Scalable microbial strain inference in metagenomic data using StrainFacts

Byron J. Smith A,B

Xiangpeng Li C

Adam Abate C,D

Zhou Jason Shi A,D

Katherine S. Pollard A,B,D,\*

A The Gladstone Institute of Data Science and Biotechnology, San Francisco, CA  
B Department of Epidemiology and Biostatistics, University of California, San Francisco, CA  
C Department of Bioengineering and Therapeutic Sciences, University of California, San Francisco, CA  
D Chan-Zuckerberg Biohub, San Francisco, CA

\* Corresponding author:  
[katherine.pollard@gladstone.ucsf.edu](mailto:katherine.pollard@gladstone.ucsf.edu)