Manuscript Title

This manuscript ([permalink](https://malariagen.github.io/manubot-test/v/9d5818cfd7a3f3083c06d17269888e9641aec647/)) was automatically generated from [malariagen/manubot-test@9d5818c](https://github.com/malariagen/manubot-test/tree/9d5818cfd7a3f3083c06d17269888e9641aec647) on May 30, 2020.

## Authors

* **John Doe** ORCID icon [XXXX-XXXX-XXXX-XXXX](https://orcid.org/XXXX-XXXX-XXXX-XXXX) · GitHub icon [johndoe](https://github.com/johndoe) · Twitter icon [johndoe](https://twitter.com/johndoe) Department of Something, University of Whatever · Funded by Grant XXXXXXXX
* **Jane Roe** ORCID icon [XXXX-XXXX-XXXX-XXXX](https://orcid.org/XXXX-XXXX-XXXX-XXXX) · GitHub icon [janeroe](https://github.com/janeroe) Department of Something, University of Whatever; Department of Whatever, University of Something

## Abstract

This manuscript is a template (aka “rootstock”) for [Manubot](https://manubot.org/), a tool for writing scholarly manuscripts. Use this template as a starting point for your manuscript.

The rest of this document is a full list of formatting elements/features supported by Manubot. Compare the input (.md files in the /content directory) to the output you see below.

## Basic formatting

**Bold** **text**

Semi-bold text

Centered text

Right-aligned text

*Italic* *text*

Combined *italics and* ***bold***

~~Strikethrough~~

1. Ordered list item
2. Ordered list item
   1. Sub-item
   2. Sub-item
      1. Sub-sub-item
3. Ordered list item
   1. Sub-item

* List item
* List item
* List item

subscript: H2O is a liquid

superscript: 210 is 1024.

[unicode superscripts](https://www.google.com/search?q=superscript+generator)⁰¹²³⁴⁵⁶⁷⁸⁹

[unicode subscripts](https://www.google.com/search?q=superscript+generator)₀₁₂₃₄₅₆₇₈₉

A long paragraph of text. Lorem ipsum dolor sit amet, consectetur adipiscing elit, sed do eiusmod tempor incididunt ut labore et dolore magna aliqua. Ut enim ad minim veniam, quis nostrud exercitation ullamco laboris nisi ut aliquip ex ea commodo consequat. Duis aute irure dolor in reprehenderit in voluptate velit esse cillum dolore eu fugiat nulla pariatur. Excepteur sint occaecat cupidatat non proident, sunt in culpa qui officia deserunt mollit anim id est laborum.

Putting each sentence on its own line has numerous benefits with regard to [editing](https://asciidoctor.org/docs/asciidoc-recommended-practices/#one-sentence-per-line) and [version control](https://rhodesmill.org/brandon/2012/one-sentence-per-line/).

Line break without starting a new paragraph by putting  
two spaces at end of line.

## Document organization

Document section headings:

# Heading 1

## Heading 2

### Heading 3

#### Heading 4

### A heading centered on its own printed page

Horizontal rule:

Heading 1’s are recommended to be reserved for the title of the manuscript.

Heading 2’s are recommended for broad sections such as *Abstract*, *Methods*, *Conclusion*, etc.

Heading 3’s and Heading 4’s are recommended for sub-sections.

## Links

Bare URL link: <https://manubot.org>

[Long link with lots of words and stuff and junk and bleep and blah and stuff and other stuff and more stuff yeah](https://manubot.org)

[Link with text](https://manubot.org)

[Link with hover text](https://manubot.org)

[Link by reference](https://manubot.org)

## Citations

Citation by DOI.[1](#ref-IhliSZDo)

Citation by PubMed Central ID.[2](#ref-mSMVRkoc)

Citation by PubMed ID.[3](#ref-126Wi5Us4)

Citation by Wikidata ID.[4](#ref-QhC8yJ7V)

Citation by ISBN.[5](#ref-zBPP9YKu)

Citation by URL.[6](#ref-6acIRW4q)

Citation by alias.[7](#ref-PZMP42Ak)

Multiple citations can be put inside the same set of brackets.[1](#ref-IhliSZDo),[5](#ref-zBPP9YKu),[7](#ref-PZMP42Ak) Manubot plugins provide easier, more convenient visualization of and navigation between citations.[2](#ref-mSMVRkoc),[3](#ref-126Wi5Us4),[7](#ref-PZMP42Ak),[8](#ref-YuJbg3zO)

Citation tags (i.e. aliases) can be defined in their own paragraphs using Markdown’s reference link syntax:

## Referencing figures, tables, equations

Figure [1](#fig:square-image)

Figure [2](#fig:wide-image)

Figure [3](#fig:tall-image)

Figure [4](#fig:vector-image)

Table [1](#tbl:bowling-scores)

Equation [1](#eq:regular-equation)

Equation [2](#eq:long-equation)

## Quotes and code

Quoted text

Quoted block of text

Two roads diverged in a wood, and I—  
I took the one less traveled by,  
And that has made all the difference.

