Date of Preparation: September 20, 2021

Academic Appoin	tments and Other Work Experience
06/2021 – Present	Infectious Diseases Division, Columbia Ur

06/2021 - Present	Infectious Diseases Division, Columbia University Associate Research Scientist	New York, NY
12/2018 – Present	Infectious Diseases Division, Columbia University Postdoctoral Research Fellow	New York, NY
06/2017 – 11/2018	Infectious Diseases Division, Columbia University Postdoctoral Research Scientist	New York, NY
08/2012 – 05/2017	Dept. of Earth and Environ. Eng., Columbia University Graduate Research Assistant	New York, NY
09/2016 – 12/2016	Dept. of Earth and Environ. Eng., Columbia University Teaching Assistant	New York, NY
08/2011 – 05/2012	Dept. of Biological Sciences, Carnegie Mellon University Teaching Assistant	Pittsburgh, PA
10/2010 – 05/2012	Dept. of Chemistry, Carnegie Mellon University Undergraduate Research Assistant	Pittsburgh, PA
06/2010 – 08/2010	Proctor & Gamble, Home Care Research & Development Products Research Intern	Cincinnati, OH

Education

08/2012 – 05/2017	Columbia University, School of Engineering and Applied Sciences	
	MS, May 2014	New York, NY
	PhD, May 2017	New York, NY
	Sponsor: Dr. Kartik Chandran	
	Thesis: "Meta-omics-derived structure, function, and activity of mixed microbial communities driving biological nutrient removal and recovery"	
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Carnegie Mellon University, Mellon College of Science 08/2008-12/2011

BS, December 2011 Pittsburgh, PA

Honors and Awards

2019	Winning Team, Up-Goer Five Thing Symposium, ASM Microbe 2019
	Science communication symposium where two teams presented research
	abstracts using only 1,000 most commonly used English words. I was on the
	winning team (Team People Still Learning) and won both team MVP and best
	phrase substitution based on audience voting.
2019	Outstanding Abstract Award, ASM Microbe 2019

2018	Conference Participation Grant, International Microbiome in HIV Workshop
2017	Herbert H. Kellogg Fellowship, Columbia School of Engineering and Applied
	Sciences
	Recognizing distinguished ability as a teaching assistant
2012 – 2016	Presidential Fellowship, Columbia School of Engineering and Applied Sciences
	Selected by a panel of SEAS faculty for the distinguished Presidential Fellowship
	as an incoming MS/PhD candidate due to "outstanding potential for graduate
	research and training." The fellowship provided full financial support to pursue my
	doctoral research with intellectual freedom in any field within SEAS.
2014 – 2015	International Fellows Program, Columbia School of International and Public
	Affairs
	Selected as part of an exclusive multidisciplinary group of 30 students to
	participate in discussions on the role of the US in international affairs.
2012	Phi Beta Kappa Honors Society induction, Carnegie Mellon University
2012	Phi Kappa Phi Honors Society induction, Carnegie Melon University
2012	University Honors, Carnegie Mellon University
2012	Mellon College of Science Honors, Carnegie Mellon University
2012	Outstanding Undergraduate Research, Carnegie Mellon University
2012	Outstanding Academic Achievement, Carnegie Mellon University
2008 – 2011	Science & Humanities Scholar, Carnegie Mellon University
2011	Howard Hughes Medical Institute (HHMI) Researcher Grant
2011	Howard Hughes Medical Institute (HHMI) Small Undergraduate Research
	Grant (SURG)

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Academic Service

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2019 Biotech without Borders Workshop Instructor

New York, NY

"Genomics in the human body: host genetics and the gut microbiome": Created and taught a 2.5-hour workshop open to the public at the community biotech lab, covering whole-exome data analysis to identify potentially pathogenic variants and metagenomics of the gut microbiome.

2009 - 2011 AmeriCorps Scholar in Service

Pittsburgh, PA

Selected as the Community Service Intern at Carnegie Mellon University through AmeriCorps. I networked with local non-profit organizations and organized targeted local service experiences for all campus student groups.

