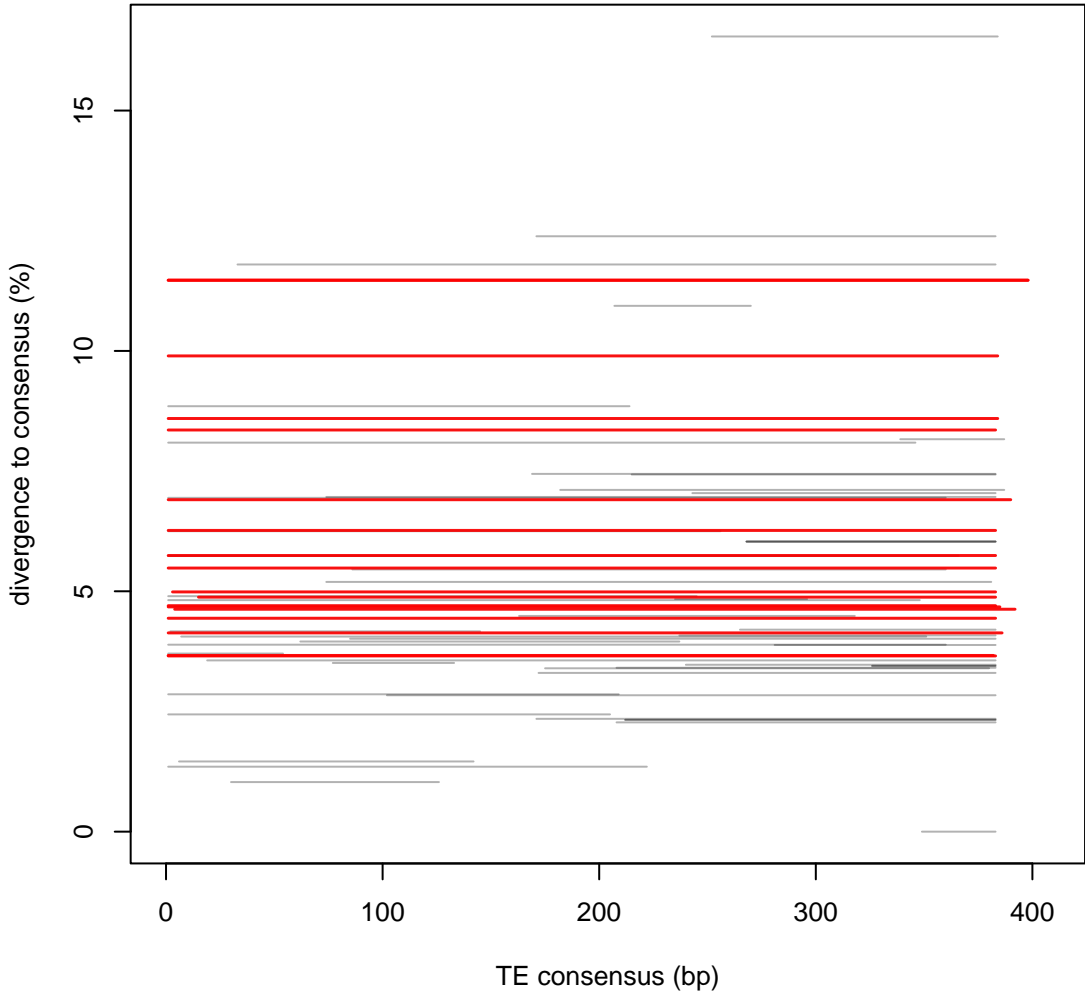
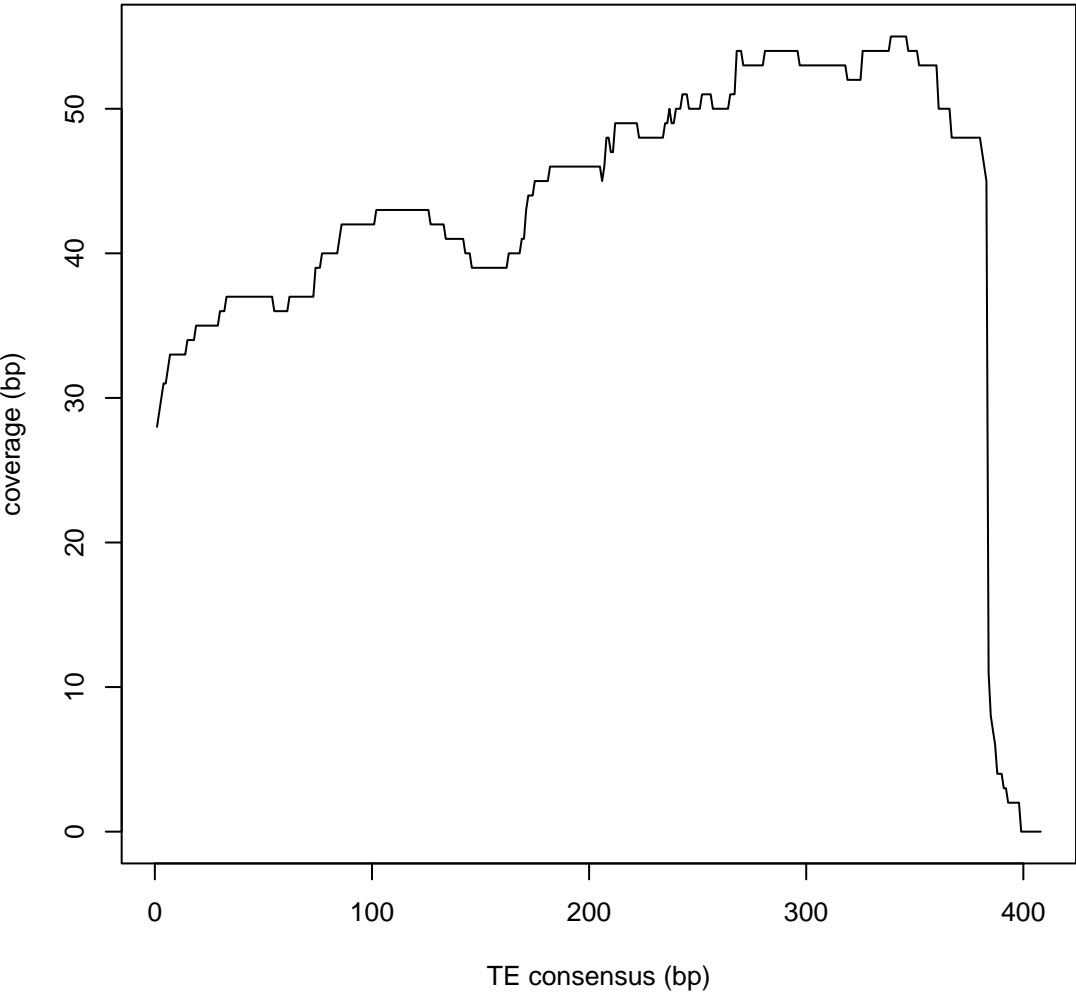


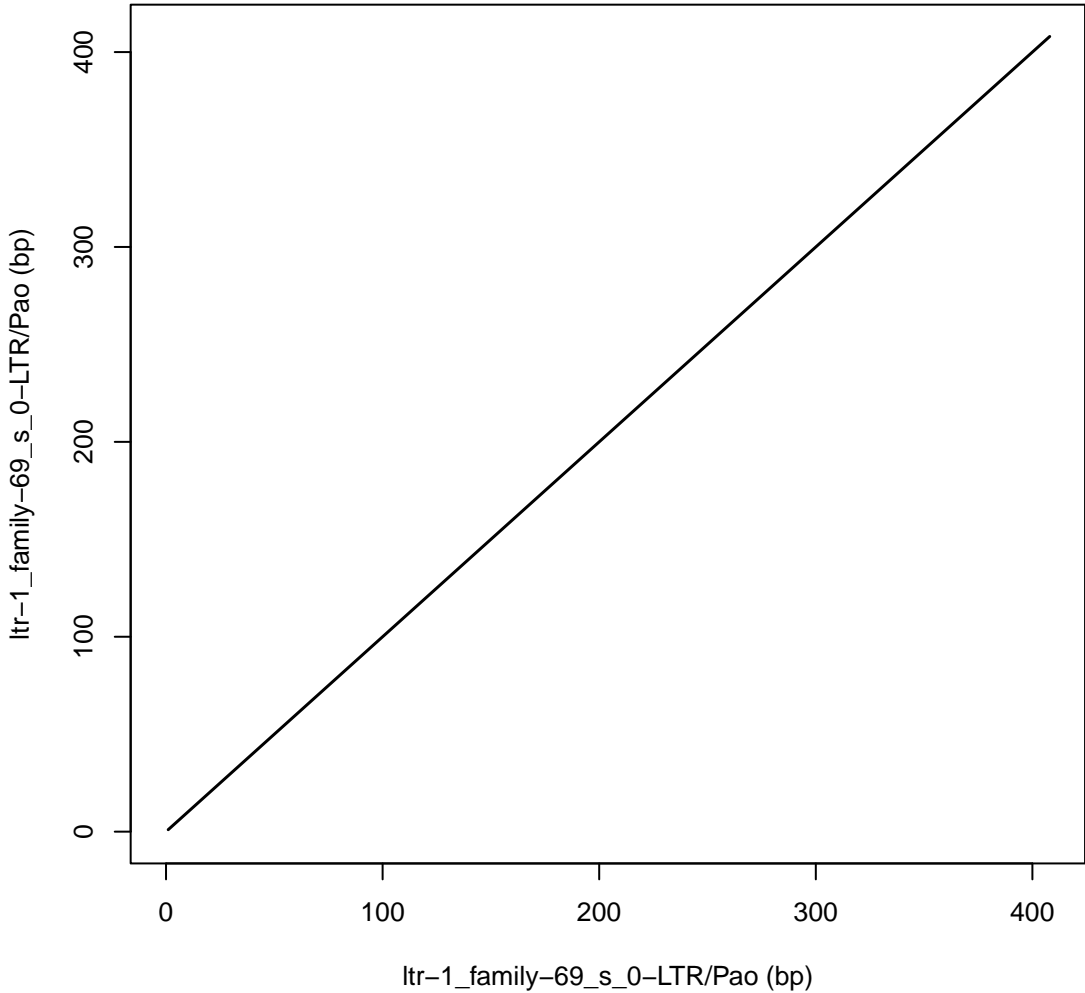
TE: ltr-1_family-69_s_0-LTR/Pao
consensus size: 408bp; fragments: 71; full length: 18 (>=367.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