Code in the middle of normal text, aka inline code.

Code block with Python syntax highlighting:

from manubot.cite.doi import expand\_short\_doi  
  
def test\_expand\_short\_doi():  
 doi = expand\_short\_doi("10/c3bp")  
 # a string too long to fit within page:  
 assert doi == "10.25313/2524-2695-2018-3-vliyanie-enhansera-copia-i-insulyatora-gypsy-na-sintez-ernk-modifikatsii-hromatina-i-svyazyvanie-insulyatornyh-belkov-vtransfetsirovannyh-geneticheskih-konstruktsiyah"

Code block with no syntax highlighting:

Exporting HTML manuscript  
Exporting DOCX manuscript  
Exporting PDF manuscript

## Figures



Figure 1: **A square image at actual size and with a bottom caption.** Loaded from the latest version of image on GitHub.



Figure 2: **An image too wide to fit within page at full size.** Loaded from a specific (hashed) version of the image on GitHub.



Figure 3: **A tall image with a specified height.** Loaded from a specific (hashed) version of the image on GitHub.

Figure 4: A vector .svg image loaded from GitHub. The parameter sanitize=true is necessary to properly load SVGs hosted via GitHub URLs. White background specified to serve as a backdrop for transparent sections of the image.

Figure 4: **A vector .svg image loaded from GitHub.** The parameter sanitize=true is necessary to properly load SVGs hosted via GitHub URLs. White background specified to serve as a backdrop for transparent sections of the image.

## Tables

Table 1: A table with a top caption and specified relative column widths.

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| *Bowling Scores* | Jane | John | Alice | Bob |
| Game 1 | 150 | 187 | 210 | 105 |
| Game 2 | 98 | 202 | 197 | 102 |
| Game 3 | 123 | 180 | 238 | 134 |

Table 2: A table too wide to fit within page.

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
|  | Digits 1-33 | Digits 34-66 | Digits 67-99 | Ref. |
| pi | 3.14159265358979323846264338327950 | 288419716939937510582097494459230 | 781640628620899862803482534211706 | [piday.org](https://www.piday.org/million/) |
| e | 2.71828182845904523536028747135266 | 249775724709369995957496696762772 | 407663035354759457138217852516642 | [nasa.gov](https://apod.nasa.gov/htmltest/gifcity/e.2mil) |

Table 3: A table with merged cells using the attributes plugin.

|  |  |  |
| --- | --- | --- |
|  | **Colors** |  |
| **Size** | **Text Color** | **Background Color** |
| big | blue | orange |
| small | black | white |

## Equations

A LaTeX equation:

An equation too long to fit within page:

## Special

WARNING *The following features are only supported and intended for .html and .pdf exports.* *Journals are not likely to support them, and they may not display correctly when converted to other formats such as .docx.*

[Link styled as a button](https://manubot.org)

Adding arbitrary HTML attributes to an element using Pandoc’s attribute syntax:

Manubot Manubot Manubot Manubot Manubot. Manubot Manubot Manubot Manubot. Manubot Manubot Manubot. Manubot Manubot. Manubot.

Adding arbitrary HTML attributes to an element with the Manubot attributes plugin (more flexible than Pandoc’s method in terms of which elements you can add attributes to):

Manubot Manubot Manubot Manubot Manubot. Manubot Manubot Manubot Manubot. Manubot Manubot Manubot. Manubot Manubot. Manubot.

Available background colors for text, images, code, banners, etc:

white lightgrey grey darkgrey black lightred lightyellow lightgreen lightblue lightpurple red orange yellow green blue purple

Using the [Font Awesome](https://fontawesome.com/) icon set:

**Light Grey Banner** useful for *general information* - [manubot.org](https://manubot.org/)

**Blue Banner** useful for *important information* - [manubot.org](https://manubot.org/)

**Light Red Banner** useful for *warnings* - [manubot.org](https://manubot.org/)

## Tables (misc. test)

|  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 |
| 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 |

At 6pt by wrapping in <div> with a style:

|  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 |
| 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 |

## References (misc. test)

URL citation overridden by manual-references.json.[9](#ref-OvL2rC9o)

## Test table from CSV hard-coded into Markdown content

Using the pantable Pandoc filter.