2010 - 2012 Biology Outreach Programs

Pittsburgh, PA

Introduced local high school students to biology laboratory practices through the Carnegie Mellon University Department of Biological Sciences.

Professional Organizations and Societies

Memberships

2019 – Present Member, Association for Clinical and Translational Science (ACTS)

2018 – Present Member, American Society of Microbiology (ASM)

2016 – Present	Member, American	Association for the	Advancement of Science	(AAAS)

2016 – Present Member, Water Environment Federation (WEF)

2016 – Present Member, NY Water Environment Association (NYWEA)

Ad hoc Reviewer

2020, 2021 Antimicrobial Agents & Chemotherapy

2019 mSphere, Journal of Global Antimicrobial Resistance

2017 Environmental Science & Technology

2016 Chemical Engineering Journal

Fellowship and Grant Support

Current Support

NIH/NIDA U01DA 053949-01 (Uhlemann, Chandran MPI) 01/01/2021 - 12/31/2023

"Optimizing SARS-CoV-2 wastewater-based surveillance in urban and university campus

settings."

Total direct cost: \$3,021,159

Role: Co-Investigator

Past Support

CTSA/Irving Institute TRANSFORM TL1TR001875 (Annavajhala) 12/01/2018 – 11/30/2020

"Microbial metagenomic pathways contributing to chronic inflammation in patients with HIV."

Total direct cost: \$123,220 Role: Postdoctoral Fellow

Teaching Experience and Responsibilities

06/2017 – 06/2019	Microbiome Working Group Research seminar series Organized monthly educational conferences on topics research at Columbia University Medical Center	New York, NY of microbiome
09/2016 – 12/2016	Dept. of Earth and Environ. Eng., Columbia University Teaching Assistant, Aquatic Chemistry	New York, NY
01/2012 – 05/2012	Dept. of Biological Sciences, Carnegie Mellon University Teaching Assistant, Experimental Techniques in Molecular	Pittsburgh, PA r <i>Biology</i>
09/2011 – 12/2011	Dept. of Biological Sciences, Carnegie Mellon University Teaching Assistant, Experimental Biochemistry	Pittsburgh, PA
08/2010 – 12/2010	Dept. of Philosophy, Carnegie Mellon University Teaching Assistant, Ethical Judgments in Professional Life	Pittsburgh, PA

Public Health Activities

04/2019	Supervisory Scientist, Blueprint Earth	Mojave National Preserve, CA
05/2014 - 08/2015	Water Programs Officer, FACE Africa	Rivercess, Liberia; New York, NY

A complete list of publications in My NCBI Bibliography can be found at:

