

I am a PhD in computational biology using data science to impact society.

EDUCATION	Massachusetts Institute of Technology , <i>Cambridge, MA</i> Ph.D., Biological Engineering, January 2019	2014 – 2019
	Columbia University , <i>New York, NY</i> B.S., Biomedical Engineering	2009 – 2013
EXPERIENCE	Biobot Analytics <i>Data Scientist</i>	2019 – present
	Biobot Analytics measures opioids in sewage to estimate consumption in cities. As Biobot's first data scientist, I lead our data analytics and visualization efforts, communicate the value of our novel opioid data, and serve as the main point of contact for our customers, public health and city officials.	
RESEARCH	Massachusetts Institute of Technology Supervisor: Eric J. Alm, Ph.D. Ph.D., <i>Department of Biological Engineering</i> (2015 – Jan. 2019) Postdoc (Feb. 2019 – Apr. 2019)	2015 – 2019
	I studied the relationship between the microbiome and health and disease, mining large clinical and biological datasets to extract scientific insight.	
	Columbia University Supervisor: Samuel L. Sia, Ph.D. <i>Molecular and Microscale Bioengineering Laboratory</i>	2011 – 2013
	As an undergraduate researcher, I worked on developing a point-of-care microfluidic device to diagnose multi-drug resistant tuberculosis in resource-limited settings.	
	Ecole Polytechnique Supervisor: Cedric Norais, Ph.D. <i>Laboratoire de Biochimie</i>	Summer 2012
	As an international undergraduate research intern, I studied the acquired-immunity CRISPR system in <i>E. coli</i> .	
TEACHING EXPERIENCE	Teaching Assistant 20.106 Systems Microbiology, <i>Massachusetts Institute of Technology</i>	Fall 2015
	I was a TA for seven advanced undergraduate students in a new course on the human microbiome, emerging disease, phylogenetics, and host-microbe interactions.	
	Lecturer Biomedical Equipment Technology Department, <i>University of Puthisastra</i> Engineering World Health, <i>Phnom Penh, Cambodia</i>	2013 – 2014
	As a Luce Scholar, I was one of the first lecturers for Engineering World Health's new Associate Bachelors program in Biomedical Equipment Technology at the University of Puthisastra, a private university in Phnom Penh.	
	Teaching Assistant The Art of Engineering, <i>Columbia University</i>	2012 – 2013
	My senior year at Columbia, I TAed the biomedical engineering section of the introductory engineering course for freshmen.	

PUBLICATIONS

1. Sole author. (*in press*)
“Data detectives, self-love, and humility: a research parasite’s perspective.”
GigaScience.
2. Middle author (of seven). (*accepted*)
“Rapid assessment of opioid exposure and treatment in cities through robotic collection and chemical analysis of wastewater.”
Journal of Medical Toxicology.
3. Second author (of nine). (2019)
“24-hour multi-omics analysis of residential sewage reflects human activity and informs public health.”
bioRxiv, doi: [10.1101/728022](https://doi.org/10.1101/728022).
4. First author (of six). (2019)
“Framework for rational donor selection in fecal microbiota transplant clinical trials.”
PLoS ONE, doi: [10.1371/journal.pone.0222881](https://doi.org/10.1371/journal.pone.0222881).
5. Middle author (of hundreds). (2019)
“Reproducible, interactive, scalable and extensible microbiome data science using QIIME 2.”
Nature Biotechnology, doi: [10.1038/s41587-019-0209-9](https://doi.org/10.1038/s41587-019-0209-9).
6. Second author (of twelve). (2019)
“Multi-site sampling and risk prioritization reveals the public health relevance of antibiotic resistance genes found in sewage environments.”
bioRxiv, doi: [10.1101/562496v1](https://doi.org/10.1101/562496v1).
7. First author (of nine). (2019)
“Aerodigestive sampling reveals altered microbial exchange between lung, oropharyngeal, and gastric microbiomes in children with impaired swallow function.”
PLoS ONE, doi: [10.1371/journal.pone.0216453](https://doi.org/10.1371/journal.pone.0216453).
8. Co-second author (of ten). (2019)
“A practical guide to methods controlling false discoveries in computational biology.”
Genome Biology, doi: [10.1186/s13059-019-1716-1](https://doi.org/10.1186/s13059-019-1716-1).
9. Middle author (of forty). (2018)
“Predictability and persistence of prebiotic dietary supplementation in a healthy human cohort.”
Scientific Reports. doi: [10.1038/s41598-018-30783-1](https://doi.org/10.1038/s41598-018-30783-1).
10. Second author (of three). (2018)
“Correcting for batch effects in case-control microbiome studies.”
PLoS Computational Biology. doi: [10.1371/journal.pone.0176335](https://doi.org/10.1371/journal.pone.0176335).
11. Sole author. (2018)
“Meta-analysis generates and prioritizes hypotheses for translational microbiome research.”
Microbial Biotechnology. doi: [10.1111/1751-7915.13047](https://doi.org/10.1111/1751-7915.13047).
12. First author (of five). (2017)
“Meta-analysis of gut microbiome studies identifies disease-specific and shared responses.”
Nature Communications. doi: [10.1038/s41467-017-01973-8](https://doi.org/10.1038/s41467-017-01973-8).

