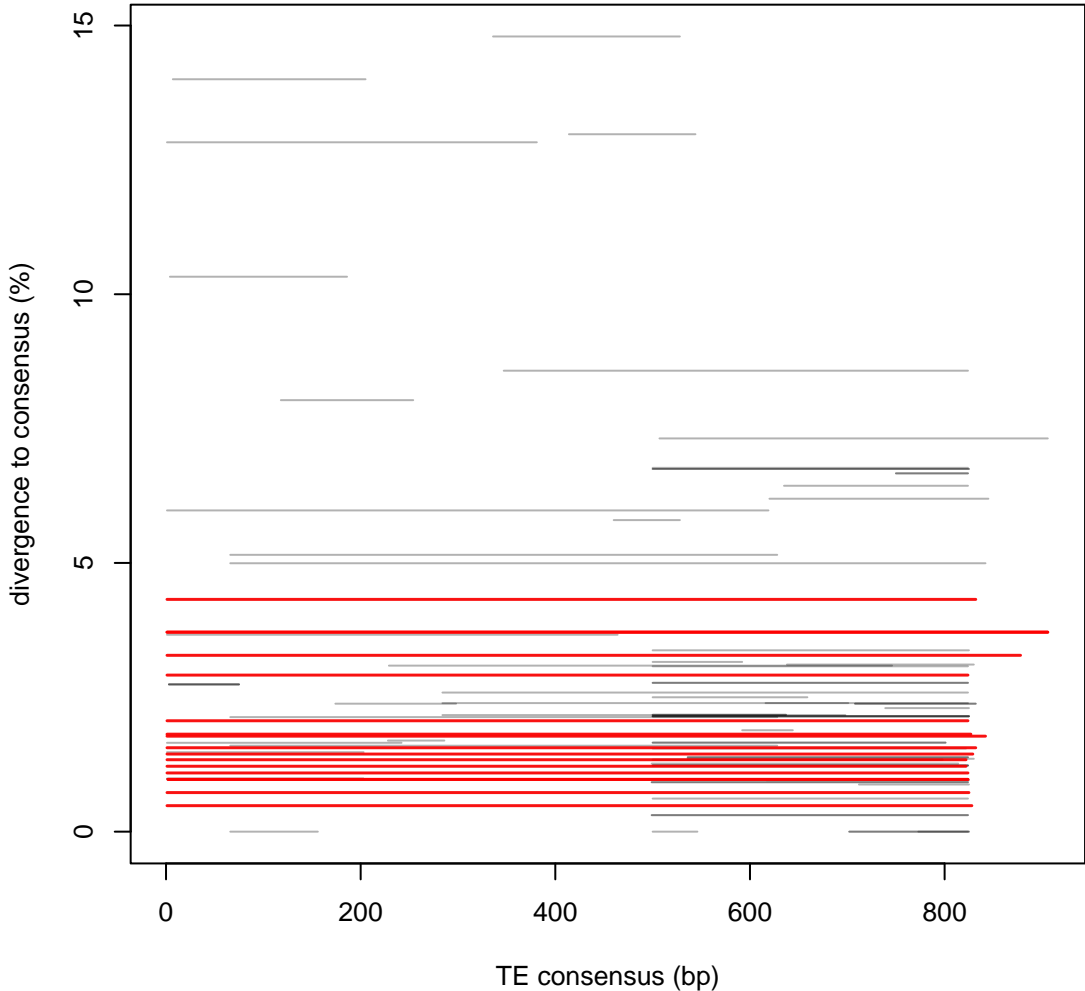
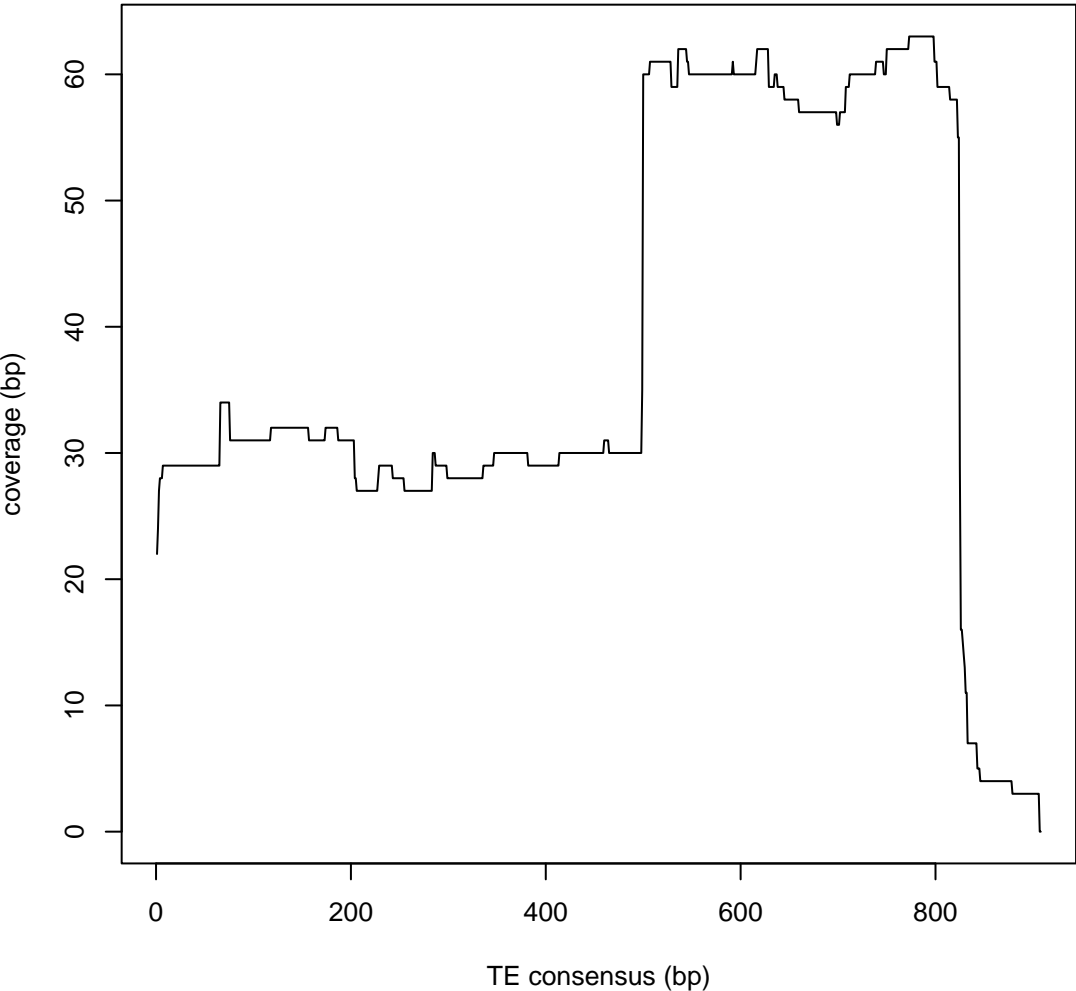


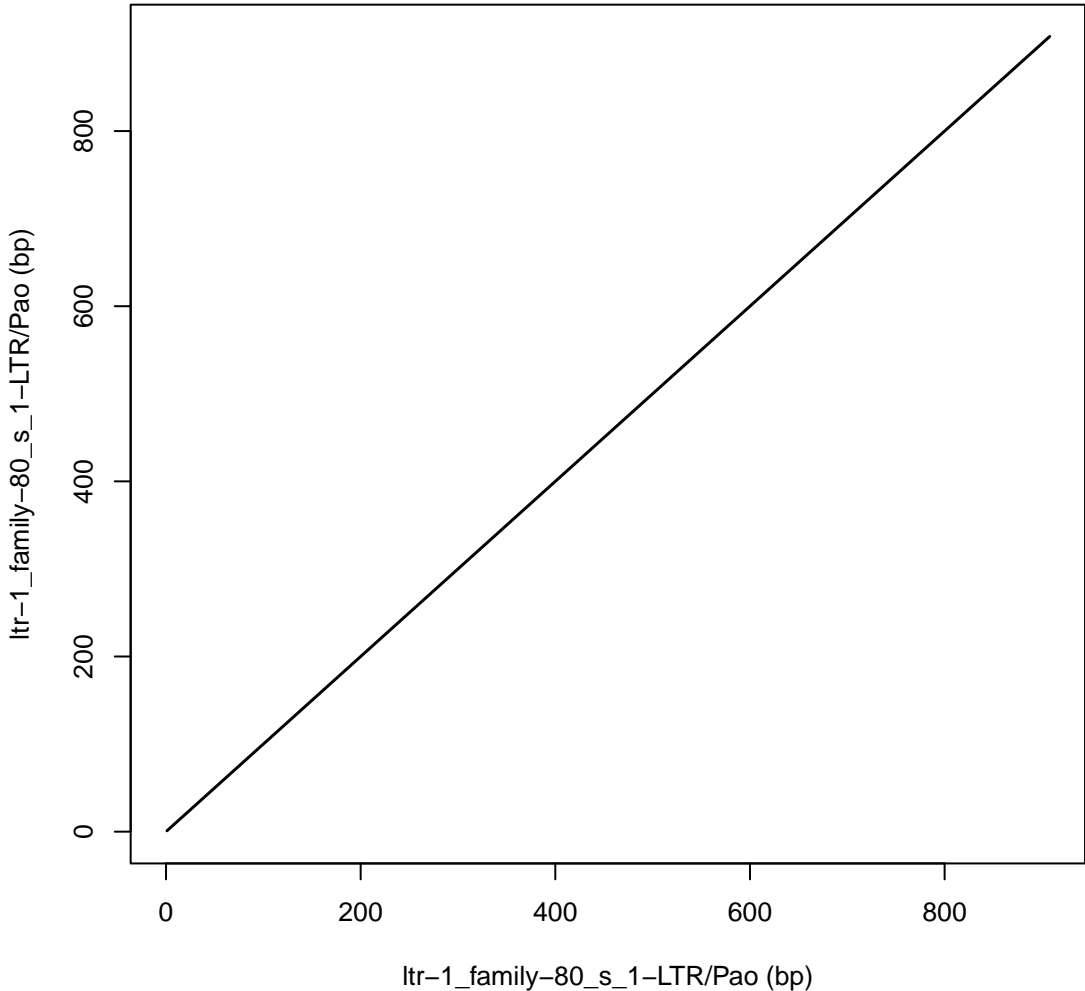
TE: ltr-1_family-80_s_1-LTR/Pao
consensus size: 908bp; fragments: 94; full length: 17 (>=817.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