https://www.ncbi.nlm.nih.gov/myncbi/medini.annavajhala.1/bibliography/public/

A. Original, Peer-Reviewed Research Publications

- Annavajhala MK*, Mohri H*, Wang P, Nair M, Zucker JE, Sheng Z, Gomez-Simmonds A, Kelley AL, Tagliavia M, Bedford T, Ho DD, Uhlemann A-C. Emergence and expansion of SARS-CoV-2 B.1.526 after identification in New York. *Nature* 2021. In Press. *contributed equally
- 2. Purpura LJ, Chang M, **Annavajhala MK**, Mohri H, Liu L, Shah J, Cantos A, Medrano N, Laracy J, Scully B, Miko BA, Pereira MR, Tsuji M, Ho DD, Uhlemann A-C, Yin MT. Prolonged severe acute respiratory syndrome coronavirus 2 persistence, attenuated immunologic response, and viral evolution in a solid organ transplant patient. *Am. J. Transplant.* 2021. In Press.
- 3. Grohs E, Hill-Ricciuti A, Kelly N, Messina M, Green DA, Geng W, **Annavajhala MK**, Zachariah P, Mathema B, Uhlemann A-C, Saiman L. *Spa* typing of *Staphylococcus aureus* in a neonatal intensive care unit during routine surveillance. *J. Pediatric Infect. Dis. Soc.* 2021; piab014. doi: 10.1093/jpids/piab014.
- Ahn D, Bhushan G, McConville TH, Annavajhala MK, Soni RK, Wong Fok Lung T, Hofstaedter CE, Shah SS, Chong AM, Castano VG, Ernst RK, Uhlemann A-C, Prince A. An acquired acyltransferase promotes *Klebsiella pneumoniae* ST258 respiratory infection. *Cell Rep.* 2021; 35(9): 109196. doi: 10.1016/j.celrep.2021.109196
- Tomlinson KL, Wong Fok Lung T, Dach F, Annavajhala MK, Gabryszewski SJ, Groves RA, Drikic M, Francoeur NJ, Sridhar SH, Smith ML, Khanal S, Britto CJ, Sebra R, Lewis I, Uhlemann A-C, Kahl BC, Prince AS, Riquelme SA. Staphylococcus aureus induces an itaconate-dominated immunometabolic response that drives biofilm formation. Nat Commun. 2021; 12(1): 1399. doi: 10.1038/s41467-021-21718-y
- Annavajhala MK, May M, Compres G, Freedberg DE, Graham R, Korem T, Stump S, Uhlemann A-C, Abrams JA. Relationship of the Esophageal Microbiome and Tissue Gene Expression and Links to the Oral Microbiome: A Randomized Clinical Trial. Clin. Transl. Gastroenterol. 2020; 11(12): e00235. doi: 10.14309/ctg.0000000000000235
- 7. Kuo SZ, Dettmer K, **Annavajhala MK**, Chong DH, Uhlemann A-C, Abrams JA, Oefner PJ, Freedberg DE. Associations between urinary 3-indoxyl sulfate, a gut microbiome-derived biomarker, and patient outcomes after intensive care unit admission. *J Crit Care* 2021; 63:15-21. doi: 10.1016/j.jcrc.2021.01.005
- 8. Miller EH, Zucker J, Castor D, **Annavajhala MK**, Sepulveda JL, Green DA, Whittier S, Scherer M, Medrano N, Sobieszczyk ME, Yin MT, Kuhn L, Uhlemann A-C. Pre-test symptom duration and cycle threshold values for SARS-CoV-2 reverse transcription-PCR (RT-PCR) predict COVID-19 mortality. *Open Forum Infect. Dis.* 2021; doi: 10.1093/ofid/ofab003
- Agrawal S, Weissbrodt DG, Annavajhala MK, Jensen MM, Carvajal Arroyo JM, Wells G, Chandran K, Vlaeminck SE, Terada A, Smets BF, Lackner S. Time to act – assessing variations in qPCR analyses in biological nitrogen removal with examples from partial nitritation/anammox systems. Water Res. 2021; 190:116604. doi: 10.1016/j.watres.2020.116604

