# Emily R. Davenport

Assistant Professor – Pennsylvania State University

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## Education

2009 - 2014	Ph.D in Human Genetics  Certificate in University Teaching University of Chicago; Chicago, IL  Advisor: Yoav Gilad (supported by NIH training grant	
2003 - 2007	B.S. in Bacteriology (with comprehensive honors) University of Wisconsin – Madison; Madison, WI	
2005	International Study Program, selected participant National University of Ireland, Galway; Galway, Ireland	

# Academic, Research, and Industry Experience

2019 -	<b>Assistant Professor</b> . Department of Biology, Huck Institutes of the Life Sciences, Institute for Computational and Data Sciences; The Pennsylvania State University; University Park, PA.
2018 - 2019	<b>Visiting Post-doctoral Fellow</b> . Supervisor: Ruth Ley (supported by NIH NRSA) Department of Microbiome Science, Max Planck Institute for Developmental Biology; Tübingen, Germany.
2014 - 2019	<b>Post-doctoral Fellow</b> . Advisor: Andrew Clark (supported by NIH NRSA)  Department of Molecular Biology and Genetics, Cornell University; Ithaca, NY.
2008 - 2009	Sequence Capture Technician. Supervisor: Mindy Bennett. Roche NimbleGen; Madison, WI.
2007 - 2008	Forensic Scientist – DNA Analyst. Supervisor: Sherry Culhane.  DNA Unit, Department of Justice - Wisconsin State Crime Lab; Madison, WI.
2004 - 2007	<b>Undergraduate Research Assistant/Independent Research</b> . Advisor: Steven Barclay Department of Bacteriology, University of Wisconsin – Madison; Madison, WI.
2006	International Research Experience for Students (IRES) in Microbiology Summer Research Program Participant. Advisor: Sukathida Ubol Department of Microbiology, Mahidol University; Bangkok, Thailand.

## Publications - peer reviewed

\* denotes equal contribution
Underlined denotes mentee author

### Submitted

- 21. Jeganathan NA, **Davenport ER**, Yochum GS, Koltun WA. *The microbiome in diverticulitis*.
- 20. Amato KR, Arrieta M, Azad M, Bailey MT, Broussard JL, Bruggeling C, Claud EC, Costello EK, **Davenport ER**, Dutilh BE, Swain Ewald HA, Ewald P, Hanlon EC, Julion WA, Keshavarzian A, Maurice CF, Miller GE, Preidis GA, Ségurel L, Singer BH, Subramanian S, Zhao L, Kuzawa CW. *The Human Microbiome and Health Inequities*.
- 19. Nowak C, Eriksson N, van Setten J, Henry A, Stenemo M, Johansson Å, **Davenport ER**, Lagerqvist B, Becker R, Budaj A, Hagström E, Held C, Himmelmann A, Katus HA, Koenig W, Siegbahn A, Steg P, James S, Axelsson T, Syvänen A, Stewart R, Akerblom A, Storey R, Waterworth D, White H, Allayee H, Hartiala J, Hazen S, Wilson Tang WH, Asselbergs F, Howe L, McCubrey R, Tragante V, Quyyumi A, Alkhodar A, Liu C, San Y, Alver M, Metspalu A, Levin D, Mordi I, Lang C, Palmer C, Glinge C, Engstrøm T, Jabbari R, Tfelt-Hansen J, Delgado G, Kleber M, März W, Ford I, Wouter Jukema J, Scott D, Trompet S, Gijsberts C, Pasterkamp G, van der Laan S, Ingelsson E, Burgess S, Lumbers R, Patel R, Arnlov J, Wallentin L, and Fall T. Causal effect of GDF-15 on cardiovascular disease and mortality genome-wide association study and Mendelian randomization analysis.

#### 2020

- 18. Marderstein AR, **Davenport ER**, Kulm S, Van Hout C, Elemento O, Clark AG. Leveraging phenotypic variability to identify genetic interactions in human phenotypes. American Journal of Human Genetics (accepted)
- 17. <u>Nichols RG</u> and **Davenport ER.** The relationship between the gut microbiome and host gene expression: A review. Human Genetics. DOI: 10.1007/s00439-020-02237-0

#### 2019

- 16. <u>Gong X</u>, **Davenport ER**, Wang D, Clark AG. Lack of spatial and temporal genetic structure of Anguilla japonica populations. Conservation Genetics. 2019;20:467
- 15. Kachoo P\*, Eraso JM\*, Beres SB, Olsen RJ, Zhu L, Nasser W, Bernard PE, Cantu CC., Ojeda Saavedtra M, José Arredondo M, Strope B, Do H, Kumaraswami M, Vuopio J, Gröndahl-Yli-Hannuksela K, Kristinsson KG, Gottfredsson M, Pesonen M, Pensar J, **Davenport ER**, Clark AG, Corander J, Caugant DA, Gaini S, Magnusses MD, Porter AR, DeLeoFR, and Musser JM.