*Awesome* **Markdown** Table

|  |  |  |
| --- | --- | --- |
| First row | defaulted to be header row | can be disabled |
| 1 | cell can contain **markdown** | It can be aribrary block element:   * following standard markdown syntax * like this |
| 2 | Any markdown syntax, e.g. |  |

## Test table from CSV file source

* Using the pantable Pandoc filter with the include option, loading from content/csv/by\_country.csv.
* Alignment has been specified: LRRRRRRR.
* ‘Samples By Country’ is the specified caption.
* dialect: unix is passed to csv.reader() by way of csv-kwargs

Samples By Country

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
| country | is\_arabiensis | is\_coluzzii | is\_gambiae | year | latitude | longitude | contributor |
| Angola | 0 | 81 | 0 | 2009 | -8.884000000000011 | 13.302000000000008 | Joao Pinto |
| Burkina Faso | 3 | 135 | 157 | 2014 | 11.248256756756797 | -4.28192905405405 | Nora Besansky |
| Cameroon | 2 | 26 | 416 | 2013 | 5.640274774774801 | 13.536222972972988 | Nora Besansky |
| Central African Republic | 0 | 18 | 55 | 1994 | 4.367000000000001 | 18.583000000000002 | Alessandra della Torre |
| Cote d'Ivoire | 0 | 80 | 0 | 2012 | 5.8980000000000095 | -4.822999999999997 | David Weetman |
| Democratic Republic of Congo | 0 | 0 | 76 | 2015 | 4.282999999999999 | 21.017000000000035 | David Weetman |
| Equatorial Guinea | 0 | 0 | 10 | 2002 | 3.7 | 8.700000000000001 | Igor Sharakhov |
| Gabon | 0 | 0 | 69 | 2000 | 0.38400000000000023 | 9.455000000000005 | Joao Pinto |
| Gambia, The | 0 | 169 | 69 | 2012 | 13.552845878136198 | -15.294989247311792 | Martin Donnelly |
| Ghana | 0 | 64 | 36 | 2012 | 5.57405 | -0.9347799999999995 | David Weetman |
| Guinea | 0 | 11 | 123 | 2012 | 8.84470588235294 | -9.757941176470567 | Ken Vernick |
| Guinea-Bissau | 0 | 0 | 29 | 2010 | 11.942742574257442 | -15.496168316831659 | Joao Pinto |
| Kenya | 13 | 0 | 28 | 2012 | -3.5110000000000015 | 39.90900000000008 | Janet Midega |
| Lab Cross | 0 | 0 | 0 | -1 | 53.4089999999997 | -2.9690000000000367 | Martin Donnelly |
| Malawi | 41 | 0 | 0 | 2015 | -15.932999999999996 | 34.75500000000003 | Martin Donnelly |
| Mali | 2 | 91 | 131 | 2014 | 12.01431111111115 | -7.824622222222243 | Nora Besansky |
| Mayotte | 0 | 0 | 23 | 2011 | -12.85708695652174 | 45.137434782608686 | Igor Sharakhov |
| Mozambique | 0 | 0 | 74 | 2004 | -23.71599999999996 | 35.298999999999985 | Joao Pinto |
| Tanzania | 225 | 0 | 68 | 2015 | -2.508323333333342 | 33.85035666666663 | David Weetman |
| Uganda | 82 | 0 | 207 | 2012 | 0.2664965517241366 | 32.59427586206877 | Martin Donnelly |

* See that column headings are carried onto the next page.
* [Pantable](https://github.com/ickc/pantable) uses [Panflute], e.g. https://github.com/ickc/pantable/blob/master/pantable/csv\_to\_table\_markdown.py
* If we needed to apply custom formatting during the build, e.g. to round those floats, we could probably write our own Pandoc filter.

