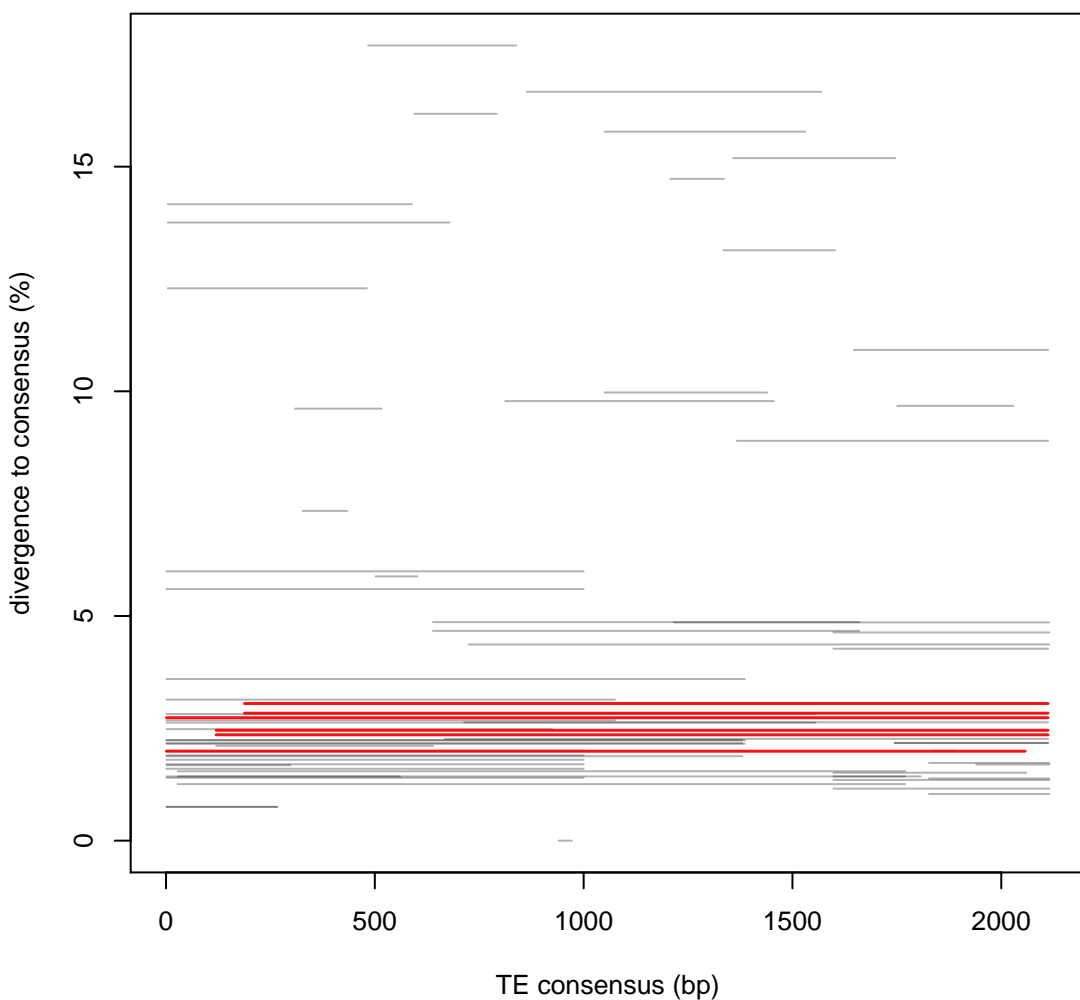
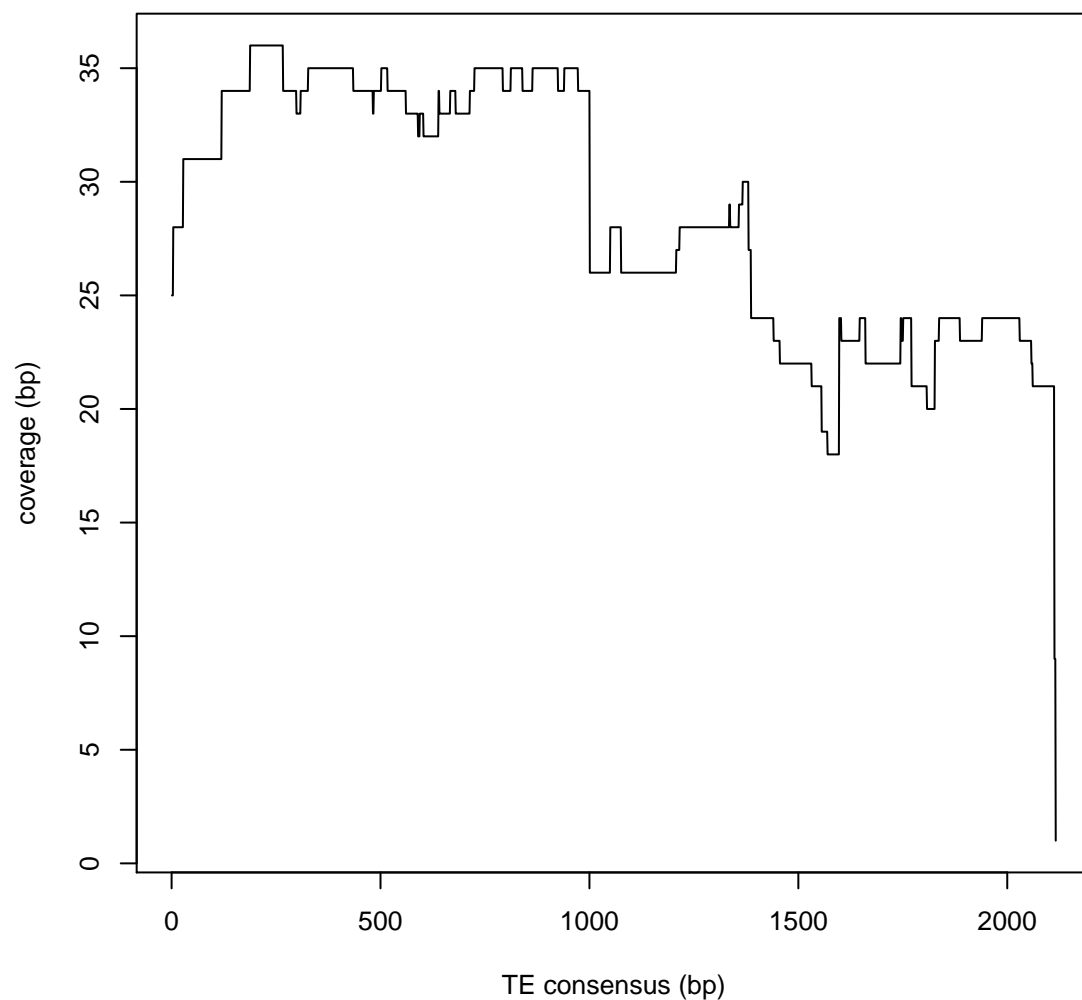


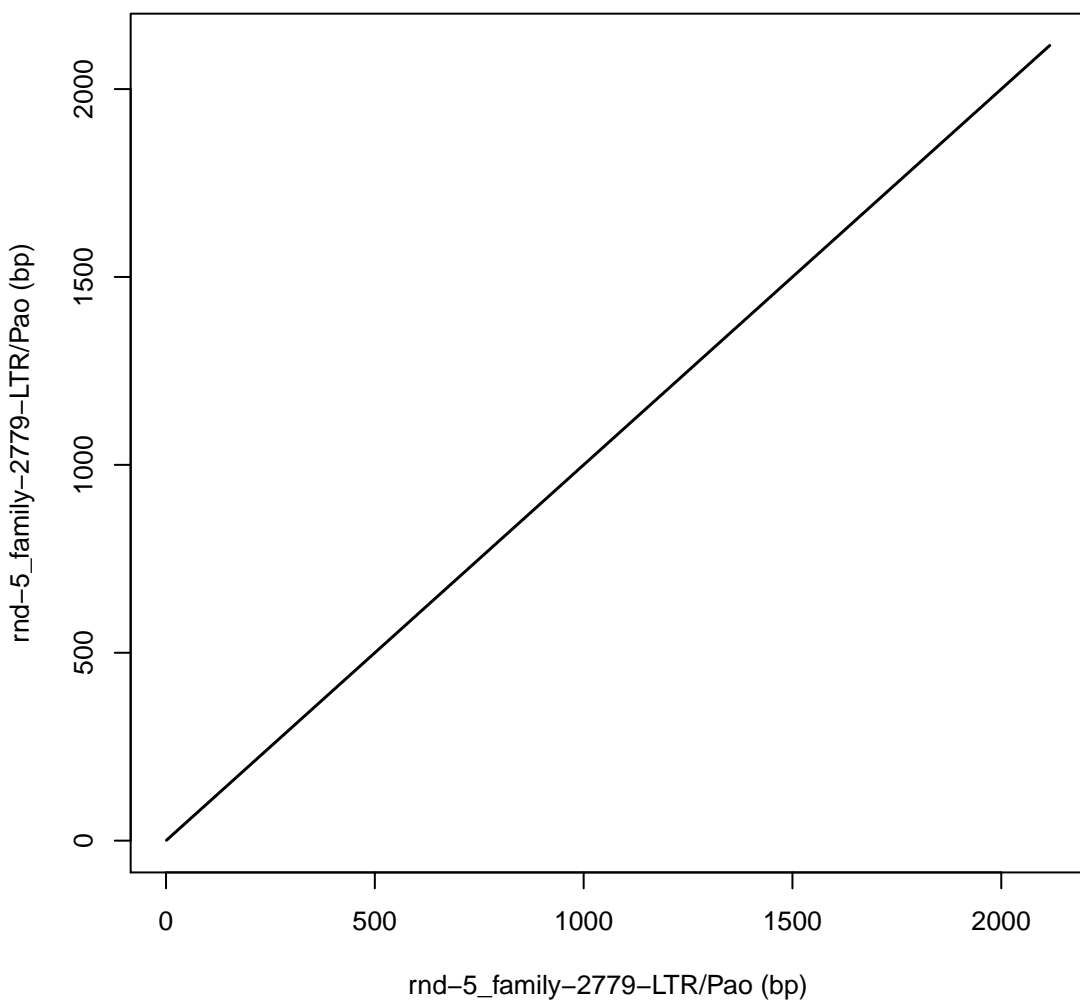
TE: rnd-5_family-2779-LTR/Pao
consensus size: 2116bp; fragments: 71; full length: 6 (≥ 1904.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