- 10. Gomez-Simmonds A, Annavajhala MK, McConville T, Dietz D, Shoucri S, Laracy J, Rozenberg F, Nelson B, Greendyke W, Furuya EY, Whittier Susan, Uhlemann A-C. Carbapenemase-producing Enterobacterales causing secondary infections during the COVID-19 crisis at a New York City hospital. *J. Antimicrob. Chemother.* 2020; doi: 10.1093/jac/dkaa466
- 11. McConville TH, **Annavajhala MK**, Giddins MJ, Macesic N, Herrera CM, Rozenberg FD, Bhushan GL, Ahn D, Mancia F, Trent MS, Uhlemann A-C. CrrB positively regulates high-level polymyxin-resistance and virulence in *Klebsiella pneumoniae. Cell Rep.* 2020; 3:108313. doi: 10.1016/j.celrep.2020.108313
- 12. Kundu S, Shen L, Somasundar Y, **Annavajhala MK**, Ryabov A, Collins T. TAML- and Buffer-Catalyzed Oxidation of Picric Acid by H₂O₂: Products, Kinetics, DFT and the Mechanism of the Dual Catalysis. *Inorg. Chem.* 2020; 59(18):13223-13232. doi: 10.1021/acs.inorgchem.0c01581
- 13. Papapanou PN, Park H, Cheng B, Kokaras A, Paster B, Burkett S, Watson CW-M, **Annavajhala MK**, Uhlemann A-C, Noble JM. Subgingival microbiome and clinical periodontal status in an elderly cohort: The WHICAP ancillary study of oral health. *J. Periodontol.* 2020; 91(S1):1-12. doi: 10.1002/JPER.20-0194
- 14. Annavajhala MK, Khan SD, Sullivan SB, Shah J, Pass L, Kister K, Kunen H, Chiang V, Monnot G, Ricupero CT, Mazur R, Gordon P, de Jong A, Wadhwa S, Yin MT, Demmer RT, Uhlemann A-C. Oral and gut microbial diversity and immune regulation in patients with HIV on antiretroviral therapy. mSphere 2020; 5(1):e00798-19. doi: 10.1128/mSphere.00798-19
- 15. **Annavajhala MK**, Gomez-Simmonds A, Macesic N, Sullivan SB, Kress A, Khan SD, Giddins MJ, Stump S, Kim GI, Narain R, Verna EC, Uhlemann A-C. Colonizing multidrugresistant bacteria and the longitudinal evolution of the intestinal microbiome after liver transplantation. *Nat. Commun.* 2019; 10(1):4715. doi: 10.1038/s41467-019-12633-4
- 16. LaCarpia F, Wojczyk B, **Annavajhala MK**, Rebbaa A, Culp-Hill R, D'Alessandro A, Freedberg D, Uhlemann A-C, Hod E. Transfusional iron overload and intravenous iron infusions modify the mouse gut microbiota similarly to dietary iron. *NPJ Biofilms Microbiomes* 2019; 5:26. doi: 10.1038/s41522-019-0097-2
- 17. Macesic N, Nelson B, McConville TH, Giddins MJ, Green DA, Stump S, Gomez-Simmonds A, Annavajhala MK, Uhlemann A-C. Emergence of polymyxin resistance in clinical Klebsiella pneumoniae through diverse genetic adaptations: a genomic, retrospective cohort study. Clin. Inf. Dis. 2020; 70(10):2084-2091. doi: 10.1093/cid/ciz623
- Macesic N, Khan S, Giddins MJ, Freedberg D, Whittier S, Green D, Furuya EY, Verna EC, Annavajhala MK, Gomez-Simmonds A, Uhlemann A-C. Escherichia coli harboring mcr-1 in a cluster of liver transplant recipients: detection through active surveillance and whole genome sequencing. Antimicrob. Agents Chemother. 2019; 63(6):e02680-18. doi: 10.1128/AAC.02680-18
- Gabryszewski SJ, Wong Fok Lung T, Annavajhala MK, Tomlinson KL, Riquelme SA, Khan IN, Noguera LP, Wickersham M, Zhao A, Mulenos AM, Peaper D, Koff JL, Uhlemann AC, Prince A. Metabolic adaptation supports persistent methicillin-resistant *Staphylococcus aureus* pulmonary infection. *Am. J. Respir. Cell Mol. Biol.* 2019; 61(2):185-197. doi: 10.1165/rcmb.2018-0389OC
- 20. Brotto AC*, **Annavajhala MK***, Chandran K. Metatranscriptomic investigation of adaptation in NO and N₂O production from a lab-scale nitrification process upon repeated exposure to anoxic-aerobic cycling. *Front. Microbiol.* 2018; 9:3012. doi:10.3389/fmicb.2018.03012

