# Marcus William Fedarko

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Education

Ph.D. Student, Computer Science University of California, San Diego

GPA: 3.60/4.0

M.S., Computer Science University of California, San Diego

GPA: 3.60/4.0

B.S. WITH HIGH HONORS, Computer Science University of Maryland

GPA: 3.81/4.0

9/2014-5/2018 College Park, MD

9/2018-Present

San Diego, CA

9/2018-6/2022

San Diego, CA

# Refereed Publications

- 7. **Fedarko MW**, Kolmogorov M, and Pevzner PA (2022). "Analyzing rare mutations in metagenomes assembled using long and accurate reads." *Genome Research*, 32(11-12):2119–2133.
- 6. Cantrell K\*, Fedarko MW\*, Rahman G, McDonald D, Yang Y, Zaw T, Gonzalez A, Janssen S, Estaki M, Haiminen N, Beck KL, Zhu Q, Sayyari E, Morton JT, Armstrong G, Tripathi A, Gauglitz JM, Marotz C, Matteson NL, Martino C, Sanders JG, Carrieri AP, Song SJ, Swafford AD, Dorrestein PC, Andersen KG, Parida L, Kim H-C, Vázquez-Baeza Y, and Knight R (2021). "EMPress Enables Tree-Guided, Interactive, and Exploratory Analyses of Multi-omic Data Sets." mSystems, 6(2):e01216-20. (\* = contributed equally)
- 5. Huey SL, Jiang L, **Fedarko MW**, McDonald D, Martino C, Ali F, Russell DG, Udipi SA, Thorat A, Thakker V, Ghugre P, Potdar RD, Chopra H, Rajagopalan K, Haas JD, Finkelstein JL, Knight R, and Mehta S (2020). "Nutrition and the Gut Microbiota in 10- to 18-Month-Old Children Living in Urban Slums of Mumbai, India." mSphere, 5(5):e00731-20.
- 4. Fedarko MW, Martino C, Morton JT, González A, Rahman G, Marotz CA, Minich JJ, Allen EA, and Knight R (2020). "Visualizing 'omic feature rankings and log-ratios using Qurro." NAR Genomics and Bioinformatics, 2(2):lqaa023.
- 3. Sanders JG, Nurk S, Salido RA, Minich J, Xu ZZ, Martino C, Fedarko M, Arthur TD, Chen F, Boland BS, Humphrey GC, Brennan C, Sanders K, Gaffney J, Jepsen K, Khosroheidari M, Green C, Liyange M, Dang JW, Phelan VV, Quinn RA, Bankevich A, Chang JT, Rana TM, Conrad DJ, Sandborn WJ, Smarr L, Dorrestein PC, Pevzner PA, and Knight R (2019). "Optimizing sequencing protocols for leaderboard metagenomics by combining long and short reads." Genome Biology, 20(1):226.
- 2. Ghurye J, Treangen T, **Fedarko M**, Hervey WJ, and Pop M (2019). "MetaCarvel: linking assembly graph motifs to biological variants." *Genome Biology*, 20(1):174.
- Meisel JS, Nasko DJ, Brubach B, Cepeda-Espinoza V, Chopyk J, Corrada-Bravo H, Fedarko M, Ghurye J, Javkar K, Olson ND, Shah N, Allard SM, Bazinet AL, Bergman NH, Brown A, Caporaso JG, Conlan S, DiRuggiero J, Forry SP, Hasan NA, Kralj J, Luethy PM, Milton DK, Ondov BD, Preheim S, Ratnayake S, Rogers SM, Rosovitz MJ, Sakowski EG, Schliebs NO, Sommer DD, Ternus KL, Uritskiy G, Zhang SX, Pop M, and Treangen TJ (2018). "Current progress and future opportunities in applications

of bioinformatics for biodefense and pathogen detection: Report from the Winter Mid-Atlantic Microbiome Meet-up, College Park, MD January 10th, 2018." Microbiome, 6(1):197.

# Outreach Presentations

- 2. "Visualizing, Exploring, and Understanding Microbiome Sequencing Data." UC San Diego CSE Research Open House, 1/2020.
- 1. "Visualizing Metagenomic Assembly Graphs, Doing Undergrad Research at UMD, Applying to Grad Schools, and probably other stuff along the way." Guest talk for CMSC 396H (U. of Maryland undergraduate honors seminar), 4/2018.

#### Service

- 4. Mentor, UC San Diego GradWIC mentorship program 10/2021-Present
- 3. Moderator, QIIME 2 forum (https://forum.qiime2.org) 3/2020-Present
- 2019, 2020, 2021, 2022 2. Co-organizer, UC San Diego CSE Visit Day
- 1. Code Review (Co-)Organizer, Knight Lab 12/2018-8/2020

# Research Experience

#### Graduate Student Researcher

# 9/2018-Present University of California, San Diego

- San Diego, CA
- Designing software for the analysis of metagenomic sequencing data and other forms of "'omic" data.
- Assisting with various software and analysis projects.

#### Research Intern

6/2016-8/2018

# University of Maryland

College Park, MD

• Designed MetagenomeScope, an interactive visualization tool for metagenome assembly graphs: see https://marbl.github.io/MetagenomeScope.

# Teaching Experience

#### TEACHING ASSISTANT

#### University of California, San Diego

San Diego, CA

• CSE 282: Introduction to Bioinformatics Algorithms 1/2022-3/20221/2021 - 3/2021• CSE 282: Introduction to Bioinformatics Algorithms

### Course Assistant

# Marine Biological Laboratory

Woods Hole, MA

7/2018

- STAMPS: Strategies and Techniques for Analyzing Microbial 7/2018-8/2018 Population Structures
- MOLE: Workshop on Molecular Evolution

TEACHING ASSISTANT

# University of Maryland

College Park, MD

• CMSC 330: Organization of Programming Languages

8/2016-12/2016

# **Professional** Experience

#### STUDENT STAFF WRITER

1/2015-9/2017

#### University of Maryland Dept. of Computer Science College Park, MD

• Composed and edited articles for the department's website and other media.

• Assisted with the logistics of various department outreach functions.

STUDENT INTERN 5/2013 - 8/2014Axiometric Columbia, MD

- Designed a graphical interface to an RF propagation model to assist clients in planning deployments of mesh networks of utility meters.
- Aided in the creation and maintenance of other utility meter deployment management software.

# INTERN SOFTWARE ENGINEER

7/2012-8/2012

# Battlefield Telecommunications Systems

Columbia, MD

- Designed a web interface to monitor the connection strength of radio devices.
- Helped integrate this functionality into the company's existing network management user interface.

# $\begin{array}{c} \textbf{Honors and} \\ \textbf{Awards} \end{array}$

9. University of Maryland CMNS Dean's List	2014 – 2018
8. University of Maryland Honors College University Honors Citation	2017
7. Rita Colwell Travel Fellowship	2017
6. Travel Award, U. of Michigan "Explore Graduate Studies" Workshop	2017
5. John D. Gannon Endowed Scholarship	2017
4. Corporate Partners in Computing Scholarship	2016, 2017
3. Omicron Delta Kappa National Leadership Honor Society	2016
2. Northrop Grumman Scholarship for Employees' Children	2014
1. University of Maryland Dean's Scholarship	2014