

Edit me to practice contributing to a collaborative Manubot manuscript

test edit - hello world. This manuscript ([permalink](#)) was automatically generated from [manubot/try-manubot@d43a3fd](#) on November 8, 2020.

Authors

- **John Doe**

 [XXXX-XXXX-XXXX-XXXX](#) ·  [johndoe](#) ·  [johndoe](#)

Department of Something, University of Whatever · Funded by Grant XXXXXXXX

- **Jane Roe**

 [XXXX-XXXX-XXXX-XXXX](#) ·  [janeroe](#)

Department of Something, University of Whatever; Department of Whatever, University of Something

Abstract

Manubot is an open source tool for writing manuscripts on GitHub in markdown format. Manubot applies the git-based software workflow to scholarly writing, enabling enhanced transparency, collaboration, automation, and reproducibility.

This manuscript is a Manubot demo, intended to give users a playground to practice using Manubot. Everyone is encouraged to try writing with Manubot by editing this manuscript.

Manubot is described in the paper titled “Open collaborative writing with Manubot” [\[1\]](#).

Test my understanding

Just a paper [\[2\]](#)

Main text

Lorem ipsum text [\[3\]](#) is a strong introduction for any manuscript.

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.

Manubot makes it easy to cite this manuscript [\[4\]](#). It has been used to write several manuscripts that are now preprints on *bioRxiv* [\[5,6,7,8\]](#). Notice that only [\[7\]](#) has the correct name of the preprint server. Manubot allows authors to overwrite reference information, in this case with a BibTeX file.

Lorem ipsum also makes a strong conclusion [\[9\]](#) here is another reference to see if duplicate references are picked up by the program if the second author uses a different identifier for the same reference like here with the immediately preceding reference [10\]](#). Did it work?

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.

I’m wondering about how editing this file works within the web browser.

References

1. Open collaborative writing with Manubot

Daniel S. Himmelstein, Vincent Rubinetti, David R. Slochower, Dongbo Hu, Venkat S. Malladi, Casey S. Greene, Anthony Gitter
Manubot (2020-05-25) <https://greenelab.github.io/meta-review/>

2. Proteogenomics connects somatic mutations to signalling in breast cancer

Philipp Mertins, D. R. Mani, Kelly V. Ruggles, Michael A. Gillette, Karl R. Clauser, Pei Wang, Xianlong Wang, Jana W. Qiao, Song Cao, Francesca Petralia, ... Steven A. Carr
Nature (2016-06) <https://www.nature.com/articles/nature18003>
DOI: [10.1038/nature18003](https://doi.org/10.1038/nature18003)

3. Lorem ipsum

Wikipedia
(2020-10-23) https://en.wikipedia.org/w/index.php?title=Lorem_ipsum&oldid=985101662

4. Edit me to practice contributing to a collaborative Manubot manuscript

John Doe, Jane Roe
Manubot (2020-10-08) <https://manubot.github.io/try-manubot/>

5. GimmeMotifs: an analysis framework for transcription factor motif analysis

Niklas Bruse, Simon J. van Heeringen
bioRxiv (2018-11-20) <https://doi.org/gfxrkc>
DOI: [10.1101/474403](https://doi.org/10.1101/474403)

6. Plasmids for independently tunable, low-noise expression of two genes

João P. N. Silva, Soraia Vidigal Lopes, Diogo J. Grilo, Zach Hensel
bioRxiv (2019-01-09) <https://doi.org/gfs47c>
DOI: [10.1101/515940](https://doi.org/10.1101/515940)

7. Scaling tree-based automated machine learning to biomedical big data with a dataset selector

Trang T. Le, Weixuan Fu, Jason H. Moore
bioRxiv (2018-12) <https://www.biorxiv.org/content/10.1101/502484v1>
DOI: [10.1101/502484](https://doi.org/10.1101/502484)

8. Genotyping structural variants in pangenome graphs using the vg toolkit

Glenn Hickey, David Heller, Jean Monlong, Jonas A. Sibbesen, Jouni Sirén, Jordan Eizenga, Eric T. Dawson, Erik Garrison, Adam M. Novak, Benedict Paten
bioRxiv (2019-10-29) <https://doi.org/gf3jfm>
DOI: [10.1101/654566](https://doi.org/10.1101/654566)

9. Hypothermic machine perfusion in kidney transplantation.

Julie De Deken, Peri Kocabayoglu, Cyril Moers
Current opinion in organ transplantation (2016-06)
<https://www.ncbi.nlm.nih.gov/pubmed/26945319>
DOI: [10.1097/mot.0000000000000306](https://doi.org/10.1097/mot.0000000000000306) · PMID: [26945319](https://pubmed.ncbi.nlm.nih.gov/26945319/)

10. Hypothermic machine perfusion in kidney transplantation

Julie De Deken, Peri Kocabayoglu, Cyril Moers

Current Opinion in Organ Transplantation (2016-06) <https://doi.org/f8mswg>
DOI: [10.1097/mot.0000000000000306](https://doi.org/10.1097/mot.0000000000000306) · PMID: [26945319](https://pubmed.ncbi.nlm.nih.gov/26945319/)