

# Semester Project [rename]

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## 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 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.

## Introduction

## Methods

## Comparisons

## Example Analysis

## Conclusions

## 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.