*contributed equally

- 21. Annavajhala MK, Kapoor V, Santo-Domingo J, Chandran K. Structural and functional interrogation of selected biological nitrogen removal systems in the United States, Denmark, and Singapore using shotgun metagenomics. *Front. Microbiol.* 2018; 9:2544. doi: 10.3389/fmicb.2018.02544
- 22. Gomez-Simmonds A, Stump S, Giddins MJ, **Annavajhala MK**, Uhlemann AC. Clonal background, resistance gene profile, and porin gene mutations modulate *in vitro* susceptibility to imipenem/relebactam in diverse Enterobacteriaceae. *Antimicrob. Agents Chemother.* 2018; 62(8):e00573-18. doi: 10.1128/AAC.00573-18
- 23. Freedberg DE, Zhou MJ, Cohen ME, Annavajhala MK, Khan S, Moscoso DI, Brooks C, Whittier S, Chong DH, Uhlemann AC, Abrams JA. Pathogen colonization of the gastrointestinal microbiome at intensive care unit admission and risk for subsequent death or infection. *Intensive Care Med.* 2018; 44:1203-1211. doi: 10.1007/s00134-018-5268-8
- 24. Gomez-Simmonds A*, **Annavajhala MK***, Wang Z*, Macesic N, Hu Yue, Giddins MJ, O'Malley A, Toussaint NC, Whittier S, Torres VJ, Uhlemann A-C. Genomic and geographic context for the evolution of high-risk carbapenem-resistant *Enterobacter cloacae* complex clones ST171 and ST78. *mBio* 2018; 9:e00542-18. doi: 10.1128/mBio.00542-18 *contributed equally
- Annavajhala MK, Kapoor V, Santo-Domingo J, Chandran K. Comammox Functionality Identified in Diverse Engineered Biological Wastewater Treatment Systems. *Environ. Sci. Technol. Lett.* 2018; 5(2):110-116. doi: 10.1021/acs.estlett.7b00577
- 26. Giddins MJ, Macesic N, **Annavajhala MK**, Stump S, Khan S, McConville TH, Gomez-Simmonds A, Uhlemann A-C. Successive emergence of ceftazidime-avibactam resistance through distinct genomic adaptations in *bla*_{KPC}-2-harboring *Klebsiella pneumoniae* ST307. *Antimicrob. Agents Chemother.* 2018; 62(3):e02101-17. doi: 10.1128/AAC.02101-17
- 27. Kundu S, **Annavajhala MK**, Kurnikov IV, Ryabov AD, Collins TJ. Experimental and Theoretical Evidence for Multiple FeIV Reactive Intermediates in TAML Activator Catalysis: Rationalizing a Counterintuitive Reactivity Order. *Chem. Eur. J.* 2012; **18**(33):10244-10249.

B. Case Reports

1. DiFranza LT, **Annavajhala MK**, Uhlemann A-C, Green D. The Brief Case: A Maggot Mystery: *Ignatzschineria* larvae sepsis secondary to an infested wound. *J Clin Microbiol* 2020; 59(3):e02279-20.

C. Pre-prints

 Park M-R, Annavajhala MK, Chandran K. Meta-azotomics of engineered wastewater treatment processes reveals differential contributions of established and novel models of N-cycling. bioRxiv 2020. doi: https://doi.org/10.1101/2020.08.25.229054

D. Reviews

 Annavajhala MK, Gomez-Simmonds A, Uhlemann A-C. Multidrug-Resistant Enterobacter cloacae Complex Emerging as a Global, Diversifying Threat. Front. Microbiol. 2019; 10:44. doi: 10.3389/fmicb.2019.00044

