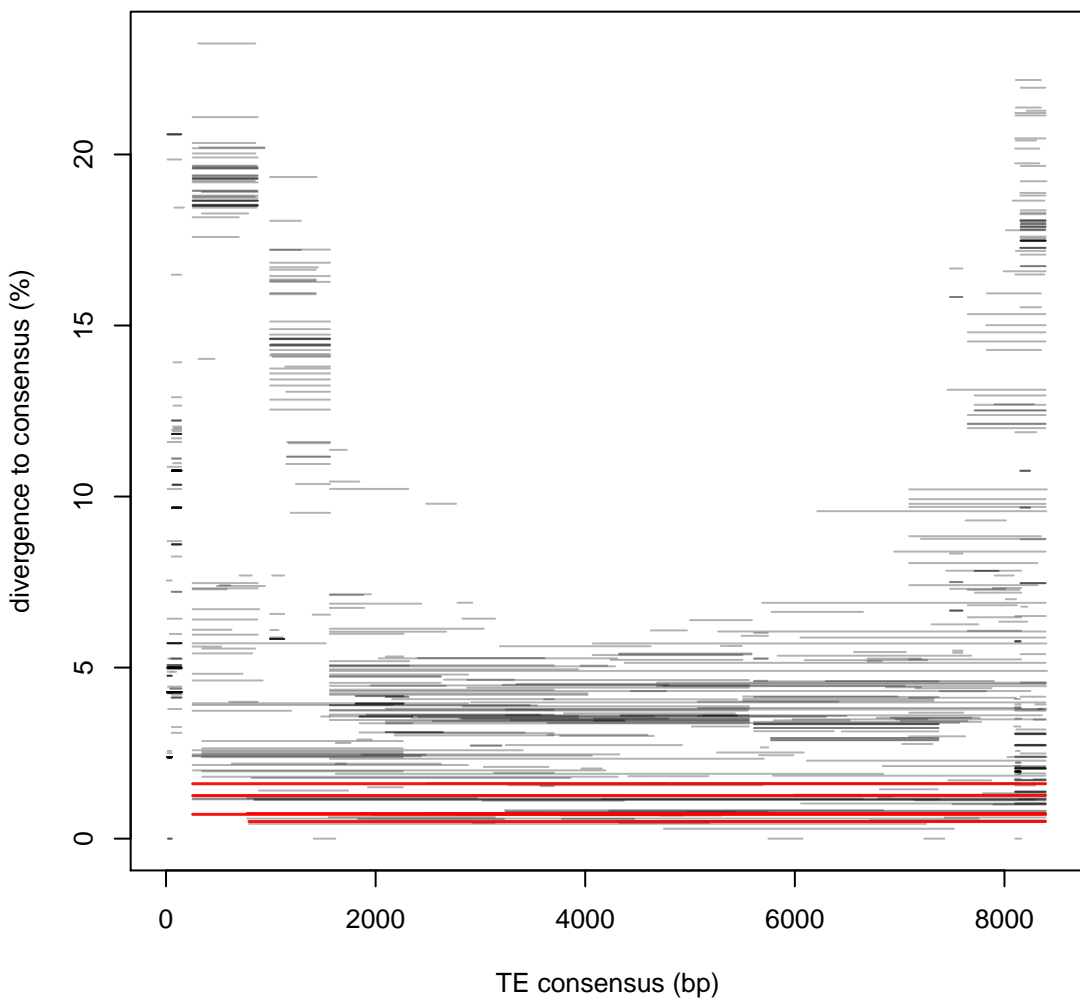
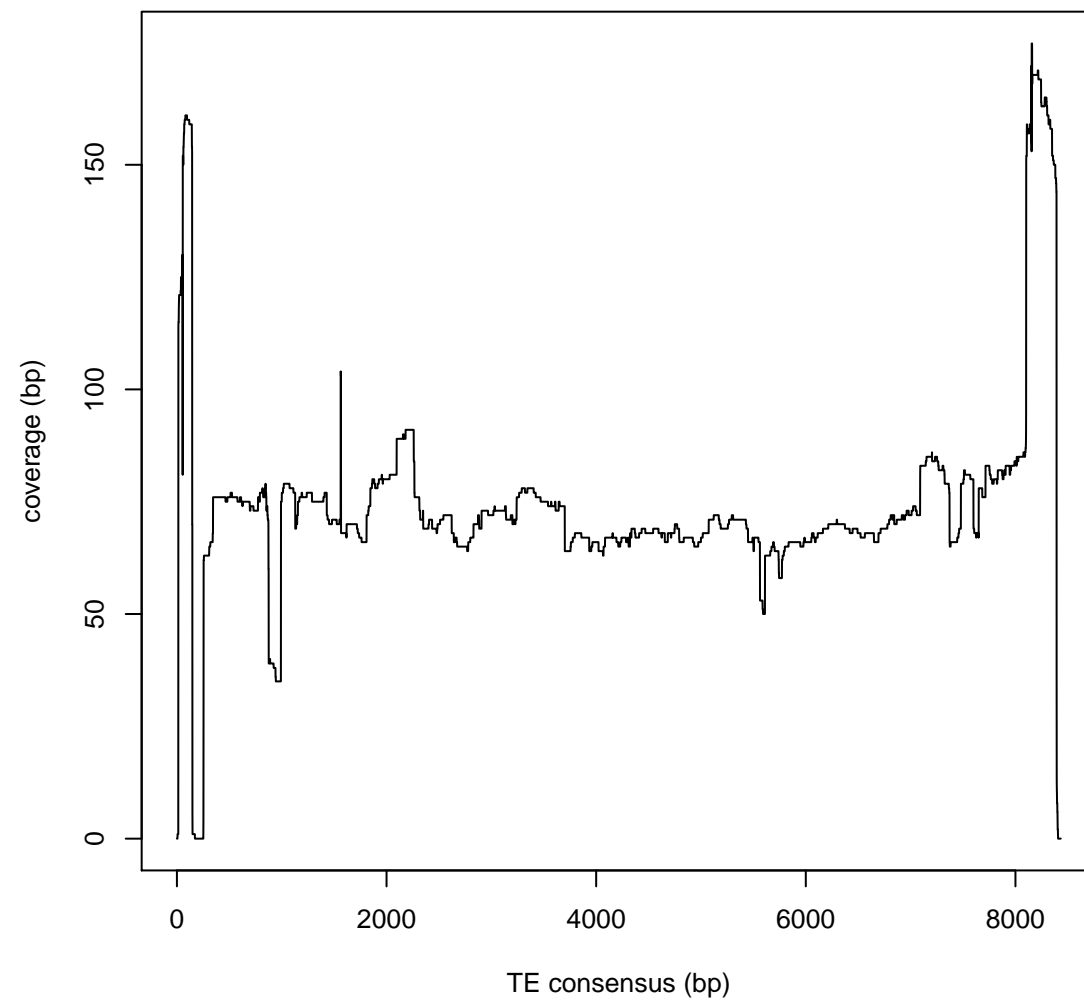


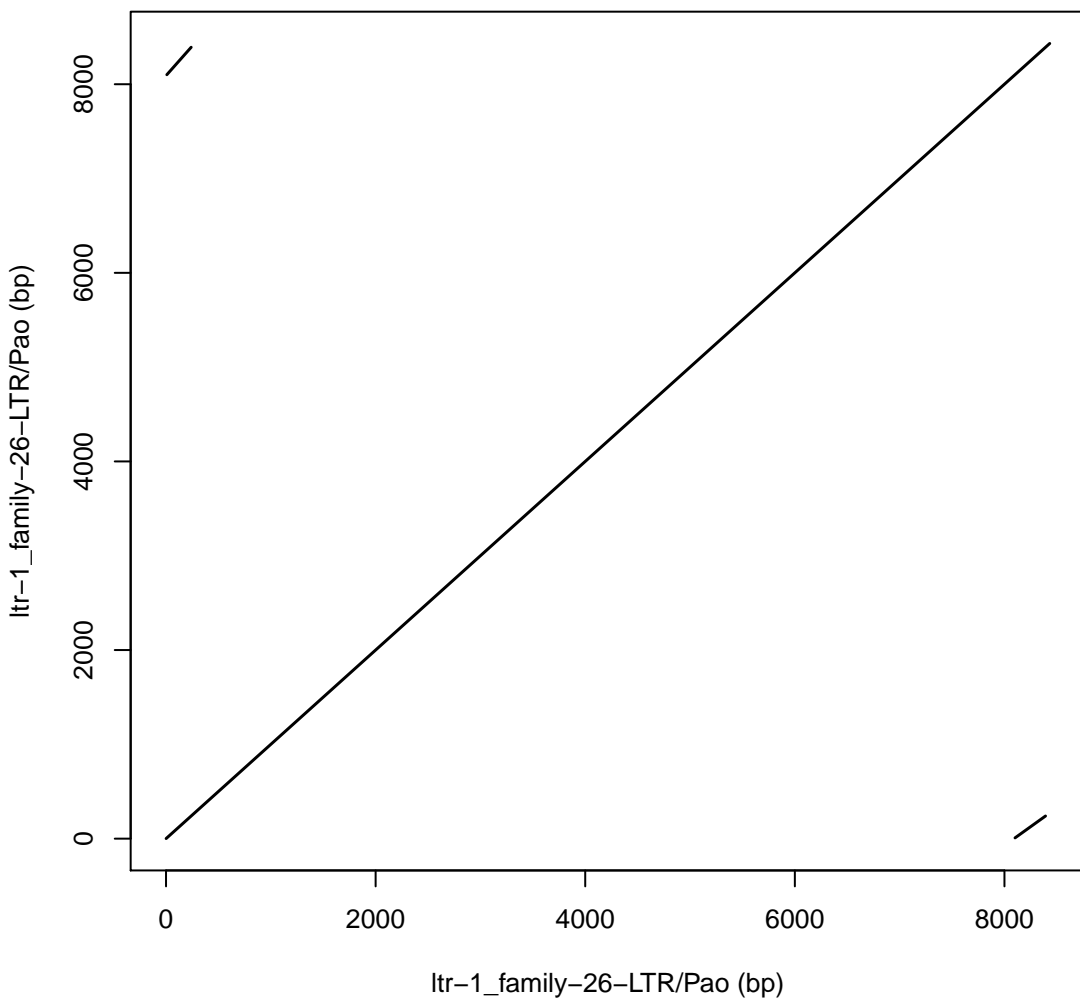
TE: ltr-1_family-26-LTR/Pao
consensus size: 8432bp; fragments: 764; full length: 5 (≥ 7588.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

