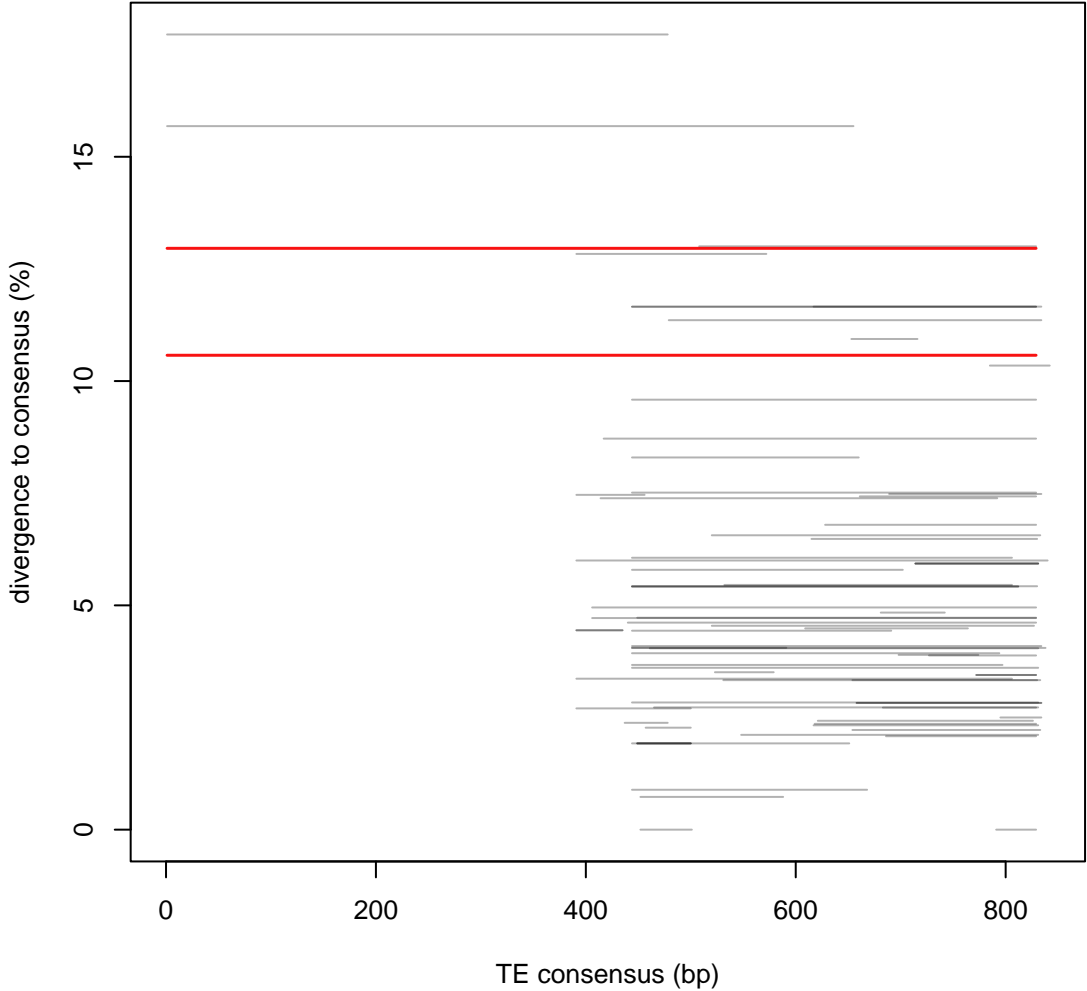
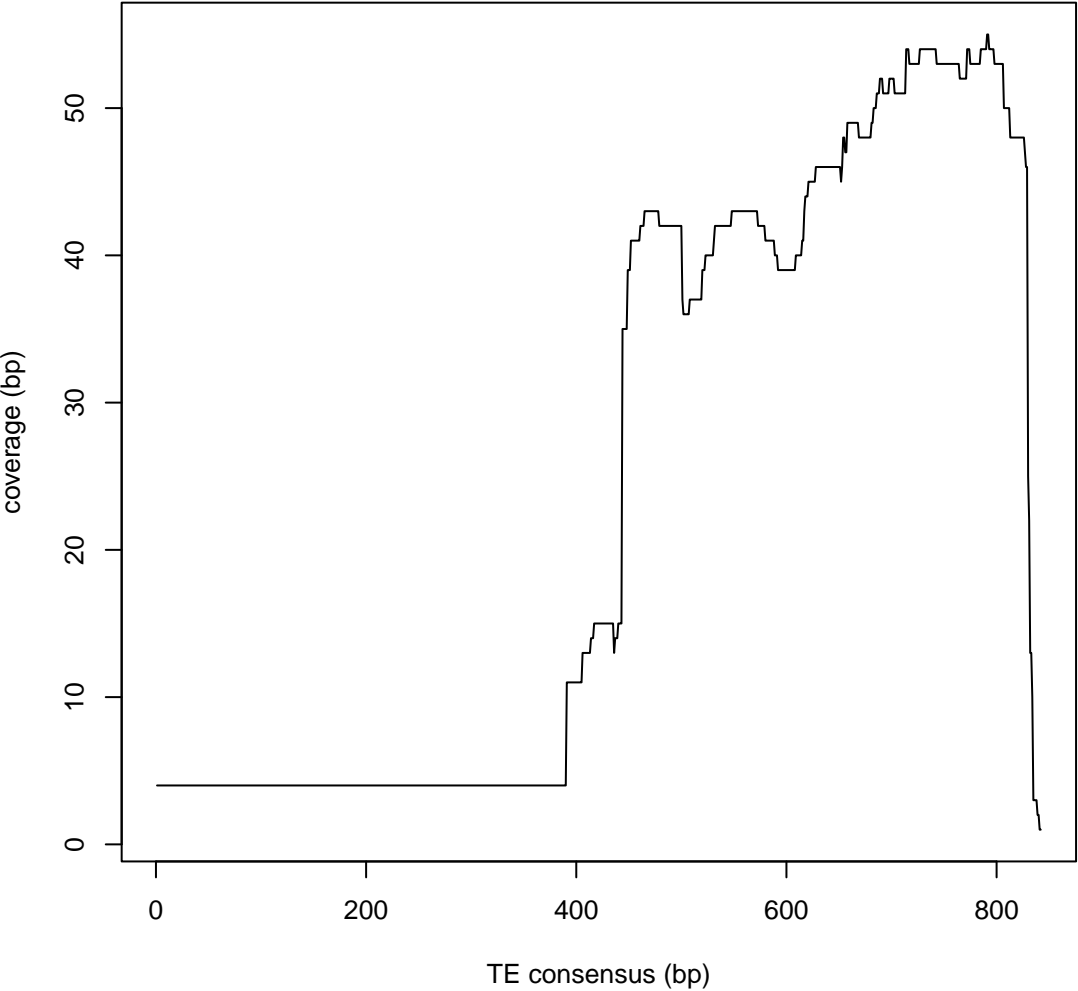


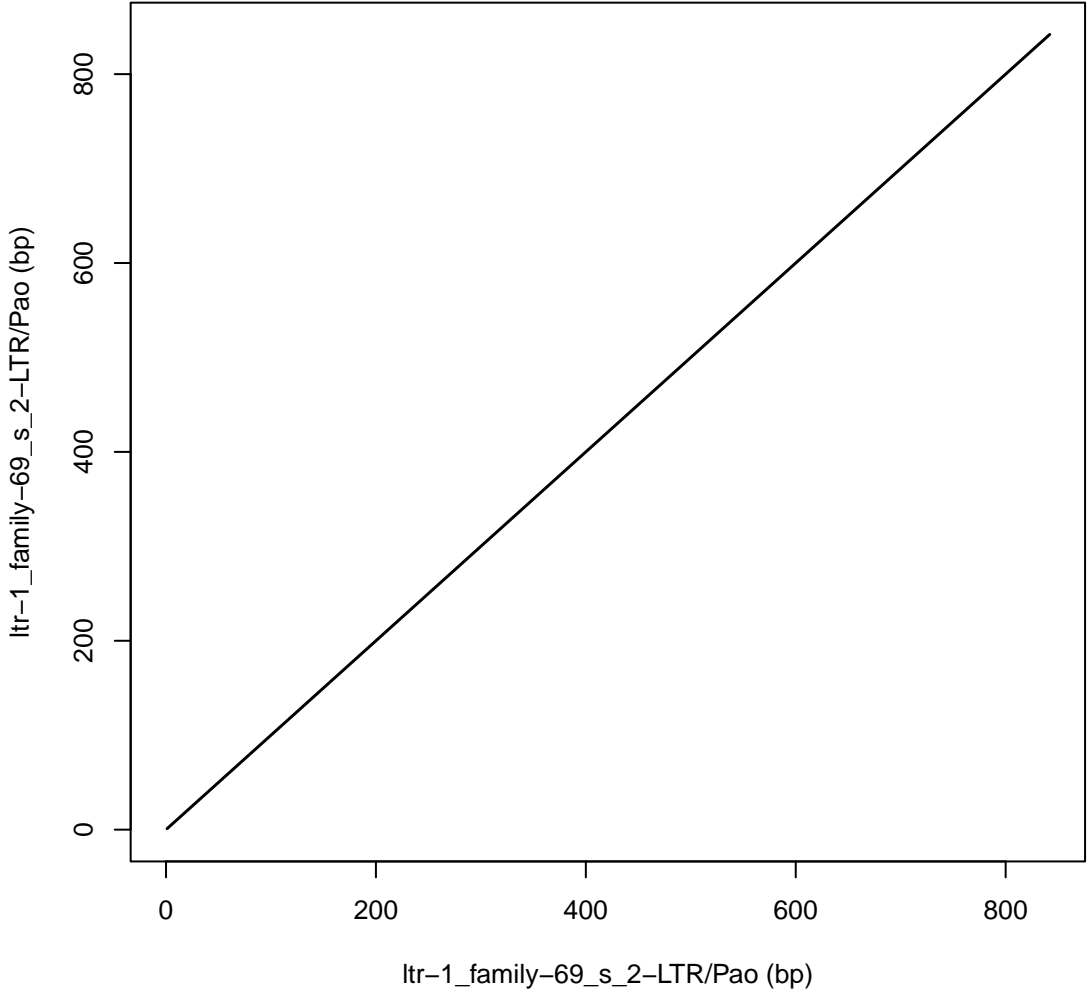
TE: ltr-1_family-69_s_2-LTR/Pao
consensus size: 842bp; fragments: 81; full length: 2 (≥ 757.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

