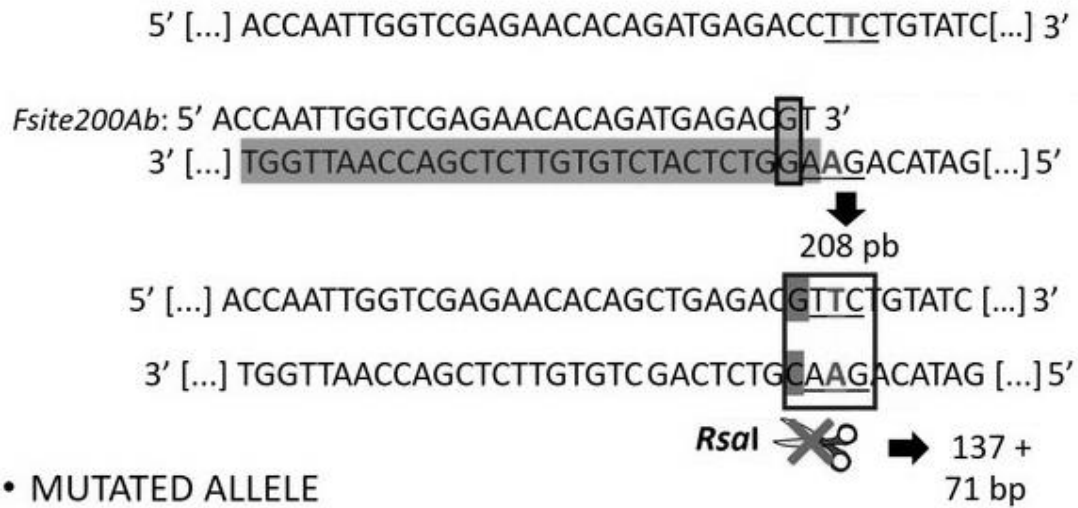
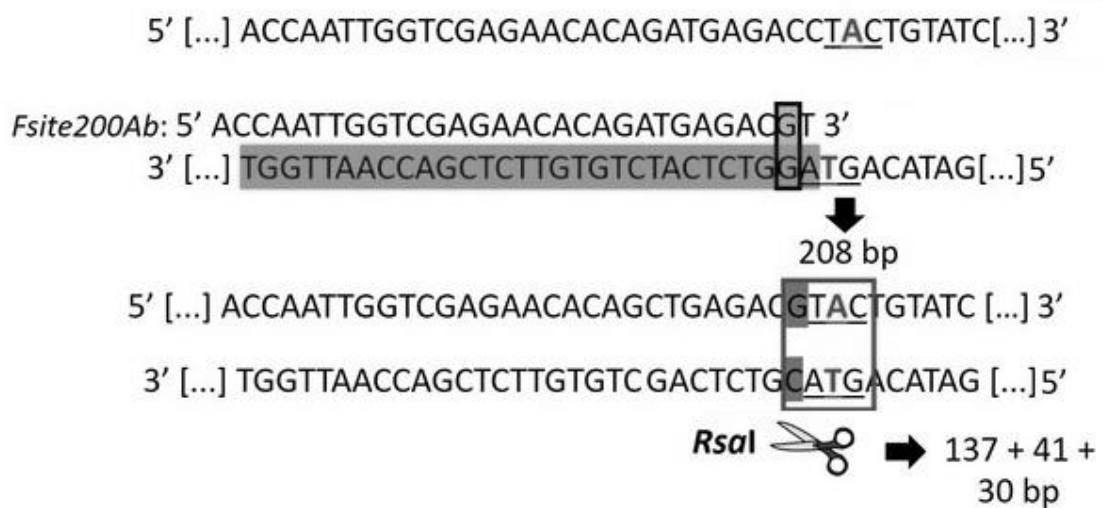


- UNMUTATED ALLELE



- MUTATED ALLELE



**Additional file 3: Figure S2.** Representation of the methodology used to analyze codon 200 of the beta-tubulin isotype 1 gene from *Ancylostoma braziliense*. The *Fsite200Ab* primer was designed to add a mutation in the amplicon, regardless of whether there was mutation at codon 200. The presence of the mutation at codon 200 in combination with the primer-introduced change creates a site for *RsaI*. If the allele does not harbor a mutation, even the altered primer cannot create a *RsaI* site. Codon 200 is underlined, with the base of interest in bold.