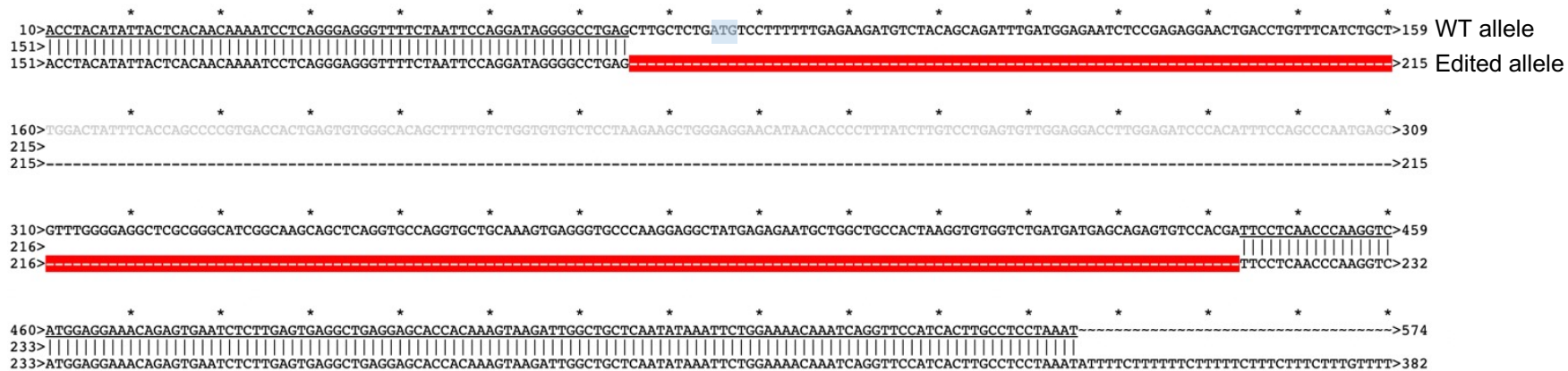
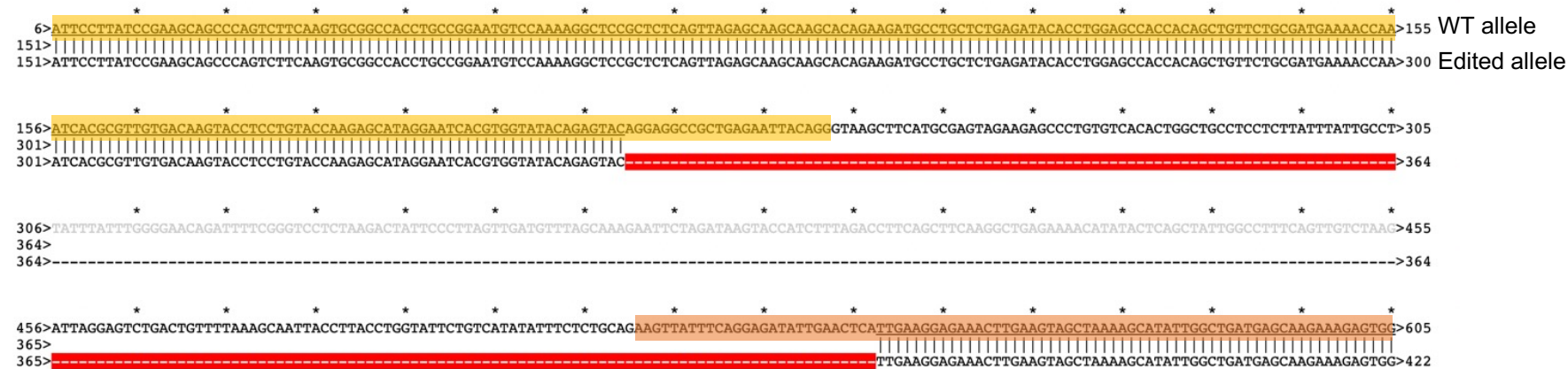


(A) Colony "95" containing a 368-base pair deletion for *TRIML1* flanking the start codon. No wild-type alleles were detected.



Start codon

(B) Colony "153" containing a 328-base pair deletion for *TRIML2* on exons 2 and 3. No wild-type alleles were detected.



Exon 2 Exon 3