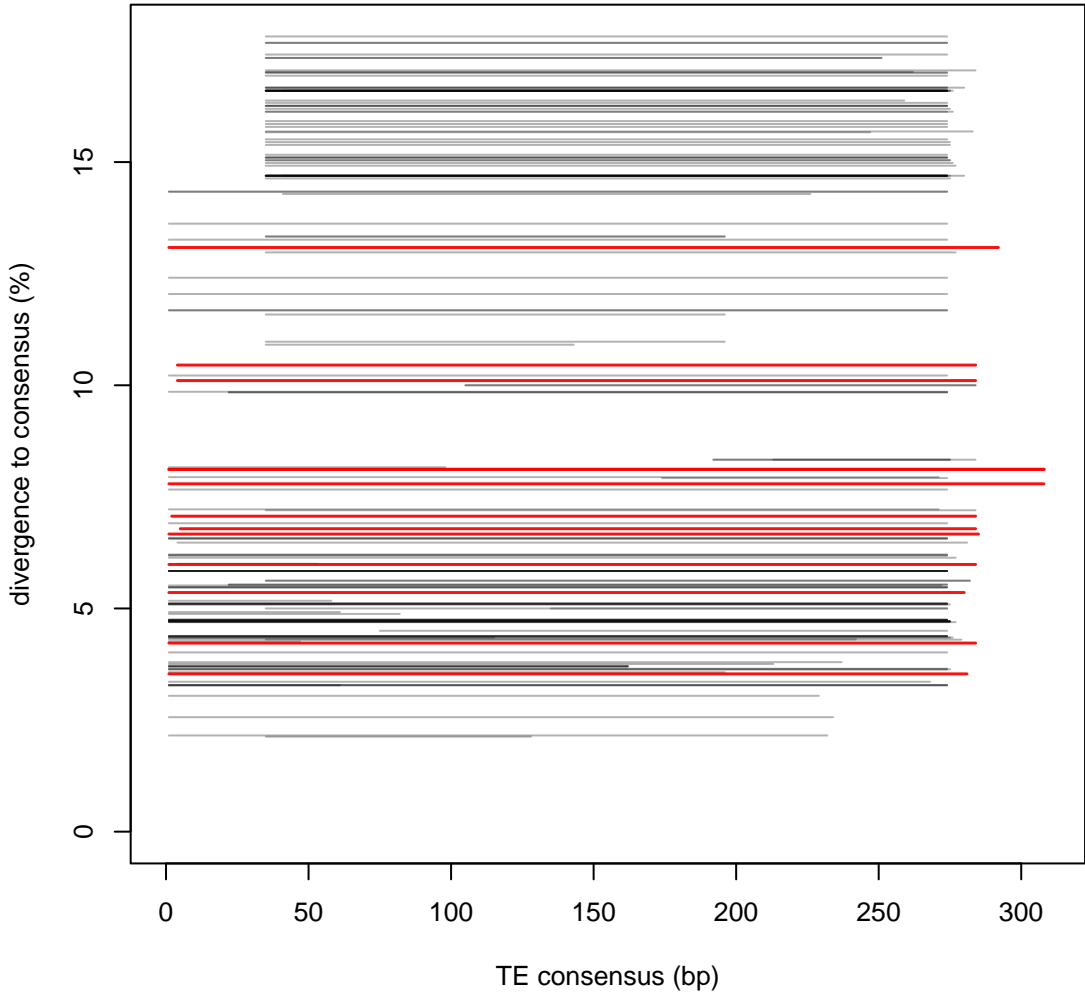
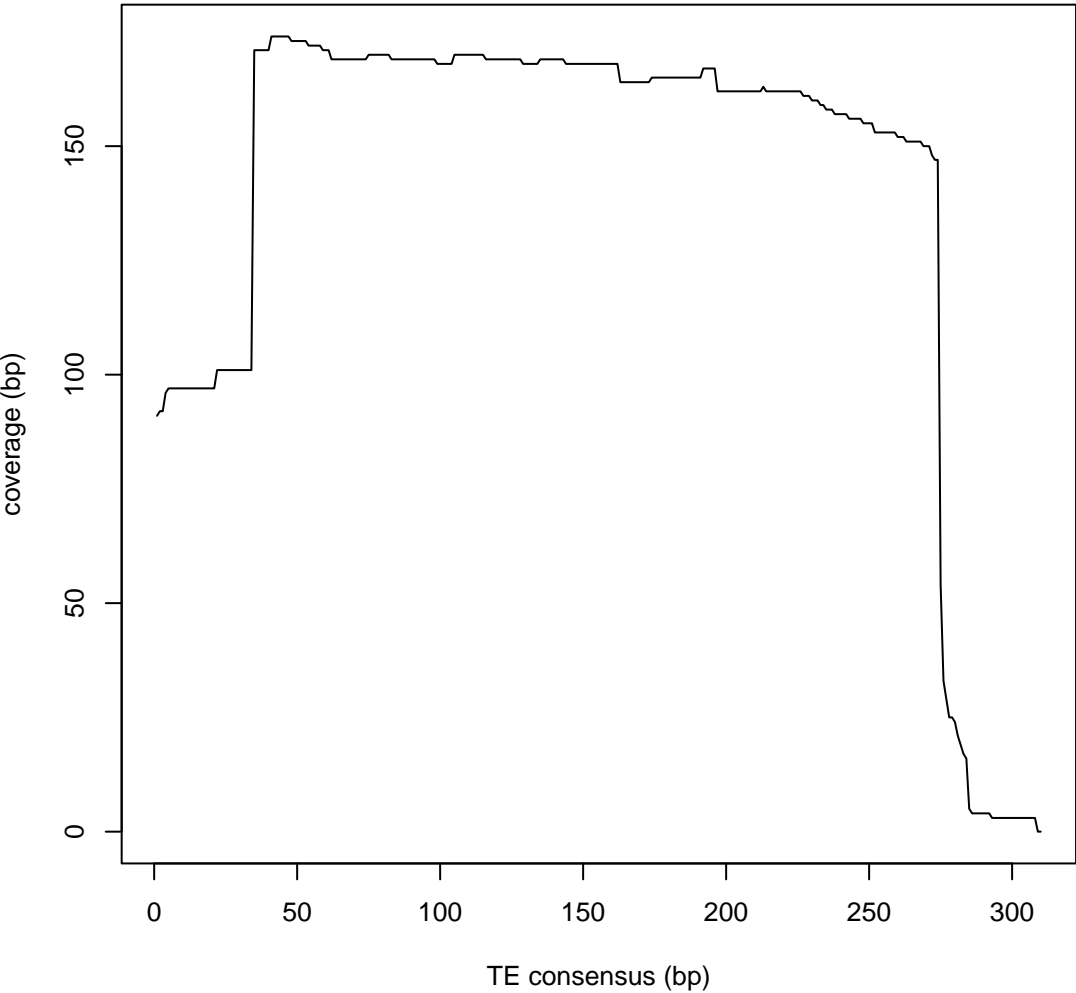


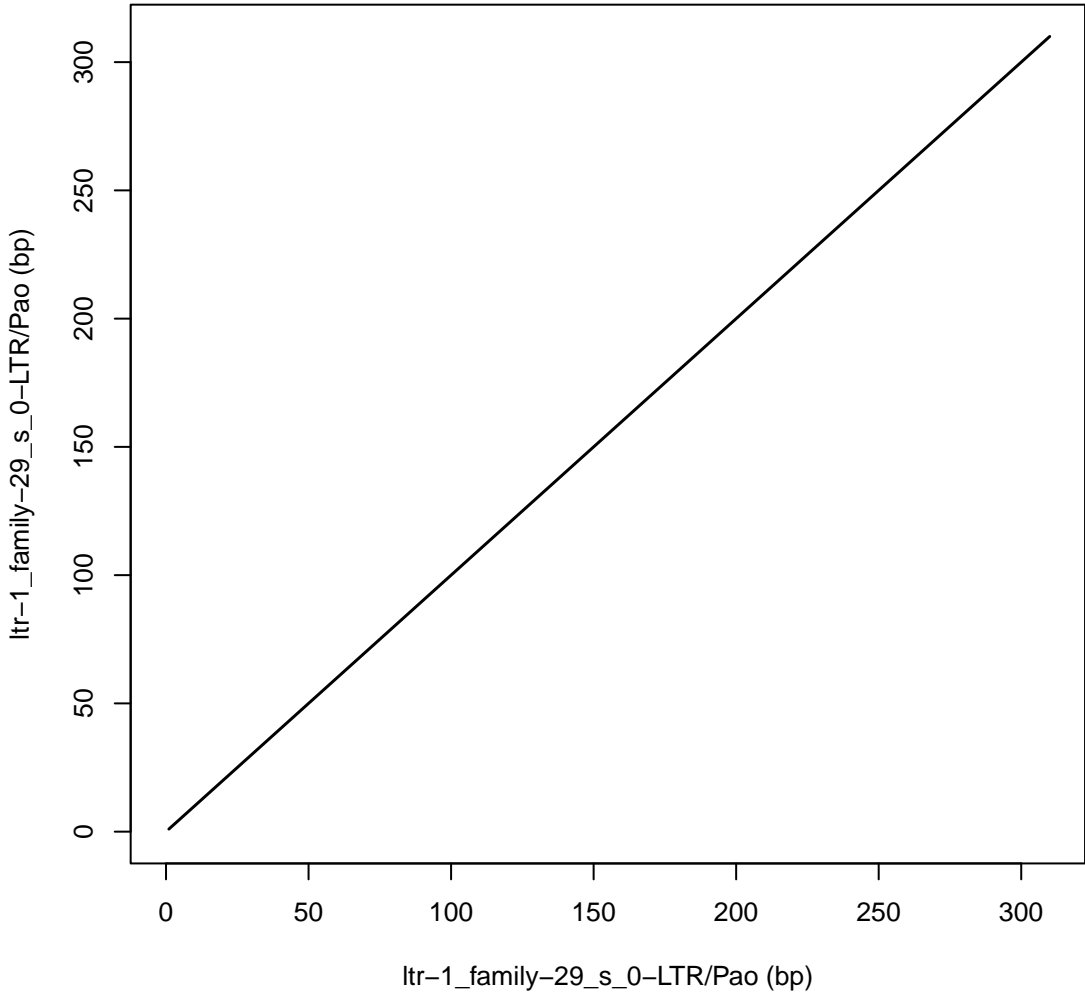
TE: ltr-1_family-29_s_0-LTR/Pao
consensus size: 310bp; fragments: 182; full length: 13 (>=279bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

