

DOMENICK J. BRACCIA

Personal Website: dbraccia.com

 [Github](#)  [Google Scholar](#)  [LinkedIn](#)  [Twitter](#)

August 15th, 2023

I am Computational Biologist with 6 years (5 academic + 1 industry) of experience with high-throughput genomic, transcriptomic, and proteomic sequencing data. I have diverse knowledge across biology, statistics, programming, mathematics, and data science.

PERSONAL INFORMATION

- Phone: (267) 664-5315 | Email: domenick@dbraccia.com
- Mailing Address: 1151 Canterbury Dr. Lansdale, PA 19446

EDUCATION

PhD	Computational Biology, Bioinformatics, and Genomics University of Maryland, College Park, MD	July 2022
BS	Physics, Minor in Math Pennsylvania State University, University Park, PA	August 2016

WORK EXPERIENCE

Scientist, Computational Biology August 2023 – Present
Cabaletta Bio, Philadelphia, PA

Bioinformatics Consultant May 2023 – Present
C M Cherry Consulting, Philadelphia, PA

Data Scientist – Bioinformatics July 2022 – Feb 2023
Scipher Medicine, Waltham, MA | References available upon request

- Designed bioinformatic pipelines to process, analyze and assess the quality of transcriptomics data
- Built cloud computing infrastructure to aid high throughput hypothesis testing and biomarker discovery
- Designed and trained classification model predicting pathology of early Rheumatoid Arthritis (RA) patients from whole blood RNA-seq gene expression data
- Performed feature selection of proteomics features correlated with early RA clinical endpoints

Graduate Research Assistant August 2017 – July 2022
University of Maryland, College Park, MD | Dr. Brantley Hall Lab

- Collected, integrated, mined, and analyzed large multi-omic datasets to characterize the progression of IBD and researched microbiome therapy options for neonatal jaundice and colorectal cancer
- Utilized DNA and RNA sequence data for hypothesis generation and experimental design
- Oversaw team Git repository for a collaborative software project to validate wearable medical device
- Performed statistical power analysis for a successful \$250,000 grant application using repeated measures ANOVA

Domenick J. Braccia

PUBLICATIONS

6. Hall B, Levy S, Default-Tompson K, Minabou G, Weiss A, **Braccia DJ**, Jenkins C, Yang Y, Arp G, Abeysinghe S, Jermain M. Discovery of the gut microbial enzyme responsible for bilirubin reduction to urobilinogen. *bioRxiv*. 2023:2023-02. **[Preprint]** 2023
5. **Braccia DJ**, Ndjite GM, Weiss A, Levy S, Abeysinghe S, Jiang X, Pop M, Hall B. Gut Microbiome–Wide Search for Bacterial Azoreductases Reveals Potentially Uncharacterized Azoreductases Encoded in the Human Gut Microbiome. *Drug Metabolism and Disposition*. 2023 Jan 1;51(1):142-53. 2023
4. **Braccia DJ**, Jiang X, Pop M, Hall AB. The capacity to produce hydrogen sulfide (H₂S) via cysteine degradation is ubiquitous in the human gut microbiome. *Frontiers in Microbiology*. 2021 Oct 20;12:705583. 2021
3. Olson ND, Kumar MS, Li S, **Braccia DJ**, Hao S, Timp W, Salit ML, Stine OC, Bravo HC. A framework for assessing 16S rRNA marker-gene survey data analysis methods using mixtures. *Microbiome*. 2020 Dec;8(1):1-8. 2020
2. Wagner J, Kancherla J, **Braccia DJ**, Matsumara J, Felix V, Crabtree J, Mahurkar A, Bravo HC. Interactive exploratory data analysis of Integrative Human Microbiome Project data using Metaviz. *F1000Research*. 2020;9. 2020
1. Paulson JN, Olson ND, **Braccia DJ**, Wagner J, Talukder H, Pop M, Bravo HC, Paulson MJ. Package ‘metagenomeSeq’. 2013

HONORS, AWARDS & RECOGNITION

- | | |
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| Front cover art of <i>Drug Metabolism and Disposition</i> Vol 51. Issue 1 | 2023 |
| National Science Foundation Research Traineeship, COMBINE , \$34,000 | 2018 – 2021 |
| Dean’s Fellowship, \$5,000 | 2020 |
| COMBINE Travel Grant, \$1,500 | 2018 |
| Dean’s Fellowship, \$5,000 | 2018 |
| Dean’s Fellowship, \$5,000 | 2017 |
| Summer Research Experience Internship, \$5,000 | 2015 |

ONLINE COURSEWORK & CERTIFICATIONS

Codecademy, <https://www.codecademy.com>

- Analyze Data with Python [April 2023](#)
- Full Stack Engineer Career Path (In Progress)
- Software Engineering for Data Scientists (In Progress)

TECHNICAL SKILLS

- **Programming:** R, Python, Bash, SQL
- **OS:** Linux, MacOS, Windows
- **Bioinformatics Pipeline Management:** NextFlow, Snakemake
- **Version Control:** GitHub, GitLab
- **Cloud Computing:** AWS, SLURM
- **Visualization:** Adobe Illustrator

GRANTS

Employing smart underwear to measure gut microbial hydrogen sulfide production 2022 – 2024

Grant Number: R21 – 1R21DK132310

Role: Other Significant Contributor (Statistical Analysis Plan)

Developing a Smart Underwear device to help living well with IBS 2021 – 2022

Grant Type: Maryland Innovation Initiative (MII)

Role: Other Significant Contributor (Statistical Analysis Plan)

PRESENTATIONS AND POSTERS

American Society of Microbiology, Washington D.C. 2022

Keystone Symposium: Harnessing the Microbiome for Disease Prevention and Therapy, Virtual 2021

Bioconductor Europe, Virtual 2020

TEACHING EXPERIENCE

BSCI 441 – Mammalian Physiology Lab 2019 – 2020

University of Maryland, College Park, PA

- Oversaw 10 rat surgical operations including cannulation, IV administration, and nerve stimulation
- Developed at-home teaching materials, labs, and lectures in response to COVID

BSCI 330 – Cell Biology and Physiology Lab 2017 – 2019

University of Maryland, College Park, PA

- Guided 200 students in cell biology procedures such as SDS-PAGE gel electrophoresis, protein quantification, and *in vivo* administration of psychoactive drugs to model organisms
- As a senior TA – coached junior TAs in teaching strategies and lab protocols (mentioned above)

PHYS 220 – Electricity and Magnetism Lab 2016

Pennsylvania State University, University Park, PA

- Led 50 students in lab techniques and lectures on the core principles of Electricity and Magnetism