E. Oral Presentations

1. Annavajhala MK, Park H, Gomez-Simmonds A, Macesic N, Khan SD, Rozenberg FD,

- Giddins MJ, Stump S, Uhlemann A-C. Gut microbiome signatures associated with colonization by high-risk multidrug-resistant *Klebsiella pneumoniae* and *Escherichia coli* clones after liver transplantation. ASM Microbe 2020 *Moved online due to COVID-19
- 2. **Annavajhala MK**, Shah J, Weidler J, Kister K, Demmer RT, Wadhwa S, Yin MT, Uhlemann A-C. Periodontal disease and the oral microbiome in antiretroviral-treated patients with HIV. ACTS 2020 *Conference cancelled due to COVID-19
- Annavajhala MK, Gomez-Simmonds A, Macesic N, Sullivan SB, Khan SD, Giddins MJ, Stump S, Verna EC, Uhlemann A-C. Microbial signatures of colonization by multidrugresistant organisms (MDRO) in liver transplant recipients. ASM Microbe 2019; San Francisco, CA.
- 4. Verna E, **Annavajhala MK**, Nenad M, Brown R, Sullivan S, Korakani G, Giddins M, Khan S, Gomez-Simmonds A, Uhlemann A-C. Intestinal Microbiome Diversity is Associated with Liver Disease Etiology and Predicts Post-Liver Transplant Mortality. ATC 2018; Seattle, WA.
- 5. Zhou MJ, Cohen ME, **Annavajhala MK**, Moscosco D, Brooks C, Whittier S, Chong DH, Uhlemann A-C, Abrams JA, Freedberg DE. Gastrointestinal Bacterial Pathogen Colonization and Risk for Subsequent Infection in the Intensive Care Unit. DDW 2018; Washington, D.C.
- 6. Verna EC, Macesic N, **Annavajhala MK**, Giddins MJ, Stump S, Brown RS, Gomez-Simmonds A, Uhlemann A-C. Dynamic adaptations of intestinal microbiota after liver transplantation. AASLD 2017; Washington, D.C.
- 7. **Annavajhala MK**, Kapoor V, Santo-Domingo J, Chandran K. Comammox functionality is ubiquitous in engineered biological wastewater treatment systems. WEFTEC 2017; Chicago, IL. * Conference Featured Presentation/Speaker
- 8. **Annavajhala MK**, Fanyin-Martin A, Taher E, Elk M, Kapoor V, Santo-Domingo J, Chandran K. Metagenomics of Anaerobic Food Waste Fermentation. WEFTEC 2017; Chicago, IL.
- 9. Park MR, **Annavajhala MK**, Park H, Chandran K. Nationwide survey of microbial structure, function and metabolic pathways driven by wastewater treatment plant operating conditions and designs revealed using metagenomic and metatranscriptomic approaches. WEFTEC 2017; Chicago, IL.
- 10. Annavajhala MK, Li Z, Chandran K. Metagenomics of a Mainstream Biofilm-Based Deammonification Process with and without Bioaugmentation from a Sidestream Deammonification System. AEESP Late-Breaking Session, WEFTEC 2016; New Orleans, LA.
- 11. Brotto AC, **Annavajhala MK**, Chandran K. Effect of Long-Term Anoxic-Aerobic Cycling on Nitrous Oxide Emissions through a Combined Metagenomics and Metatranscriptomics Approach. WEF/IWA Nutrient Removal & Recovery 2016; Denver, CO.

F. Poster Presentations

- 1. Baldwin-Hunter BL, Rozenberg FD, **Annavajhala MK**, Park H, Dimango E, Keating C, Uhlemann A-C, Abrams JA. Alterations in gut microbiome in adults with cystic fibrosis associate with presence of colon adenomas and short chain fatty acid levels. AGA 2021
- 2. Gomez-Simmonds A, **Annavajhala MK**, McConville TH, Dietz DE, Shoucri SM, Laracy JC, Nelson B, Whittier S, Uhlemann A-C. Cluster of carbapenemase-producing Enterobacterales secondary infections during the COVID-19 crisis at a New York City