- Received PSB Award for Rigorous Secondary Data Analysis

13. Second author (of three). (2017)
“dbOTU3: A new implementation of distribution-based OTU calling.”
PloS ONE. doi: [10.1371/journal.pone.0176335](https://doi.org/10.1371/journal.pone.0176335).

Non-peer reviewed

14. [Tutorial] “Updating your QIIME 2 plugin.” (2019) *QIIME 2 developer documentation*.
15. [Blog post] “Scientific discovery from a clinical study: surprises from the lung and stomach microbiomes.” (2019) *Nature Microbiology Community Forum*. go.nature.com/30rx4VZ.
16. [Tutorial] “QIIME 2 for Experienced Microbiome Researchers.” (2018) *QIIME 2 documentation*.
17. [Tutorial] “Developing a plugin for dummies.” (2018) *QIIME 2 developer documentation*.
18. [Tutorial] “Publishing your plugin on conda.” (2018) *QIIME 2 developer documentation*.
19. [Blog post] “Fuzzy zeros in percentile normalization method to correct for batch effects.” (2018) *microBEnet: the microbiology of the Built Environment network*.
20. [Blog post] “Beyond dysbiosis: disease-specific and shared microbiome responses to disease.” (2017) *Nature Microbiology Community Forum*. go.nature.com/2As9meL.
21. [Dataset] “MicrobiomeHD: the human gut microbiome in health and disease.” (2017) *Zenodo*. doi: [10.5281/zenodo.1146764](https://doi.org/10.5281/zenodo.1146764)

Other

22. [Blog post] “A well-kept secret for finding a job post-PhD.” (*in press*) *MIT Graduate Admissions Blog*.
23. [Blog post] “Learning to Engage in Deep Conversations.” (2018) *MIT Graduate Admissions Blog*.
24. [Resource] “Graduate student support resources flowchart.” (2018) *MIT Division of Student Life*.

ORAL PRESENTATIONS

1. “24-hour multi-omics analysis of residential sewage reflects human activity and informs public health.” American Chemical Society National Meeting. Wastewater-Based Epidemiology: Opportunities, Challenges & Applications to Public Health & Safety. San Diego, CA. Aug. 2019. *Invited*.
2. “Intro to microbiome data visualization.” Cornell University 2019 Microbiome Hackathon: Microbiome Hack. Ithaca, NY. Apr. 2019. *Invited*.
3. “Framework for rational donor selection in fecal microbiota transplant clinical trials.” International Conference on Microbiome Engineering (ICME 2018). Boston, MA. Nov. 2018. *Invited*.
4. “Predictive power of the microbiome.” Science on Tap! Boston College Department of Biology seminar series. Boston, MA. Aug. 2018. *Invited*.
5. “Distribution-based methods to increase power and reduce redundancy in microbiome data.” Teaching and Developing QIIME 2 Workshop. San Diego, CA. May 2018. *Selected*.
6. “Meta-analysis to identify consistent disease-associated microbiome shifts.” MIT-Harvard Microbiome Symposium. Cambridge, MA. March 2018. *Selected*.

