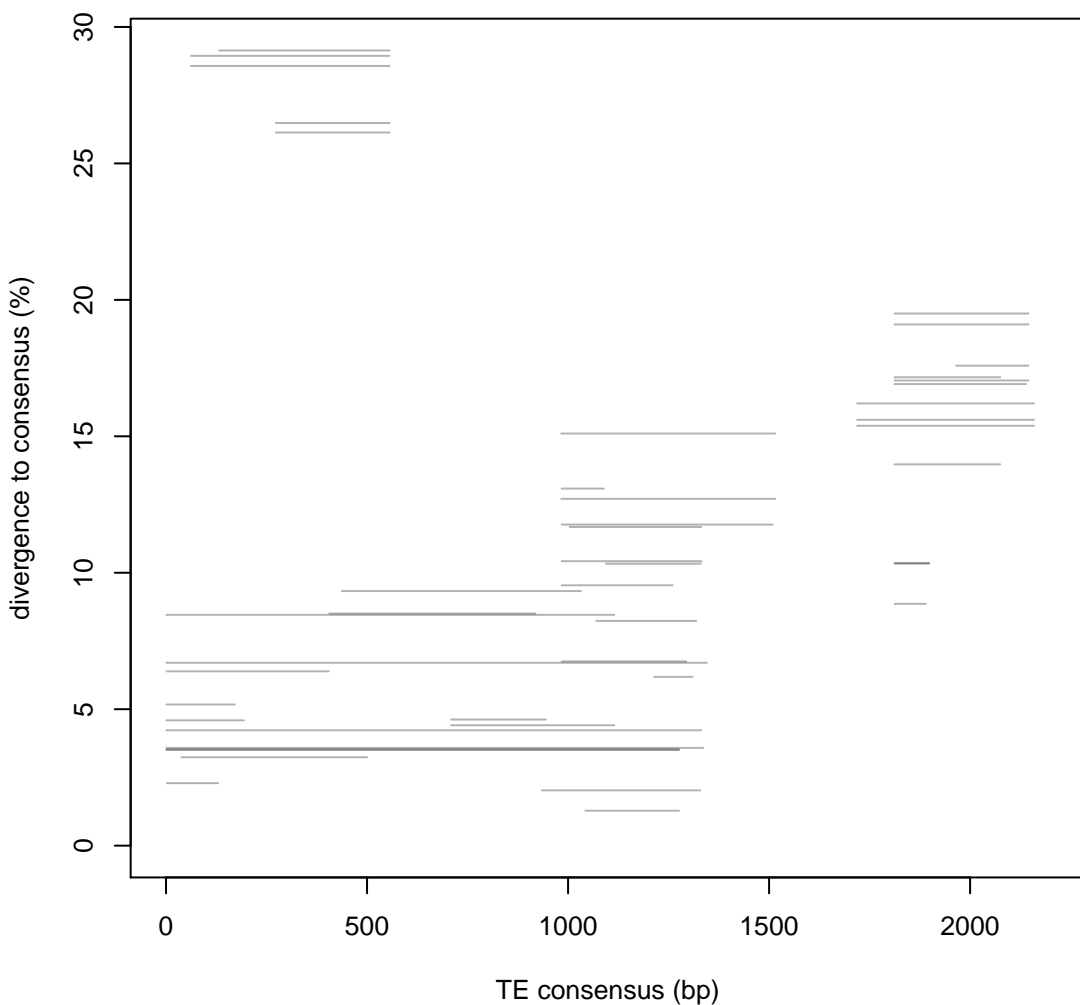
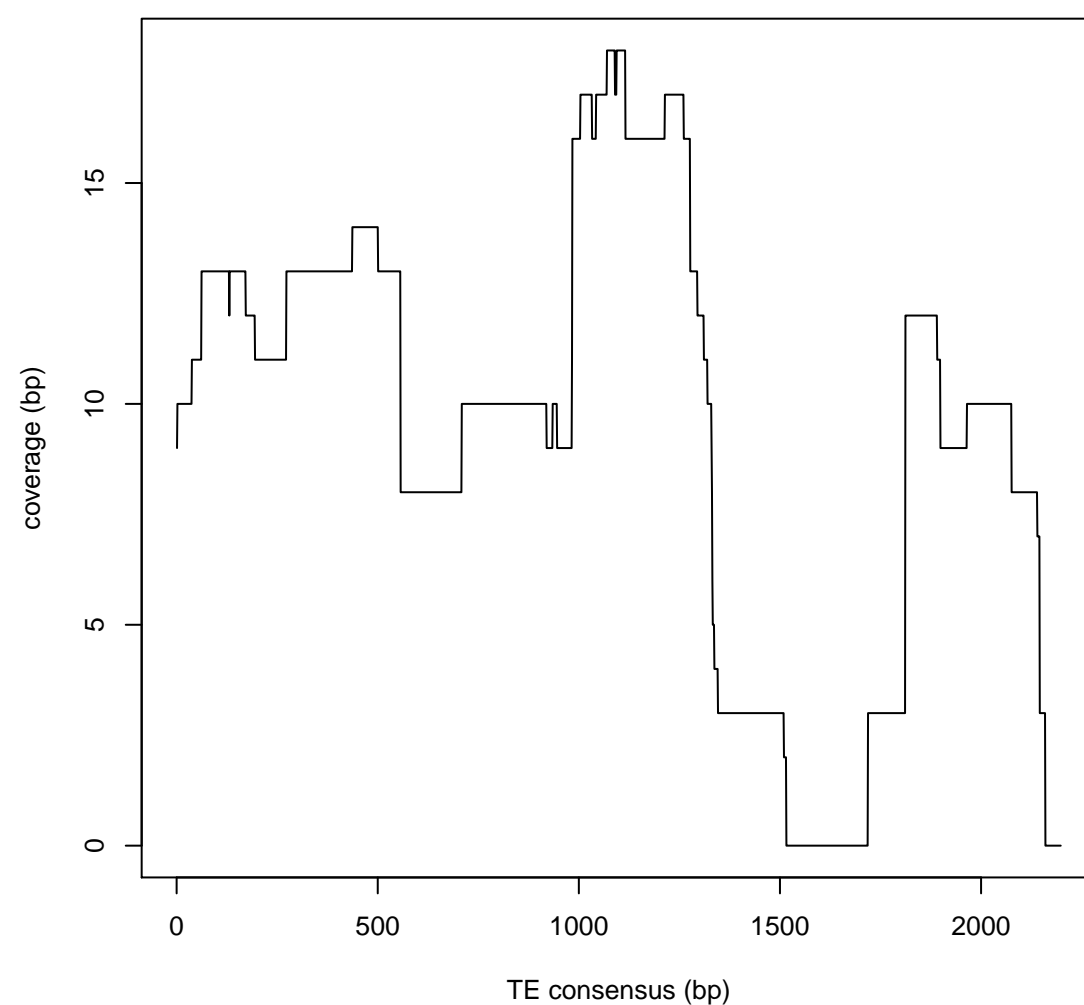


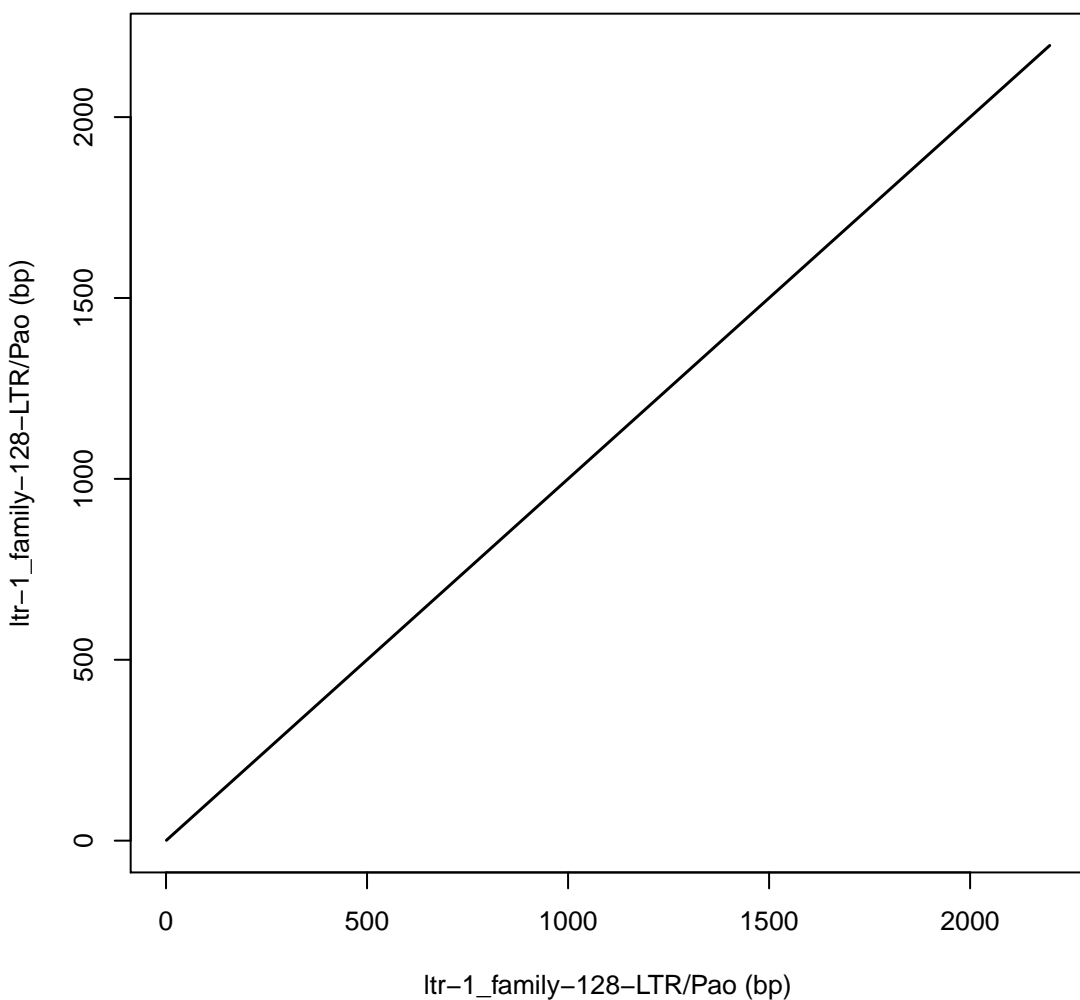
TE: ltr-1_family-128-LTR/Pao
consensus size: 2198bp; fragments: 46; full length: 0 (>=1978.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

