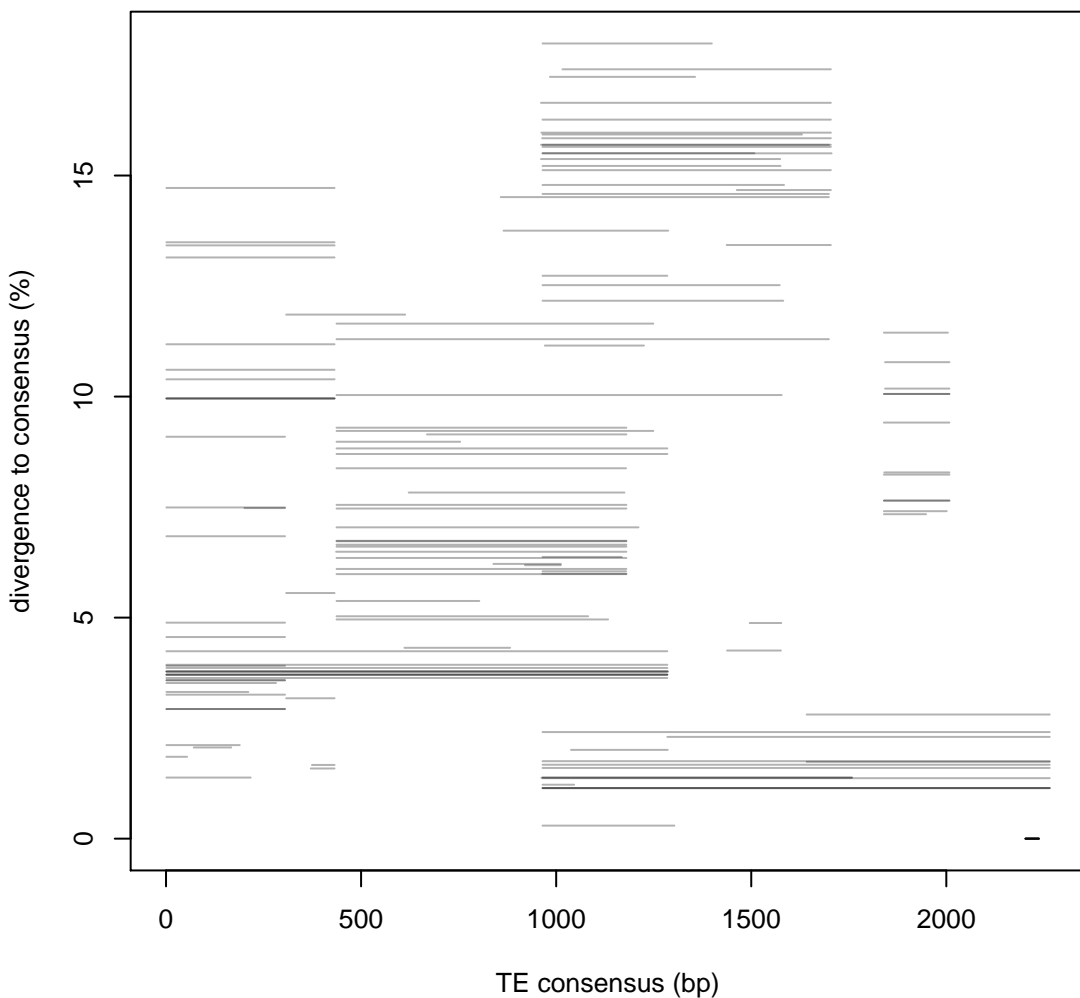
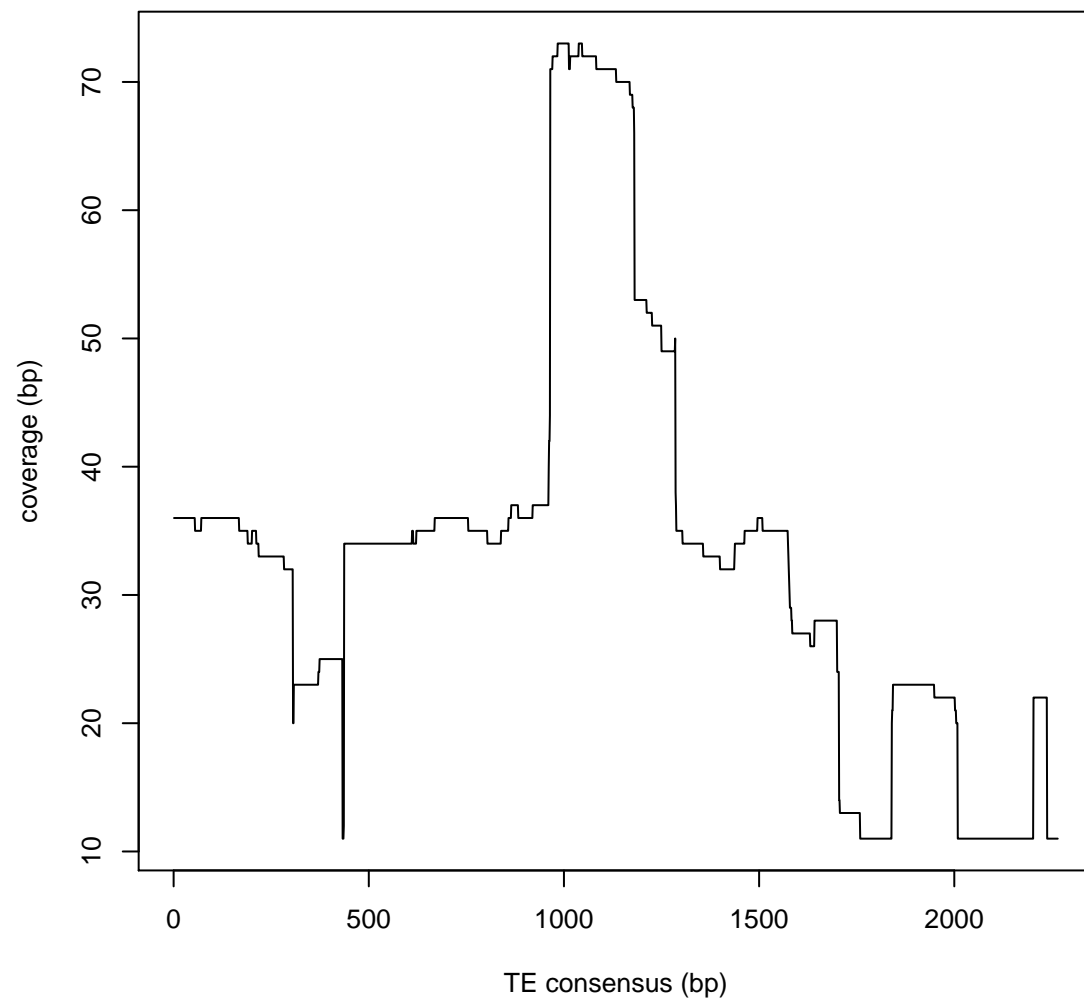


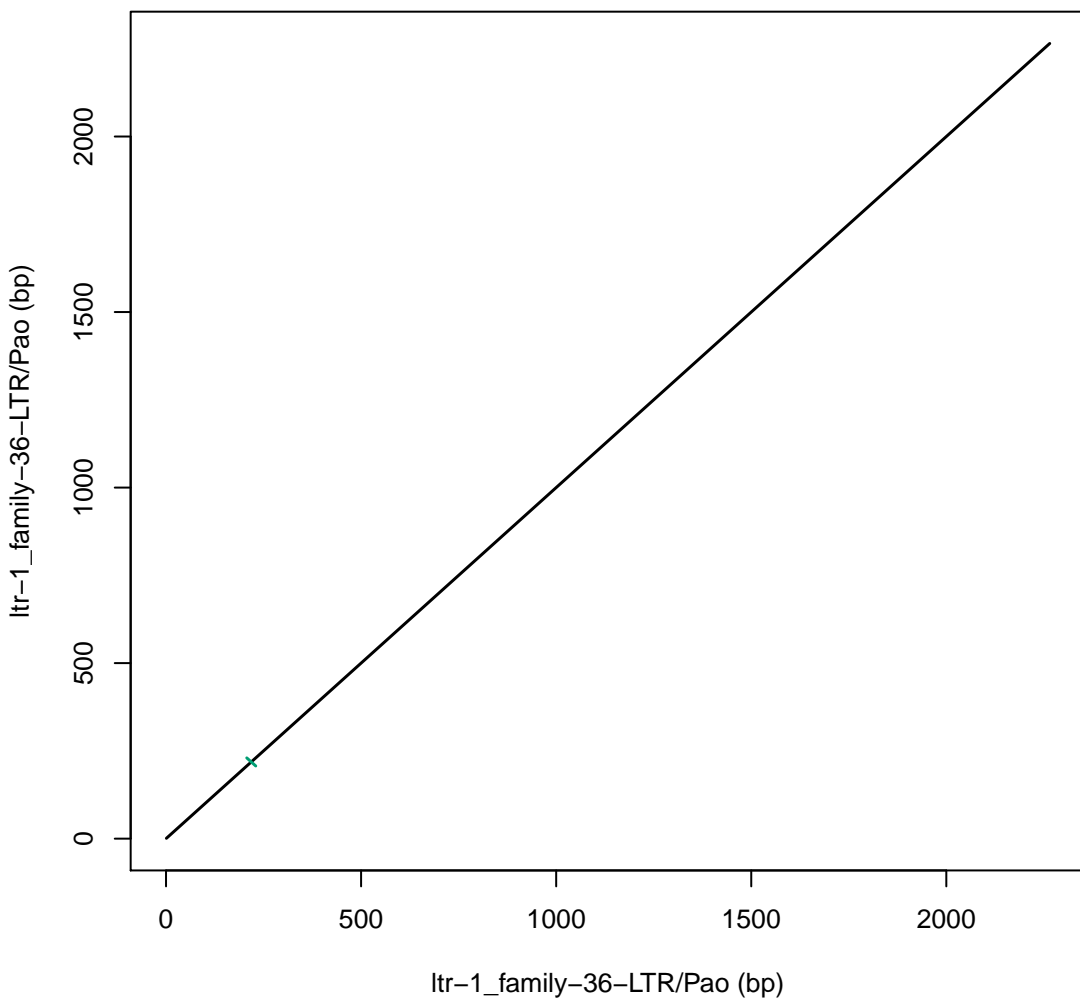
TE: ltr-1_family-36-LTR/Pao
consensus size: 2265bp; fragments: 141; full length: 0 (>=2038.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

