

# Title of this Paper

Your Name

## Abstract

Pizza [1] is an understudied yet widely utilized implement for delivering in-vivo *Solanum lycopersicum* based liquid mediums in a variety of next-generation mastications studies. Here<sup>1</sup> we describe a de novo approach for large scale *T. aestivum* assemblies based on protein folding that drastically reduces the generation time of the mutation rate.

Sentence blah.<sup>2</sup>

## Algorithm

$$f(x) = pizza^2$$

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<sup>1</sup>In-text foot note

<sup>2</sup>I'm a footnote: <https://github.com/MrShoenel/md-2-pdf-pandoc-report-bib>

## References

- [1] M. Pizza *et al.*, “Identification of vaccine candidates against serogroup b meningococcus by whole-genome sequencing,” *Science*, vol. 287, no. 5459, pp. 1816–1820, 2000.