SOFTWARE	Percentile normalization		
	Correcting batch effects in case-control microbiome studies.		
	Python implementation (<i>contributor</i>) and QIIME 2 plugin (<i>developer</i>).		
	Distribution-based OTU calling		
	New implementation of Preheim <i>et al.</i> 's distribution-based OTU clustering algorithm.		
	Python implementation (<i>contributor</i>) and QIIME 2 plugin (<i>developer</i>).		
	Amplicon sequencing pipeline		
	End-to-end pipeline to process 16S data.		
	Python implementation (<i>co-developer</i>).		
SKILLS	Technical: Python (pandas, seaborn, matplotlib, scikit-learn, etc); git and GitHub; AWS (S3, EC2, Glacier, IAM); bash; LaTeX; R (proficient); data wrangling and exploration; machine learning; NGS analysis (16S, metagenomics); untargeted metabolomics analysis.		
	Non-technical: conflict management; student advocacy; project management; oral and written communication; diversity, equity, and inclusion.		
	Languages: English (native), French (fluent), Spanish (conversational), Khmer (beginner)		
FELLOWSHIPS & AWARDS	Fellowships		
	Siebel Scholars Foundation		2018
	<i>Siebel Scholar Class of 2019</i>		
	National Defense Science and Engineering Graduate Fellowship		2015 – 2018
	<i>NDSEG Recipient</i>		
	National Science Foundation Graduate Research Fellowship		2015
	<i>Honorable Mention</i>		
	Henry Luce Foundation		2013 – 2014
	<i>Luce Scholar, Engineering World Health Cambodia</i>		
	Awards		
	PSB Award for Rigorous Secondary Data Analysis		2019
	<i>Junior Research Parasite</i>		
	MIT Graduate Women of Excellence		2017
	Salutatorian		2013
	<i>Columbia University Fu Foundation School of Engineering and Applied Science</i>		
	Richard Skalak Award in Biomedical Engineering		2013
	<i>Columbia University Department of Biomedical Engineering</i>		
	Robert E. and Claire S. Reiss Prize in Biomedical Engineering		2013
	<i>Columbia University Department of Biomedical Engineering</i>		
	King's Crown Bronze Leadership Award		2012
	<i>Columbia University</i>		
	Tau Beta Pi, The Engineering Honor Society		2012

LEADERSHIP
& SERVICE

Academic and professional

Bluebonnet Data	2019 – present
Volunteer data analytics for state and local campaigns <i>Team lead and data scientist</i>	
MIT Microbiome Club	2015 – 2018
<i>Co-founder, president, and executive board member</i>	
MIT-Harvard Microbiome Symposium	2016 and 2017
<i>Co-founder and organizing committee</i>	
MIT Biotech Group	2017 – 2018
<i>Beyond the Bench Initiative board member</i>	

Departmental and MIT

Biological Engineering Department Visiting Committee	2018
<i>Graduate student representative</i>	
Graduate Student Advisory Group for Engineering (GradSAGE)	2017 – 2018
Advisory group to the Dean of the School of Engineering <i>Advisor/Advisee Relations Subcommittee</i>	
MIT Graduate Student Council	2017 – 2019
Diversity and Inclusion Subcommittee	
<i>Vice Chair, Department and Classroom Inclusion co-coordinator</i>	
BE Resources for Easing Friction and Stress (BE REFS)	2016 – 2019
<i>Confidential conflict management coach and graduate student advocate</i>	
BE Graduate Student Board	2015 – 2018
<i>Diversity Chair</i>	
<i>Co-founder, BE Application Assistance Program</i>	
<i>Lead author, BE Departmental Values Statement</i>	
<i>Co-lead, 2016 BE Diversity Survey</i>	

Outreach & Mentorship

MIT Microbiome superUROP	2016 – 2017
<i>Mentor, supervised one undergraduate researcher</i>	
Science Club for Girls, Young Leaders in STEM	2016 and 2017
<i>Volunteer, developed and taught three-day course on microbiology and the human microbiome</i>	
MIT SPLASH	2015
<i>Volunteer, “Microbiome 101: What’s in your poop?”</i>	
E3: Empowering, Encouraging, and Eliminating Barriers for Women in STEM	2015
<i>Mentor (2015), guest presenter (2016)</i>	
ESL Program for MIT Service Employees	2015 – 2018
<i>Math GED tutor</i>	

Reviewer

MIT Summer Research Program (MSRP) Reviewer	2018 and 2019
MIT Committed to Caring Selection Committee	2017
MIT IDEAS Global Challenge Reviewer	2015 – 2018

EXTRA-
CURRICULARS

MIT Interfaith Dialogue Program
Addir fellow

2017 – 2018

sMITe, MIT Women's Ultimate Frisbee Team
B-team captain (2014 – 2015)

2014 – 2019