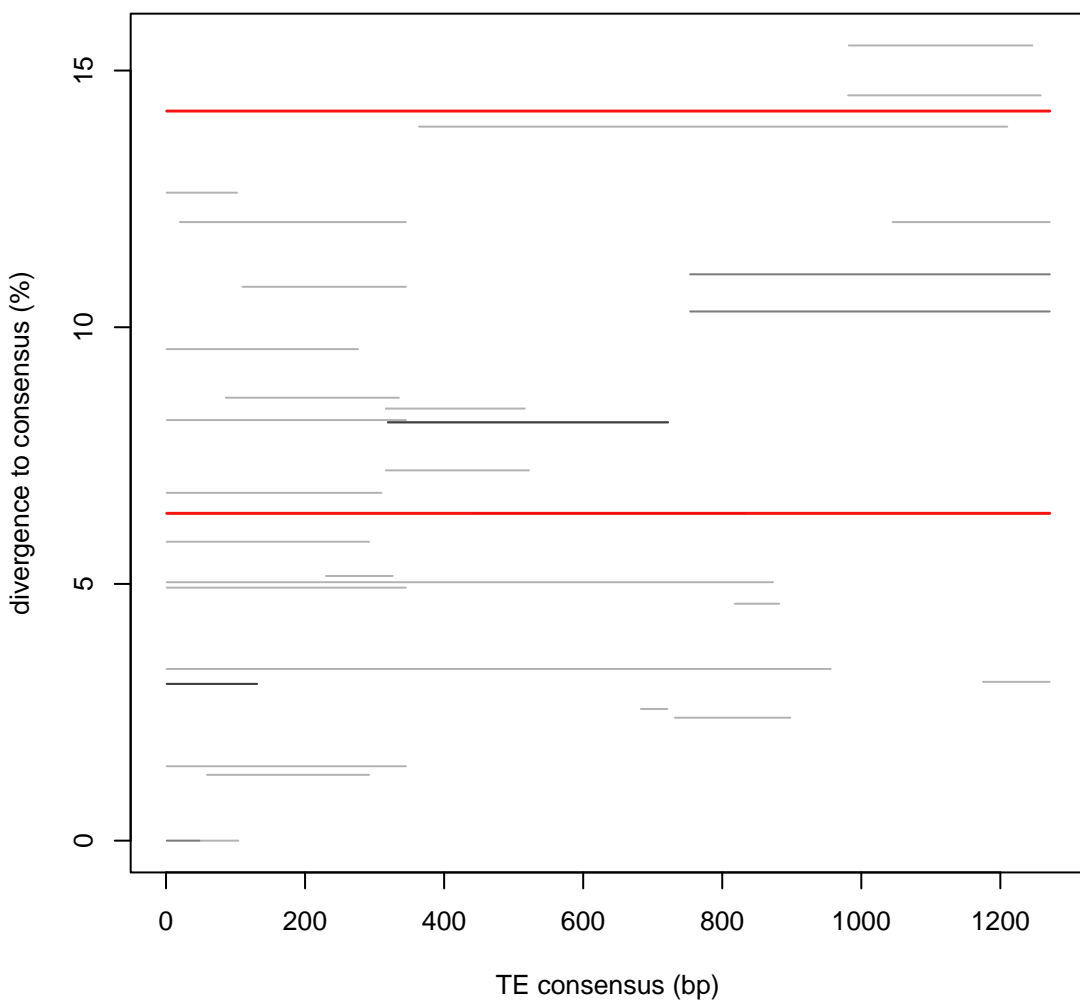
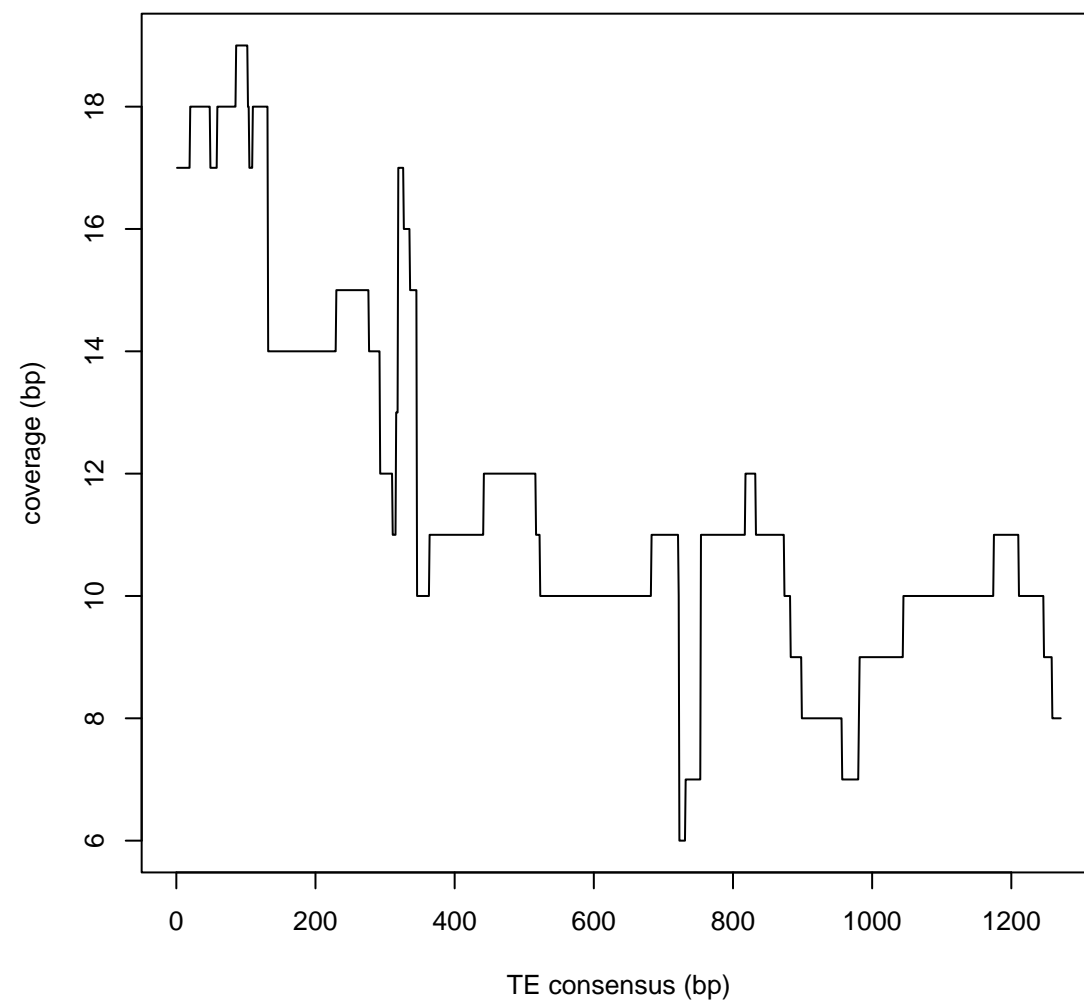


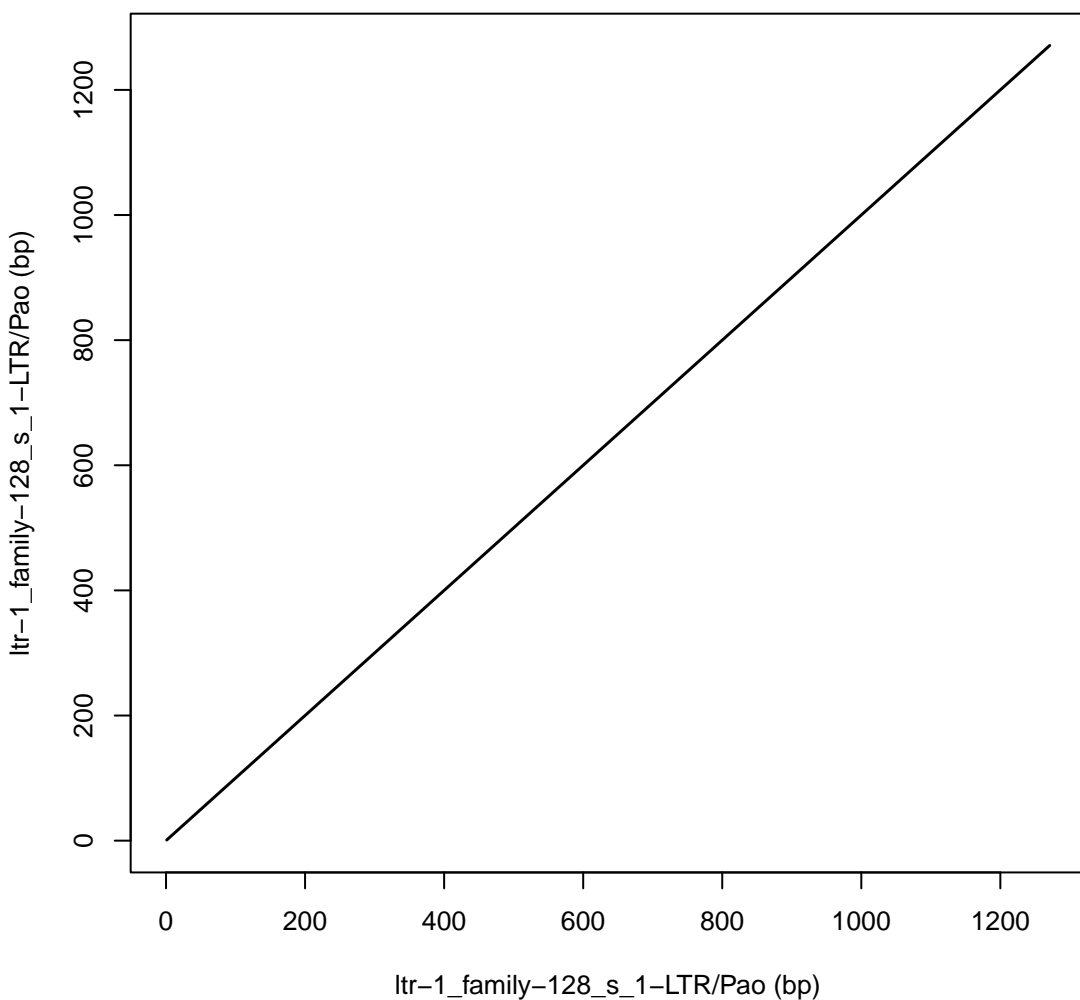
TE: ltr-1_family-128_s_1-LTR/Pao
consensus size: 1271bp; fragments: 41; full length: 2 (≥ 1143.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