- hospital. IDWeek 2020
- Rubin, LG, Beachy J, Uhlemann, A-C, Annavajhala MK, Balamohan A, Jendresky L. Prolonged Outbreak of Clonal, Mupirocin-resistant Methicillin-resistant Staphylococcus aureus (MRSA) in a Neonatal Intensive Care Unit (NICU): association with Personnel and Possible Environmental Reservoir, analyzed using Whole Genome Sequencing (WGS). PAS 2020.
- 4. Ahn D, Bhushan G, McConville TH, **Annavajhala MK**, Uhlemann A-C, Prince AS. Enhanced bioenergetics by a novel acyltransferase promotes persistence of carbapenem-resistant K. pneumoniae in the airway. AJRCCM 2020.
- 5. Wong T, Shi W, Urso A, **Annavajhala MK**, Uhlemann A-C, Prince AS. Carbapenemase-producing *Klebsiella pneumoniae* ST258 activates the T6SS in response to the host metabolite itaconate to promote bacterial adaptation to the lung. AJRCCM 2020.
- 6. Yoo K, **Annavajhala MK**, Kapoor V, Santo-Domingo J, Chandran, K. Mining antibiotic resistance genes and potential human pathogens in biological nitrogen removal processes nationwide using shot-gun metagenomic analysis. WEFTEC 2019; Chicago, IL.
- Annavajhala MK, Gomez-Simmonds A, Macesic N, Sullivan SB, Kress A, Khan SD, Giddins MJ, Stump S, Verna EC, Uhlemann A-C. Drug Class-Specific Relationship between Antibiotic Exposure and the Gut Microbiota in a High-Risk Liver Transplant Cohort. ASM Microbe 2019; San Francisco, CA. * Selected for Outstanding Abstract Award and Rapid-Fire Poster Talk
- 8. Rojas R, Macesic N, Tolari G, Guzman A, **Annavajhala MK**, Uhlemann A-C. Emergence of diverse carbapenem-resistant Enterobacteriaceae (CRE) in the Dominican Republic. ID Week 2018; San Francisco, CA.
- 9. **Annavajhala MK**, Saidu R, Tergas A, Kuhn L, Denny L, Uhlemann AC. Distinct cervical microbial diversity and community structure by human papilloma-virus and cervical disease status in HIV-positive women in South Africa. Microbiome in HIV 2018; Rockville, MD.
- 10. Annavajhala MK, Geng W, Hill-Ricciuti A, Ferguson S, Stump S, Giddins MJ, Messina M, Zachariah P, Green D, Whittier S, Saiman L, Uhlemann AC. Hybrid Sequencing and Assembly Reveals Genomic Diversity of Methicillin-susceptible Staphylococcus aureus (MSSA) from a Neonatal Intensive Care Unit (NICU) Surveillance Effort. ASM Rapid Applied Microbial Next Generation Sequencing and Bioinformatic Pipelines 2018; Tysons, VA.
- 11. May M, **Annavajhala MK**, Compres G, Freedberg DE, Graham R, Uhlemann AC, Abrams JA. A Randomized Controlled Trial to Assess the Effects of an Antimicrobial Mouthwash on the Oral and Esophageal Microbiome. DDW 2018; Washington, DC.

G. Thesis

 Annavajhala MK. Meta-omics-derived structure, function, and activity of mixed microbial communities driving biological nutrient removal and recovery. Columbia University, New York, NY, 2017.

Invited Presentations

A. Seminars

1. "Activism in the STEM Workplace." BioMediGirls Annual Advocacy Summit 2020; online.

- 2. "The Human Microbiome: Health, Illness, and Prevention." Bloomingdale Aging in Place, 2019; New York, NY.
- 3. "Genomics of methicillin-susceptible *Staphylococcus aureus* (MSSA) from a neonatal intensive care unit (NICU)." Pediatric Infectious Diseases Division Conference Rounds, Columbia University Medical Center, 2019; New York, NY.
- 4. "Microbiome 101." Microbiome Working Group Seminar Series, Columbia University Medical Center, 2017; New York, NY.
- 5. "Using Molecular Techniques and Next-Generation Sequencing to Understand and Optimize Wastewater Treatment Processes." Hot Topics in Water and Wastewater, NJAES Office of Continuing Professional Education, Rutgers University, 2015; Belvedere, NJ.
- 6. "Meta-Omics of the Engineered Water Cycle." Ion Torrent World Tour, 2015; New York, NY.

Trainings and Workshops

Exposome Bootcamp, Columbia University Mailman School of Public Health, July 2020

Machine Learning Bootcamp, Columbia University Mailman School of Public Health, June 2019

Introduction to Python, Foundations for Research Computing, Columbia University, August 2019