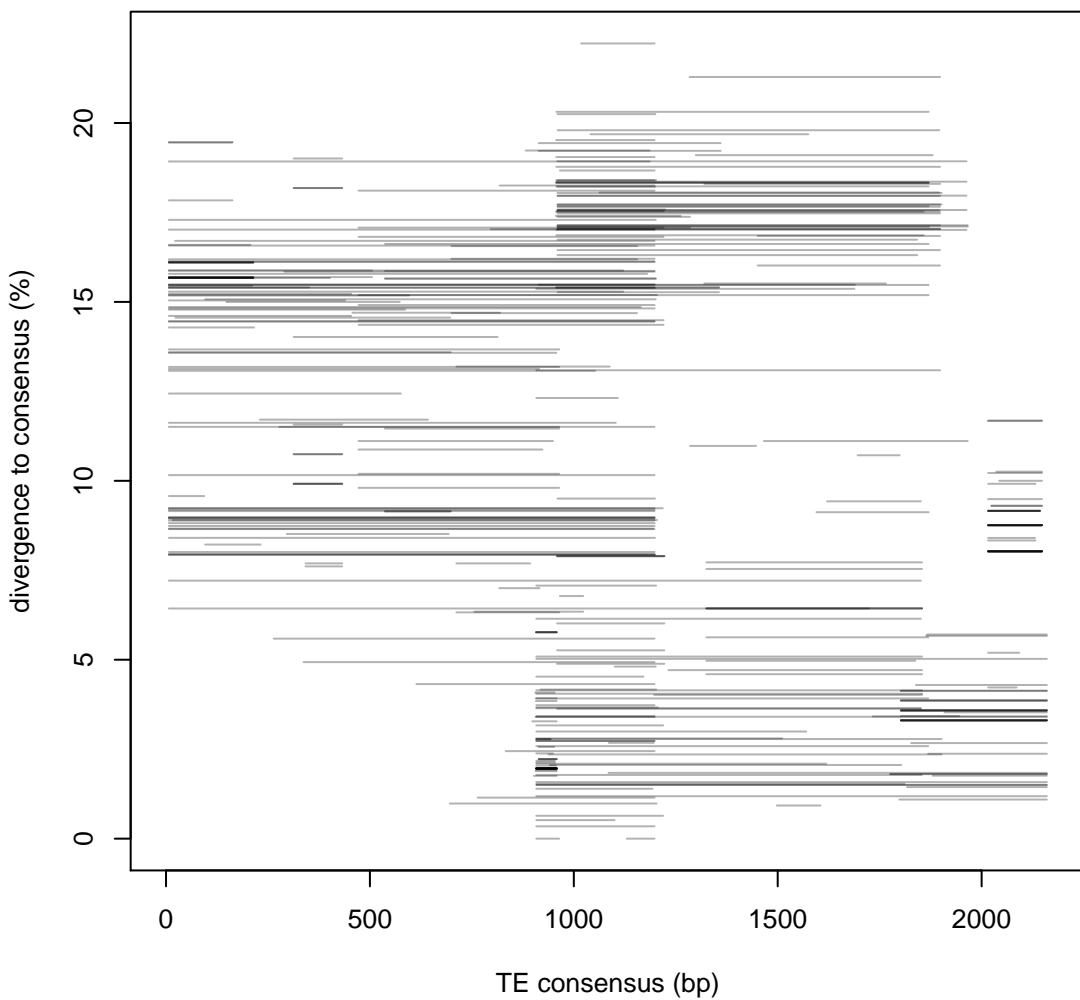
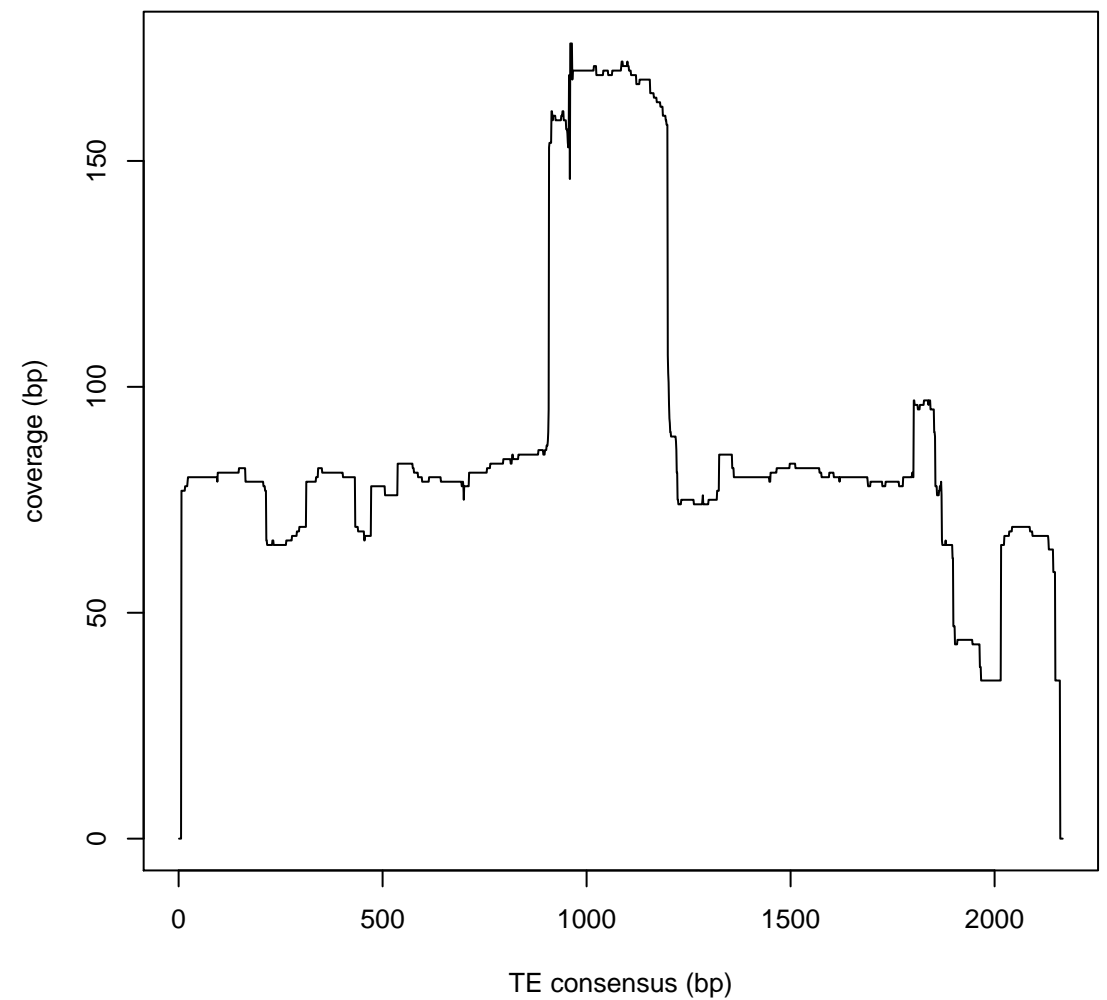


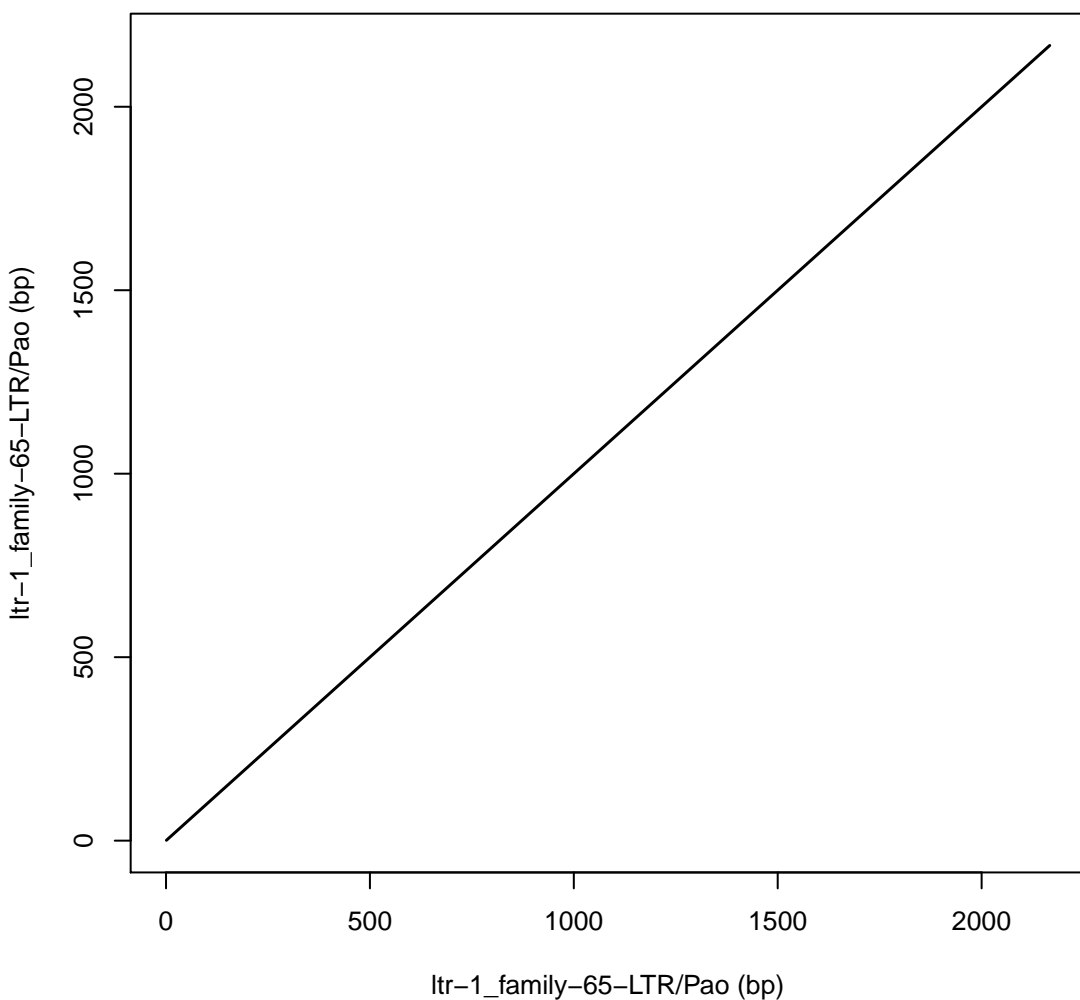
TE: ltr-1_family-65-LTR/Pao
consensus size: 2167bp; fragments: 362; full length: 0 (≥ 1950.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

