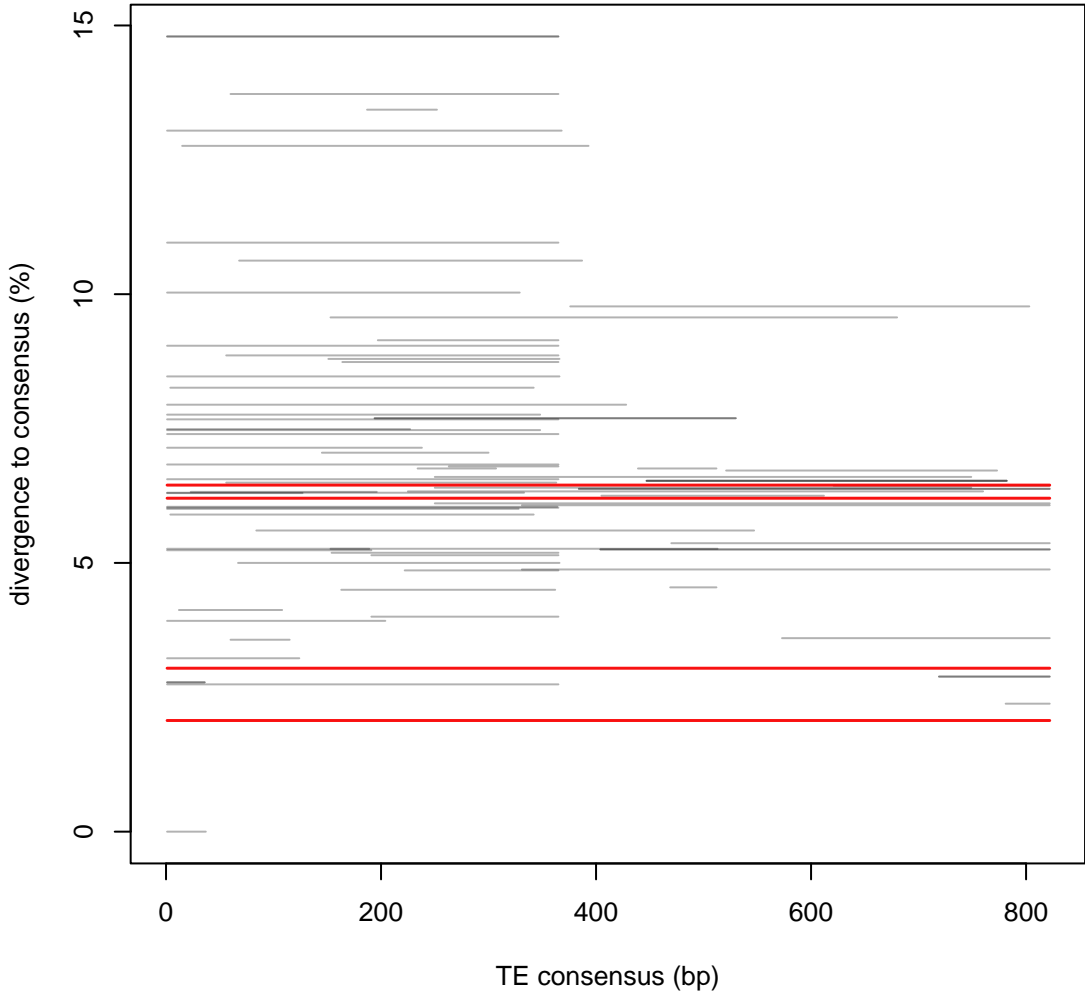
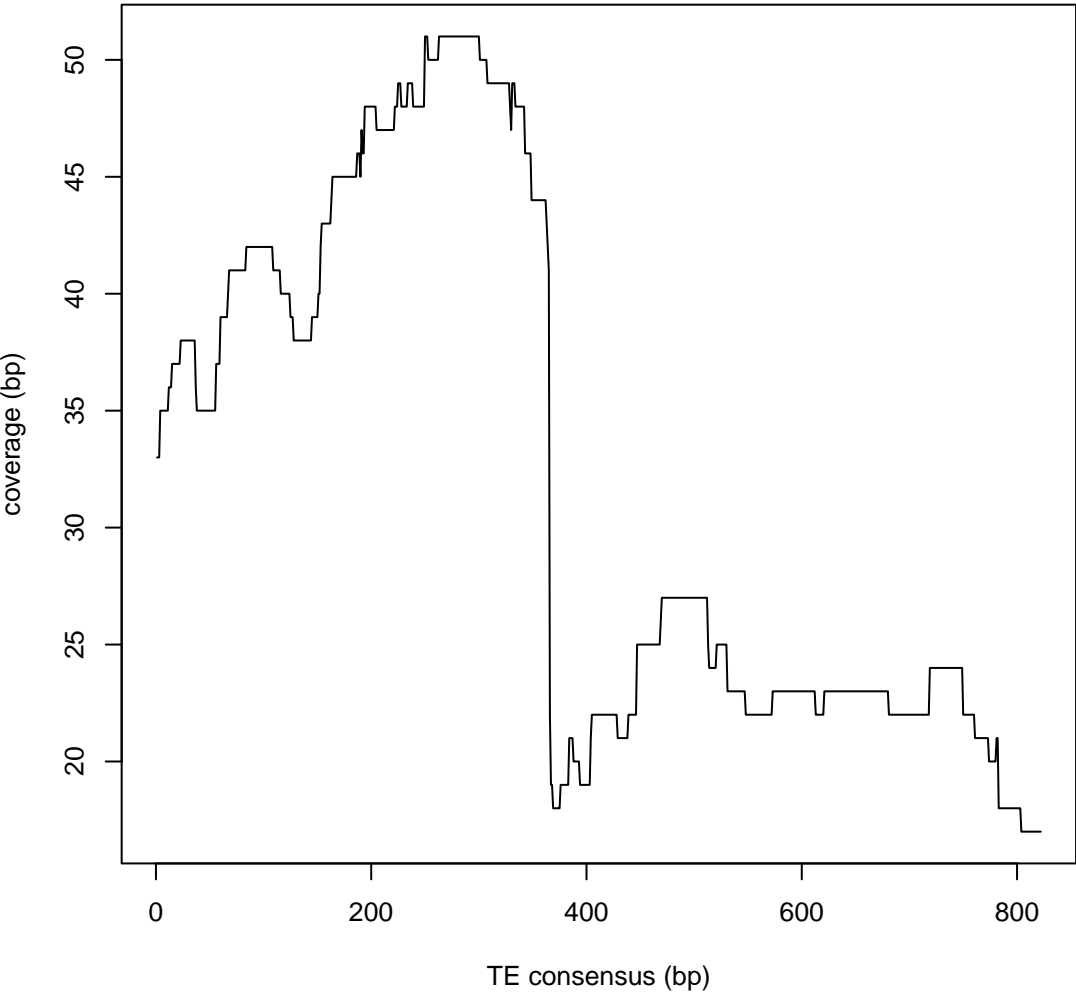


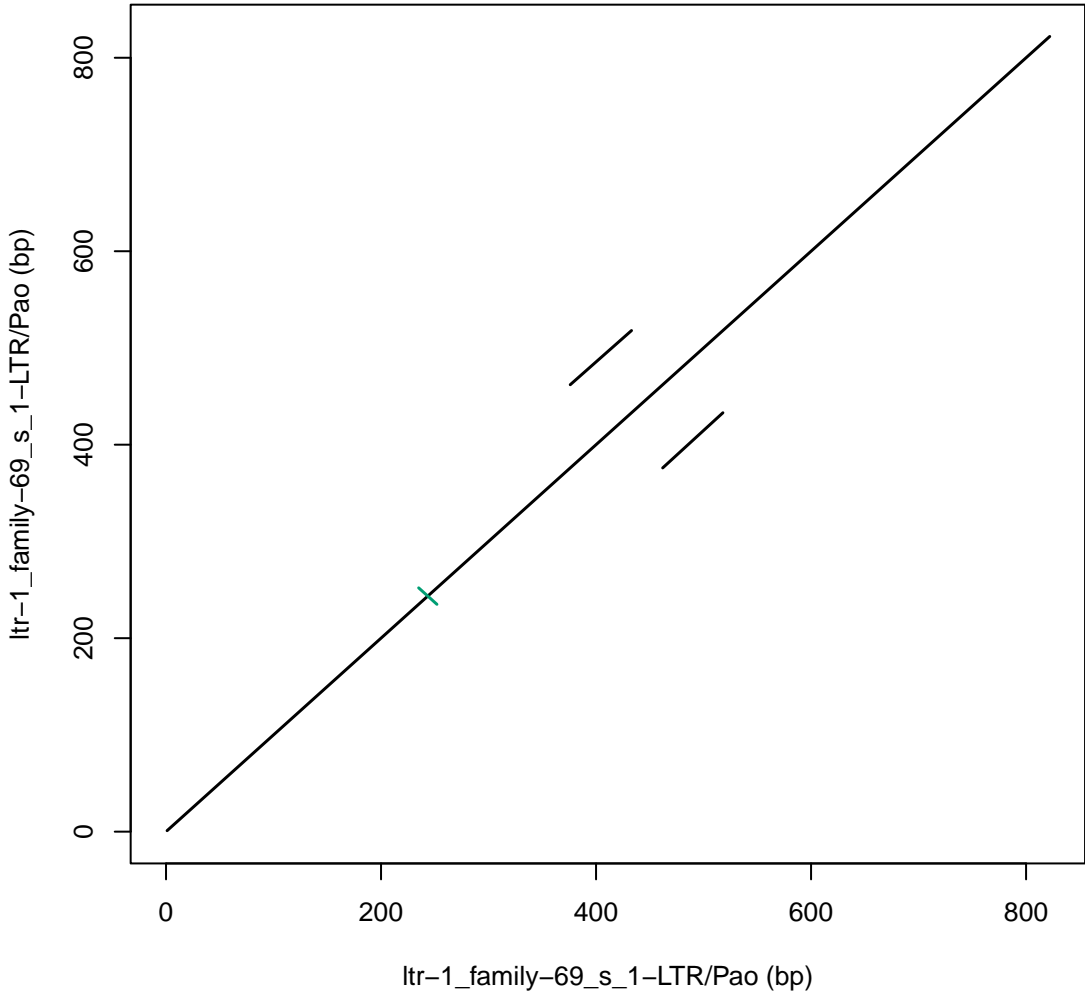
TE: ltr-1_family-69_s_1-LTR/Pao
consensus size: 822bp; fragments: 85; full length: 4 (>=739.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

