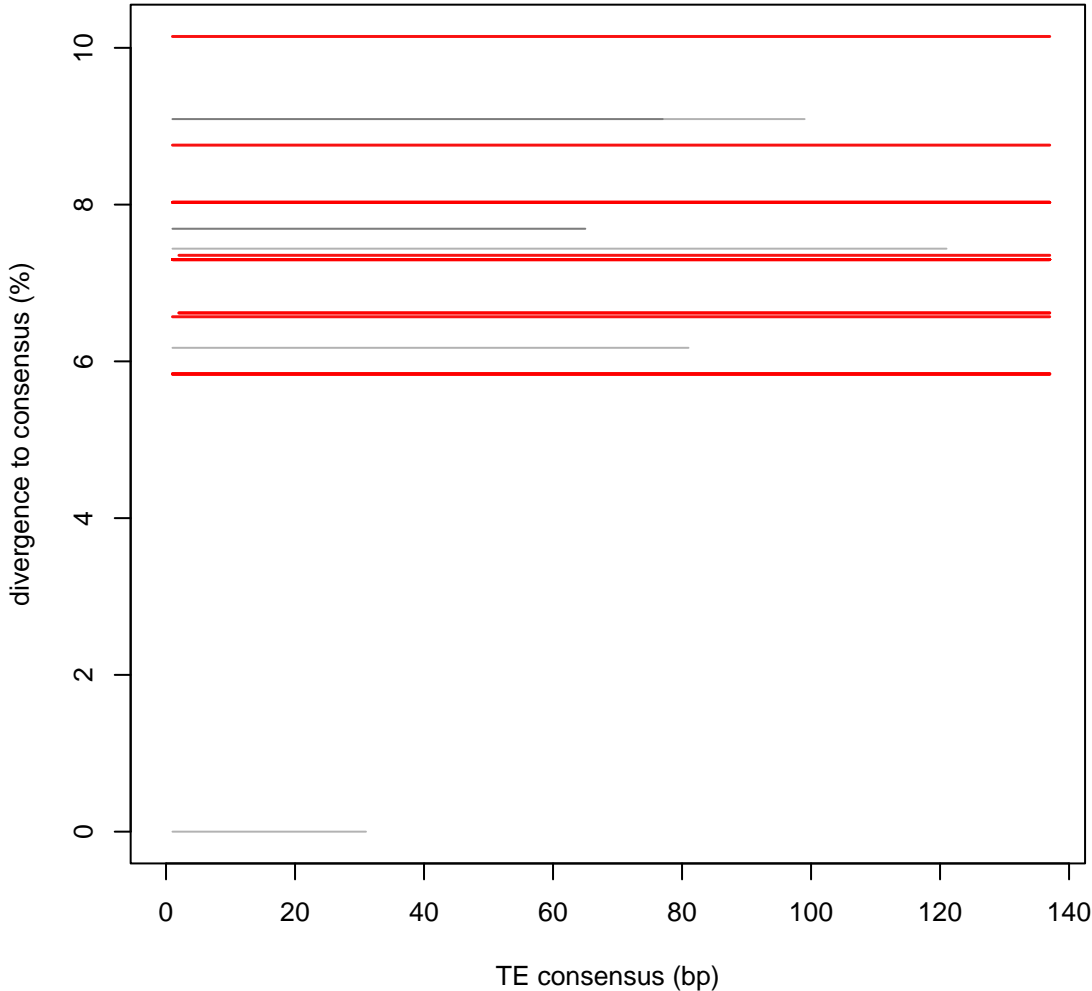
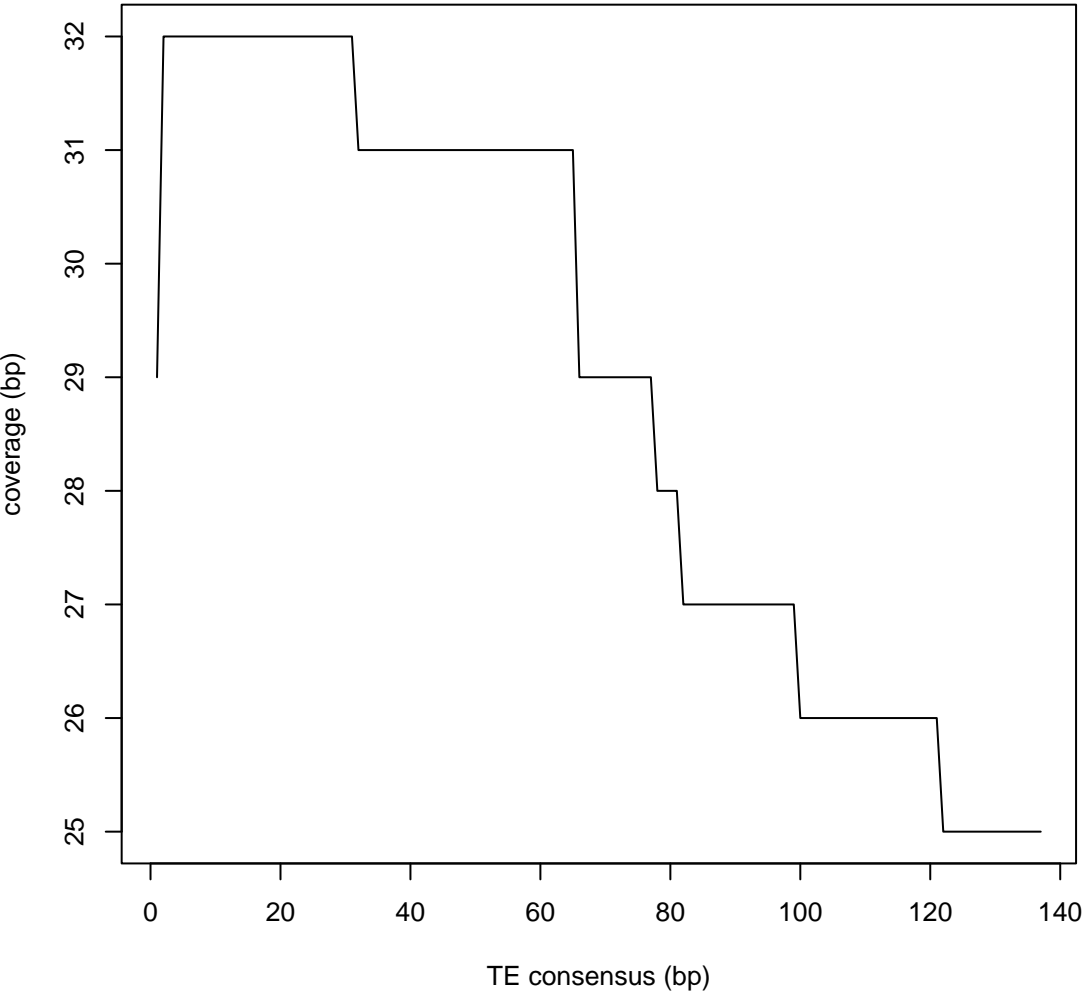


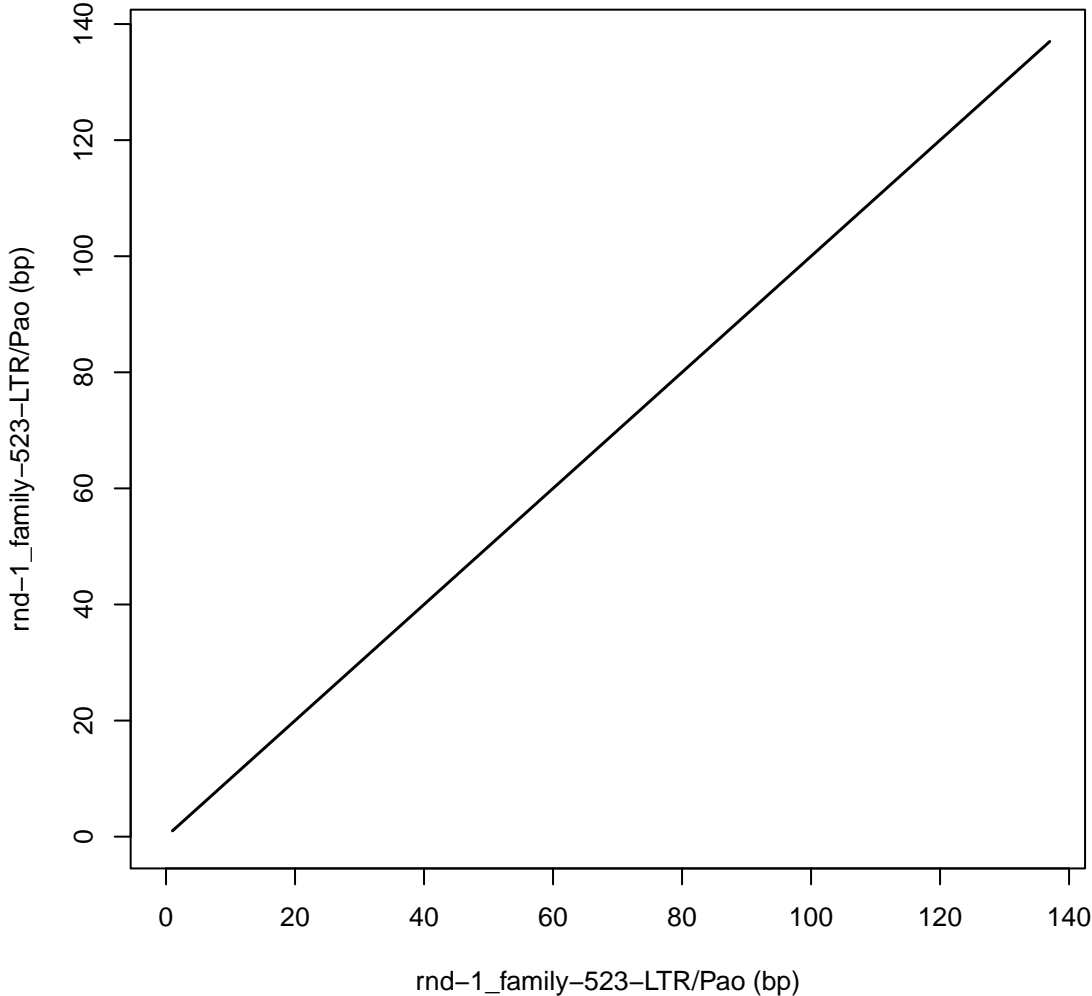
TE: rnd-1_family-523-LTR/Pao
consensus size: 137bp; fragments: 32; full length: 25 (≥ 123.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