## Test figures from image sources

* May need to use secno=1 on all figures, e.g. {#fig:id secno=1} to avoid mislabelling/miscounting bug? https://github.com/tomduck/pandoc-fignos/issues/76

### 1) Separate figures with alt text (“captions”), no resizing, refs linked by default

Figure 5: Stairway Plot

Figure 5: Stairway Plot

Figure 6: ROH Scatter Plot

Figure 6: ROH Scatter Plot

[**5**](#fig:figure-test-1a), Stairway plot of changes in population size over time. Absolute values of time and Ne are shown on alternative axes as a range of values, assuming lower and upper limits for the mutation rate μ as 2.8 × 10−9 and 5.5 × 10−9, respectively, and t = 11 generations per year. ka, thousand years ago. [**6**](#fig:figure-test-1b), Runs of homozygosity (ROH) in individual mosquitoes, highlighting recent inbreeding in Kenyan (grey) and colony (black) mosquitoes. G, Ghana; K, Kisumu; M, Mali; P, Pimperena.

* The “Figure {n}:” label before the alt text “captions”, e.g. Figure 6s: ROH Scatter Plot, is added automatically.
* Btw, {@fig:figure-test-1b} appears as {???}: when in “quotes”, e.g. “{**???**}”, which is unexpected, possibly a bug
* Curly brackets are to separate fig ref from adjacent non-whitespace strings, e.g. @fig:figure-test-1b- versus {@fig:figure-test-1b}-

### 2) Separate figures with no alt text (no “captions”), no resizing, refs linked by default

Figure 7: 

Figure 7:

Figure 8: 

Figure 8:

[**7**](#fig:figure-test-2a), Stairway plot of changes in population size over time. Absolute values of time and Ne are shown on alternative axes as a range of values, assuming lower and upper limits for the mutation rate μ as 2.8 × 10−9 and 5.5 × 10−9, respectively, and t = 11 generations per year. ka, thousand years ago. [**8**](#fig:figure-test-2b), Runs of homozygosity (ROH) in individual mosquitoes, highlighting recent inbreeding in Kenyan (grey) and colony (black) mosquitoes. G, Ghana; K, Kisumu; M, Mali; P, Pimperena.

* The “Figure {n}:” label before the empty alt text “captions”, e.g. Figure 8:, is added automatically.
  + Want a way to override this label’s template
* The colon after the Figure 8:, which anticipates text, looks odd/confusing when there is no text. It looks like it applies to the content that follows after that label, but it doesn’t/shouldn’t.

### 3) Separate subfigures using [Pandoc “divs”](https://pandoc.org/MANUAL.html#divs-and-spans), with no alt text, no layout, no resizing

**Figure 1**





**a**, Stairway plot of changes in population size over time. Absolute values of time and Ne are shown on alternative axes as a range of values, assuming lower and upper limits for the mutation rate μ as 2.8 × 10−9 and 5.5 × 10−9, respectively, and t = 11 generations per year. ka, thousand years ago. **b**, Runs of homozygosity (ROH) in individual mosquitoes, highlighting recent inbreeding in Kenyan (grey) and colony (black) mosquitoes. G, Ghana; K, Kisumu; M, Mali; P, Pimperena.

* Pandoc divs let you group content under one Manubot figure reference, in this case two simple images (not Manubot figures in their own right)
* Putting the Markdown for the figures on separate paragraphs (as required by Pandoc) causes a vertical layout by default, which is undesirable here
* Figure references for [**1**](#fig:figure-test3)**a** and [**1**](#fig:figure-test3)**b** are just references/links to [**1**](#fig:figure-test3) plus manually managed suffixes
* Can switch off the link for figure references, e.g. @fig:figure-test3{nolink=True} gives 1 rather than [1](#fig:figure-test3)
* The figure label was added in manually, i.e. \*\*Figure @fig:figure-test3{nolink=True}\*\*, otherwise there would be no visible label for this figure, even though it has a reference
* Note figure-test3 has reset to “1”, rather than [8](#fig:figure-test-2b) + 1. Bug? https://github.com/tomduck/pandoc-fignos/issues/76
  + Using secno=1 here, i.e. {#fig:figure-test3 secno=1} has made no difference, but perhaps need to use secno=1 on **all** figures in order to workaround. (?!)

### 4) Breaking of pandoc-fignos when figures are not in their own Markdown paragraph

Stairway Plot ROH Scatter Plot

**??**, Stairway plot of changes in population size over time. Absolute values of time and Ne are shown on alternative axes as a range of values, assuming lower and upper limits for the mutation rate μ as 2.8 × 10−9 and 5.5 × 10−9, respectively, and t = 11 generations per year. ka, thousand years ago. **???{nolink=True}**, Runs of homozygosity (ROH) in individual mosquitoes, highlighting recent inbreeding in Kenyan (grey) and colony (black) mosquitoes. G, Ghana; K, Kisumu; M, Mali; P, Pimperena.

