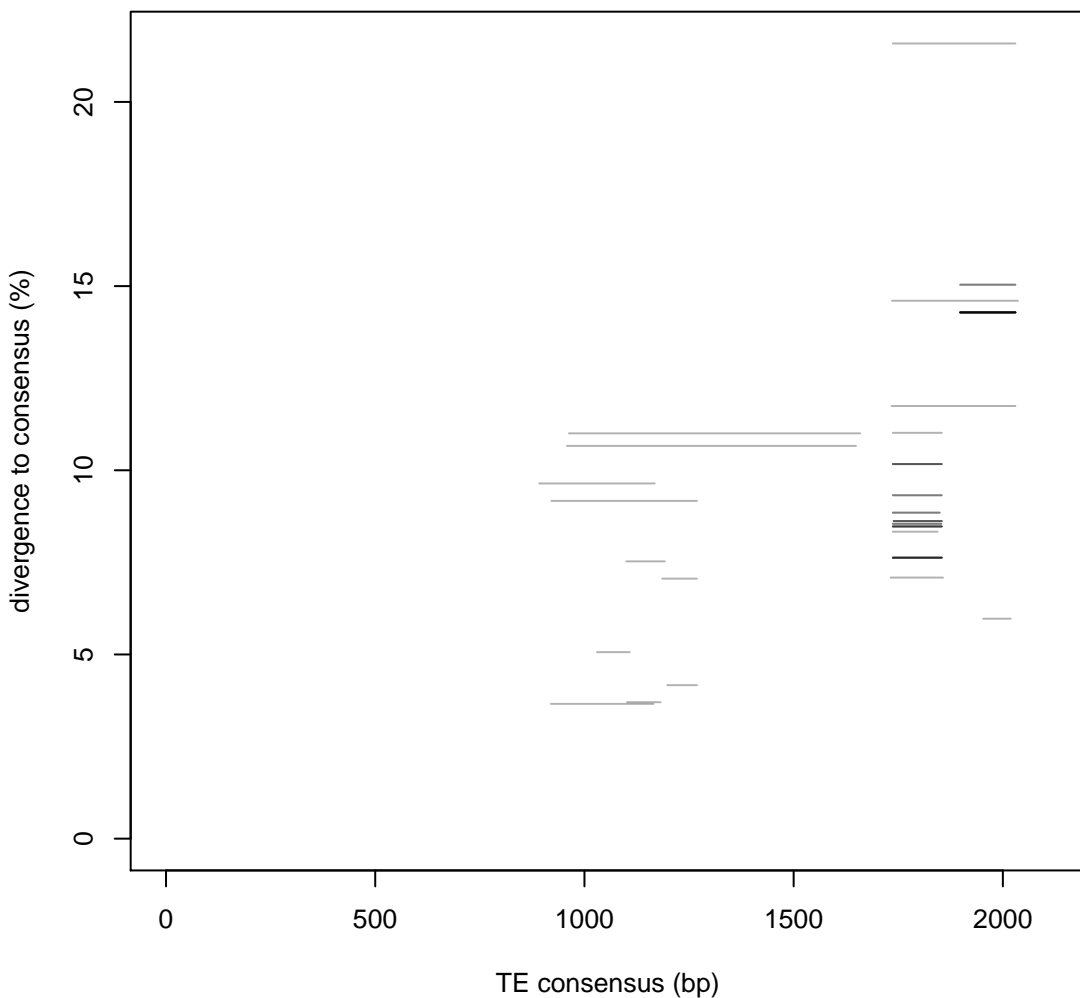
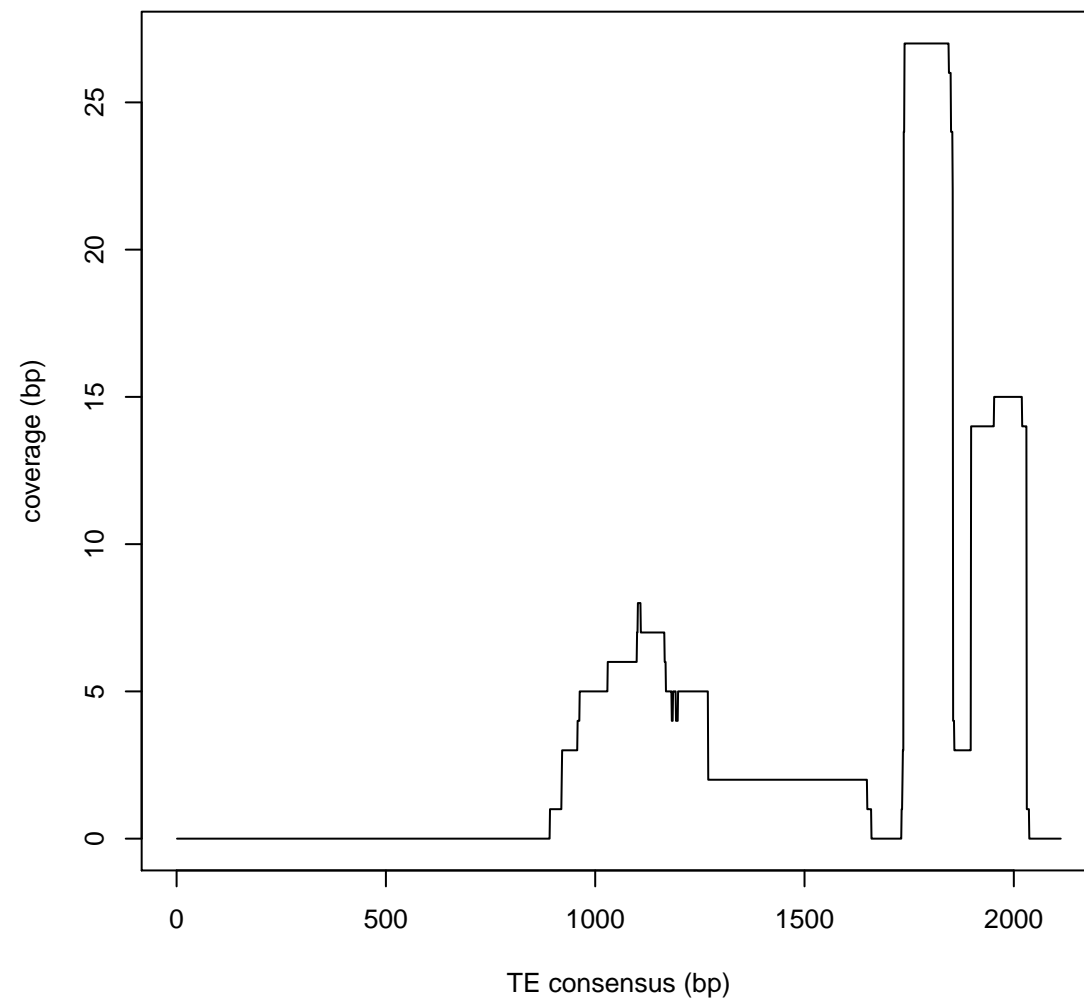


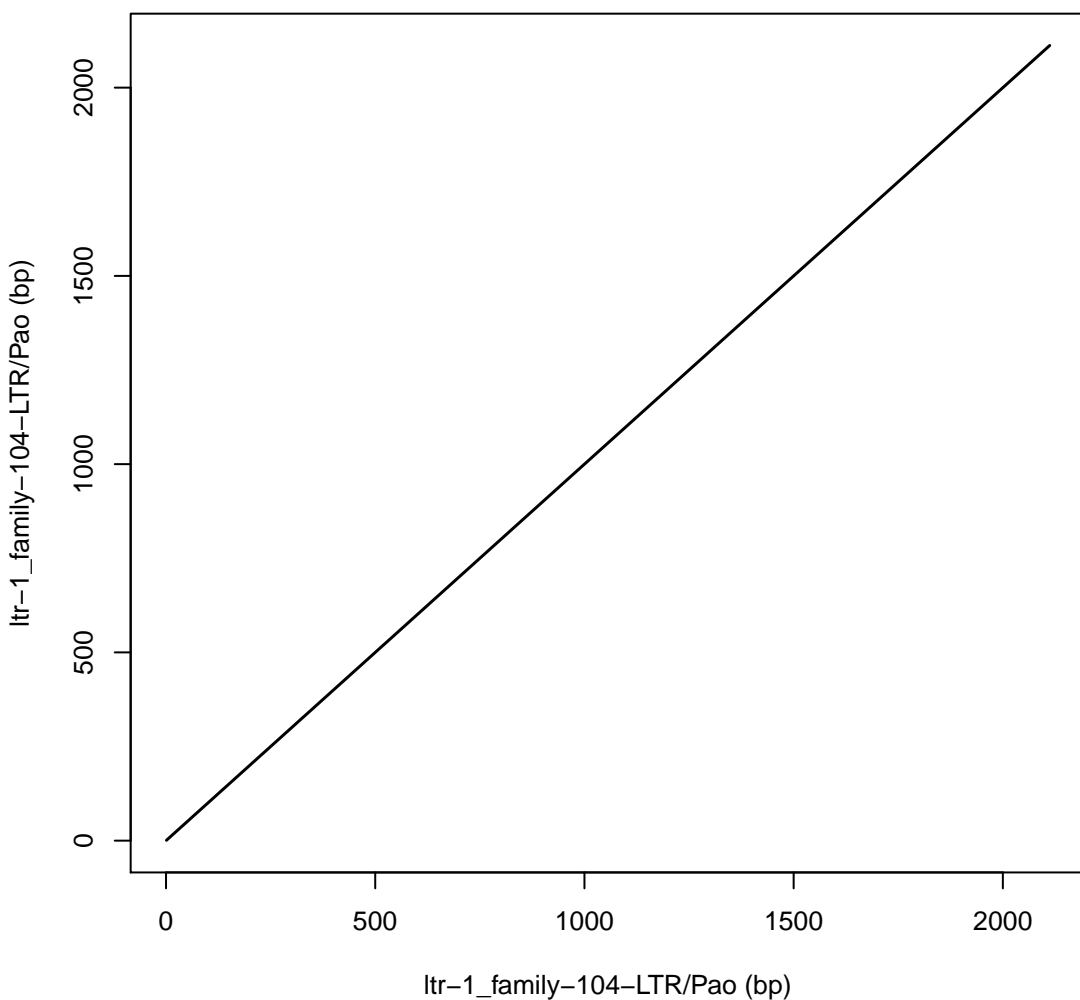
TE: ltr-1_family-104-LTR/Pao
consensus size: 2112bp; fragments: 49; full length: 0 (≥ 1900.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

no orf >400 bp detected

