

Inducible high-efficiency CRISPR-Cas9-targeted gene editing and
precision base editing in African trypanosomes

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Supplementary Information.

Supplementary Figures 1-2.

Uncropped
gel and blots
for Fig. 1.

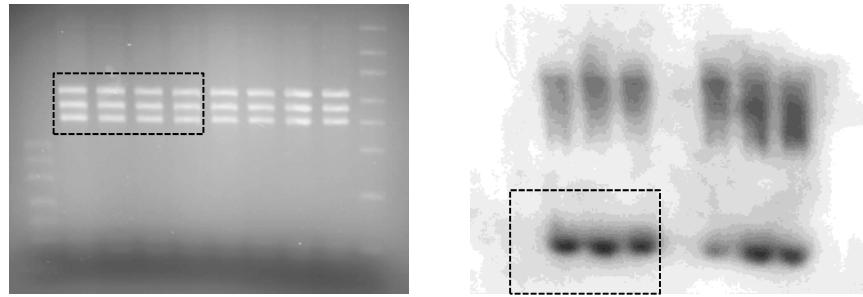
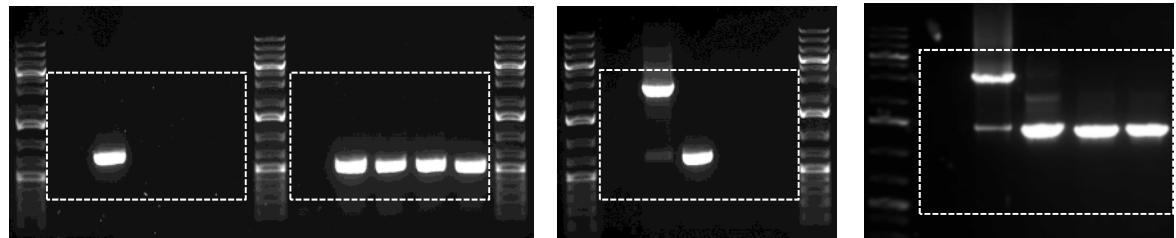


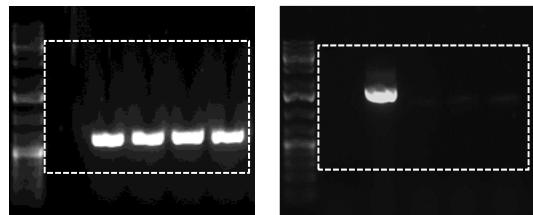
Fig. 1b.

Fig. 1e.

Uncropped
gels for
Fig. 2c.



Uncropped
gels for
Fig. 3c.



Uncropped
gels for
Fig. 4.

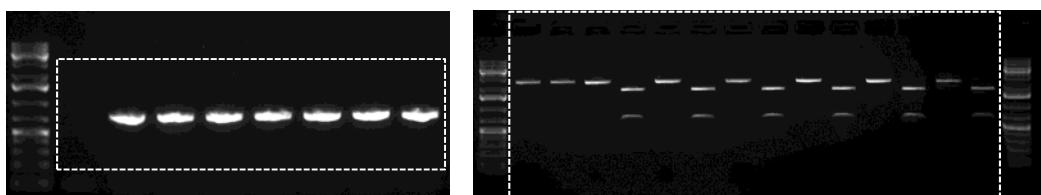
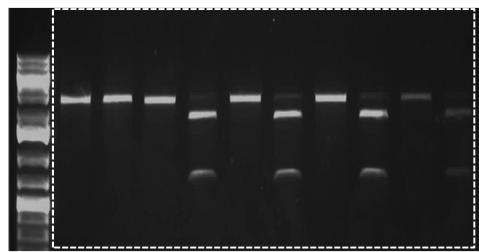
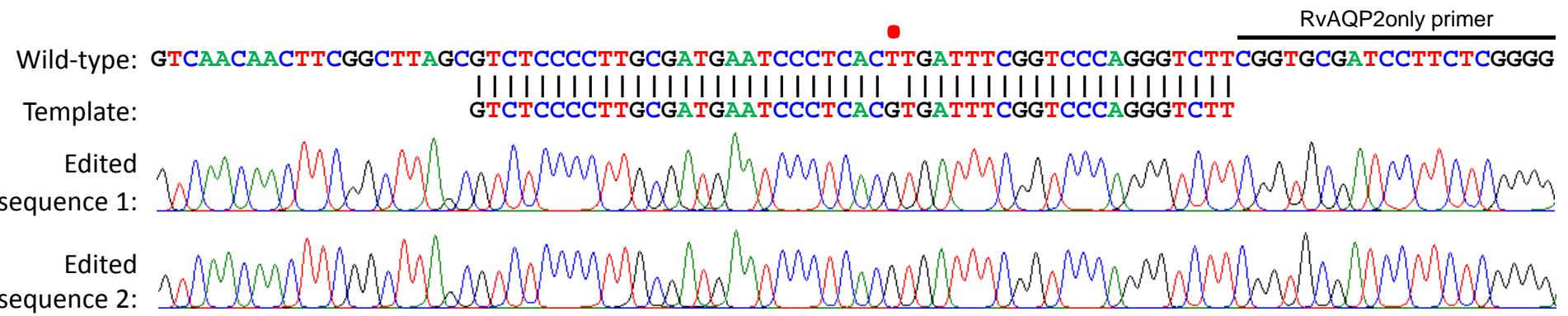


Fig. 4c.

Fig. 4d.



Supplementary Fig. 1. All uncropped gels and blots are shown for Figures 1-4. The areas shown in Figures 1-4 are indicated by dashed boxes. Data for a second clone are also shown in the Fig. 1 panels.



Supplementary Figure 2. Extended sequence traces from Fig 4b and for two independent clones, showing only the precision T⁷⁹¹G edit and no additional changes in the template or flanking regions. The location of one of the primers used to amplify fragments for sequencing is indicated.