* See broken refs as “??” and ???{nolink=True}.
* Reports errors:
  + pandoc-fignos: Bad reference: @fig:figure-test4a.
  + Generating csl\_item for 'figure-test3b' failed due to a NotImplementedError: Manubot does not know how to generate a csl\_item for 'figure-test3b'

Pandoc requires:

![Stairway Plot](images/stairway\_plot.svg){#fig:figure-test4a}  
  
![ROH Scatter Plot](images/roh\_scatter.svg){#fig:figure-test4b}

Instead of the above:

![Stairway Plot](images/stairway\_plot.svg){#fig:figure-test4a}  
![ROH Scatter Plot](images/roh\_scatter.svg){#fig:figure-test4b}

* This constraint prevents figures from being placed next to eachother by a simple inline method (as above) without breaking the managed references.

### 5) Separate figures with HTML table-based layout, with tags, no alt text, no resizing

Figure a: 

Figure a:

Figure b: 

Figure b:

**a**, Stairway plot of changes in population size over time. Absolute values of time and Ne are shown on alternative axes as a range of values, assuming lower and upper limits for the mutation rate μ as 2.8 × 10−9 and 5.5 × 10−9, respectively, and t = 11 generations per year. ka, thousand years ago. **b**, Runs of homozygosity (ROH) in individual mosquitoes, highlighting recent inbreeding in Kenyan (grey) and colony (black) mosquitoes. G, Ghana; K, Kisumu; M, Mali; P, Pimperena.

* If the HTML contains any preceding whitespace in the markup, e.g. indentation, you will see broken HTML markup
* HTML table-based layout could be used to place the figures next to each other while still meeting the Pandoc separate-paragraph rule, but it’s a bit of a hack
* Table cell borders removed using <td style="border: none">
* Manubot shows a figure label for each figure, even when no alt/caption text. Here the figure label/number is overidden usign specified “tags”
* Tags can be used to override the automatic managed figure number, e.g. here @fig:figure-test5a is labelled as “a” instead of an automatically incremented figure number. In other words, normally the figure label would be a number, which would be incremented automatically, except for bug/feature: https://github.com/tomduck/pandoc-fignos/issues/76, but here we have overridden that figure label.
* Again, the colon after the figure label, which anticipates text, looks odd/confusing when there is no text. It looks like it applies to the content that follows after that label, but it doesn’t/shouldn’t.

### 6) Breaking of separate figures with Markdown table-based layout, no alt text, no resizing

|  |  |
| --- | --- |
|  |  |

**??**, Stairway plot of changes in population size over time. Absolute values of time and Ne are shown on alternative axes as a range of values, assuming lower and upper limits for the mutation rate μ as 2.8 × 10−9 and 5.5 × 10−9, respectively, and t = 11 generations per year. ka, thousand years ago. **??**, Runs of homozygosity (ROH) in individual mosquitoes, highlighting recent inbreeding in Kenyan (grey) and colony (black) mosquitoes. G, Ghana; K, Kisumu; M, Mali; P, Pimperena.

* See broken figure references as “??”, and no figure labels under the figures, compared to (5)
  + Generates errors pandoc-fignos: Bad reference: @fig:figure-test6a. and pandoc-fignos: Bad reference: @fig:figure-test6b.
* Want to remove visible table cell borders
* Markdown table-based layout cannot be used to place the figures next to each other without breaking the Pandoc separate-paragraph rule

### 7) Separate figures using images inline (not Manubot figures), no resizing

 

**a**, Stairway plot of changes in population size over time. Absolute values of time and Ne are shown on alternative axes as a range of values, assuming lower and upper limits for the mutation rate μ as 2.8 × 10−9 and 5.5 × 10−9, respectively, and t = 11 generations per year. ka, thousand years ago. **b**, Runs of homozygosity (ROH) in individual mosquitoes, highlighting recent inbreeding in Kenyan (grey) and colony (black) mosquitoes. G, Ghana; K, Kisumu; M, Mali; P, Pimperena.

