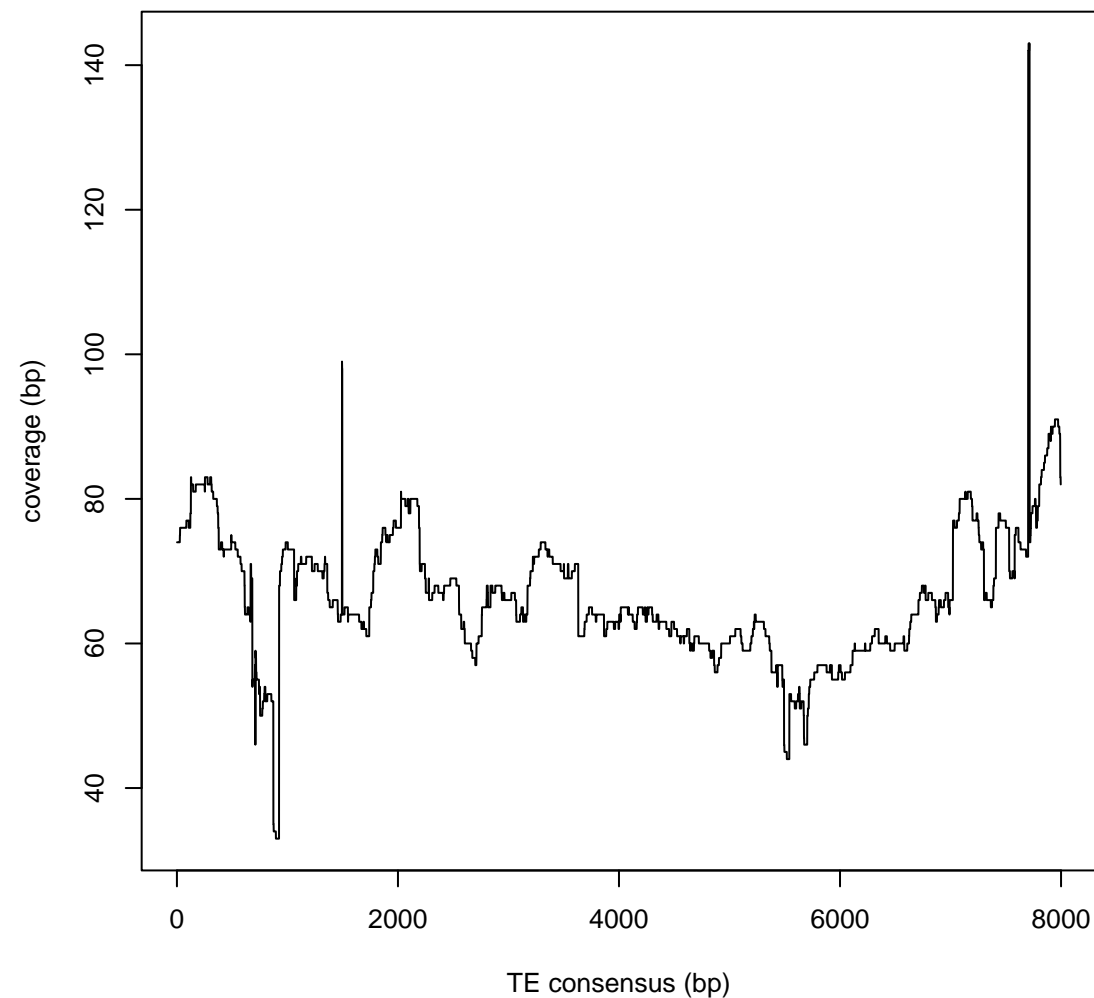


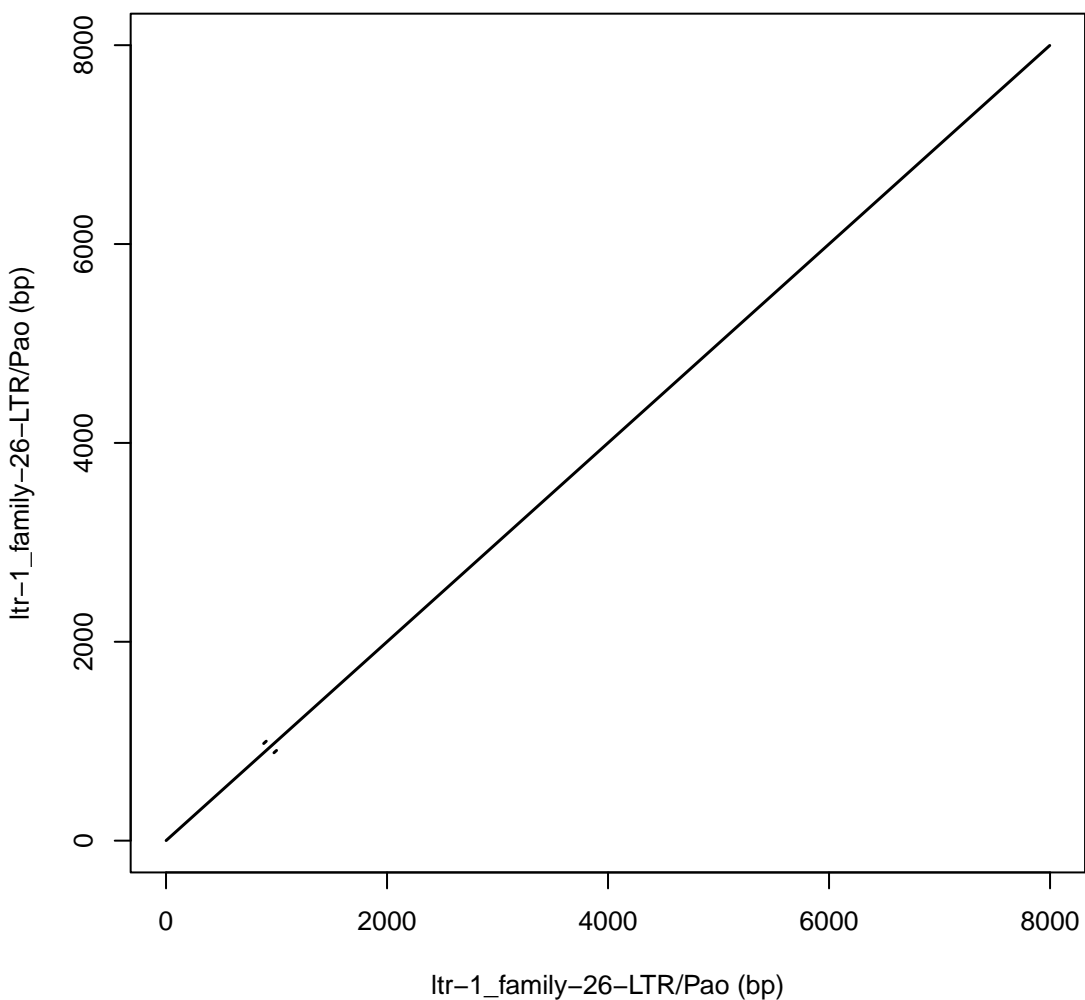
TE: ltr-1_family-26-LTR/Pao
consensus size: 7997bp; fragments: 546; full length: 1 (>=7197.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