* images can be used to simulate figures, at the cost of all the figure management features of Manubot (assuming the features worked as intended)
* references for figures would have to be managed manually. A figure reference has not been added here, and would have to be added and maintained manually (outside of Manubot’s system, by hard-coding the figure identifier/label, e.g. Figure 1, in the Markdown)

### 8) Separate figures using images inline (not Manubot figures), resized to 3in and 4in width, rearranged

 

**a**, Stairway plot of changes in population size over time. Absolute values of time and Ne are shown on alternative axes as a range of values, assuming lower and upper limits for the mutation rate μ as 2.8 × 10−9 and 5.5 × 10−9, respectively, and t = 11 generations per year. ka, thousand years ago. **b**, Runs of homozygosity (ROH) in individual mosquitoes, highlighting recent inbreeding in Kenyan (grey) and colony (black) mosquitoes. G, Ghana; K, Kisumu; M, Mali; P, Pimperena.

* Labels **a** and **b** in the description are manually coded (these are not Manubot-managed figures, only Markdown images)
* See that subfigure **b** appears before (left of) subfigure **a**, intentionally swapped around
* See that the **b** label is larger than the **a** label, because the whole **b** image has been made larger than **a**, intentionally
* embedding subfigure labels in the image means that image resizing and rearranging is frustrated
  + new images would need to be generated with new labels, or the SVG code modified by hand

### 9) Separate figures using images (not Manubot figures) with HTML table-based layout, resized to 3in and 4in width

If the labels were not embedded in the image, the labels would naturally need be added to the Markdown in some way, perhaps as figure “captions”.

Imagine that the two images below did not contain their **a** and **b** labels in their top-left corners:

**Figure 42**

Figure a: Stairway Plot

Figure a: Stairway Plot

Figure b: ROH Scatter Plot

Figure b: ROH Scatter Plot

**a**, Stairway plot of changes in population size over time. Absolute values of time and Ne are shown on alternative axes as a range of values, assuming lower and upper limits for the mutation rate μ as 2.8 × 10−9 and 5.5 × 10−9, respectively, and t = 11 generations per year. ka, thousand years ago. **b**, Runs of homozygosity (ROH) in individual mosquitoes, highlighting recent inbreeding in Kenyan (grey) and colony (black) mosquitoes. G, Ghana; K, Kisumu; M, Mali; P, Pimperena.

* This is similar to (1), except with the layout approach of (5)
* The **a** and **b** in the descriptions are taken from the Manubot figure tag labels, specified in Markdown.
* The label **Figure 42** was added manually. Only the subfigure identifiers/labels are managed by Manubot. There is no “Manubot figure 42”, so that label would have to be managed/maintained manually.

### 10) Single figure with subfigures as labelled images in a HTML table-based layout

Imagine that the following two images do not contain labels **a** and **b** in their top-left corners:

**Figure 1**

**a**



**b**



**1a**, Stairway plot of changes in population size over time. Absolute values of time and Ne are shown on alternative axes as a range of values, assuming lower and upper limits for the mutation rate μ as 2.8 × 10−9 and 5.5 × 10−9, respectively, and t = 11 generations per year. ka, thousand years ago. **1b**, Runs of homozygosity (ROH) in individual mosquitoes, highlighting recent inbreeding in Kenyan (grey) and colony (black) mosquitoes. G, Ghana; K, Kisumu; M, Mali; P, Pimperena.

* This is similar to (3), except with the layout approach of (5)
* The label **Figure 1** was added manually, but the 1 part is automatic.
* The subfigure labels **a** and **b** outside the image were added manually.
* In the description, the 1 part (of 1a and 1b) is automatic, whereas the “a” and “b” parts are added manually, i.e. hard-coded in the Markdown.

## References

1. Himmelstein, D. S. *et al.* Sci-Hub provides access to nearly all scholarly literature. *eLife* **7**, (2018).

2. Beaulieu-Jones, B. K. & Greene, C. S. Reproducibility of computational workflows is automated using continuous analysis. *Nat Biotechnol* **35**, 342–346 (2017).

3. Heaven, D. Bitcoin for the biological literature. *Nature* **566**, 141–142 (2019).

4. S, c. Plan S: Accelerating the transition to full and immediate Open Access to scientific publications. (2018).

5. Suber, P. *Open access*. (MIT Press, 2012).

6. Himmelstein, D. S. *et al.* *Open collaborative writing with Manubot*. <https://greenelab.github.io/meta-review/> (2020).

7. Ching, T. *et al.* Opportunities and obstacles for deep learning in biology and medicine. *Journal of The Royal Society Interface* **15**, 20170387 (2018).

8. Himmelstein, D. S. *et al.* Open collaborative writing with Manubot. *PLOS Computational Biology* **15**, e1007128 (2019).

9. Himmelstein, D. manubot/rootstock GitHub repository. *GitHub* <https://github.com/manubot/rootstock> (2019).