Integrated analysis of population genomics, transcriptomics and virulence provides novel insights into Streptococcus pyogenes pathogenesis. Nature Genetics. 2019;51 548-559

#### 2018

14. Jha AR, **Davenport ER**, Gautam Y, Bhandari D, Tandukar S, Ng K, Holmes S, Prasad Gautam G, Bahadur Sherchand J, Bustamante CD, and Sonnenburg JL. *Gut microbiome transition across a life gradient in Himalaya*. PLoS Biology. 2018;16(11):e2005396 (epub 2018 Nov 15)

#### 2017

- 13. **Davenport ER\***, Sanders JG\*, Song SJ, Amato KR, Clark AG, and Knight R. *The human microbiome in evolution*. BMC Biology. 2017. 15:127
- 12. Goodrich JK, **Davenport ER**, Clark AG, and Ley RE. The relationship between the human genome and microbiome comes into view. Annual Reviews Genetics. 2017. 51(1)
- 11. Igartua C, **Davenport ER**, Gilad Y, Nicolae DL, Pinto J, and Ober C. Host genetic variation in mucosal immunity pathways influences the upper airway microbiome. Microbiome. 2017 Feb 1;5:16

#### 2016

- Davenport ER, Goodrich JK, Bell JT, Spector TD, Ley RE, Clark AG. ABO antigen and secretor statuses are not associated with gut microbiota composition in 1,500 twins. BMC Genomics. 2016 Nov 21;17:941
- 9. Beaumont M, Goodrich JK, Jackson MA, Yet I, **Davenport ER**, Vieira-Silva S, Debelius J, Pallister T Mangino M, Raes J, Knight R, Clark AG, Ley RE, Spector TD, and Bell JT. *Heritable components of the human fecal microbiome are associated with visceral fat.* Genome Biology. 2016 Sep 26;17:189
- 8. Goodrich JK, **Davenport ER**, Beaumont M, Jackson MA, Knight R, Spector TD, Bell JT, Clark AG, and Ley RE. *Genetic determinants of the gut microbiome in UK twins*. Cell Host and Microbe. 2016: 19(5), 731-743
- 7. Goodrich JK\*, **Davenport ER**\*, Waters JL\*, Clark AG, and Ley RE. *Cross-species comparisons of host genetic associations with the microbiome*. Science. 2016: 352(6285), 532-535
- 6. **Davenport ER**. Elucidating the role of the host genome in shaping microbiome composition. Gut Microbes. 2016: 7(2), 178-184
- 5. Blischak JD, **Davenport ER**, and Wilson G. A quick introduction to version control with Git and GitHub. PLoS Computational Biology. 2016;12(1):e1004668 (epub 2016 Jan 19)

#### 2015

- 4. **Davenport ER,** Cusanovich DA, Michelini K, Barrerio LB, Ober C, and Gilad Y. *Genome-wide association studies of the human gut microbiota*. PLoS One. 2015;10(11):e0140301 (epub 2015 Nov 3)
  - ----> An Editor's Pick for the PLoS Microbiology special collection:

http://collections.plos.org/microbiology-picks

----> An Editor's Pick for the PLoS Experimental Biology special collection:

http://collections.plos.org/experimental-biology

#### 2014

- 3. Zhou X, Cain CE, Myrthil M, Lewellen N, Michelini K, **Davenport ER**, Stephens M, Pritchard JK, and Gilad Y. *Epigenetic modifications are associated with inter-species gene expression variation in primates*. Genome Biology. 2014 Dec 3;15(12):547
- 2. **Davenport ER**, Mizrahi-Man O, Michelini K, Barreiro LB, Ober C, and Gilad Y. Seasonal variation in human gut microbiome composition. PLoS One. 2014;9(3):e90731 (epub 2014 Mar 11)

#### 2013

1. Mizrahi-Man O, **Davenport ER**, and Gilad Y. *Taxonomic classification of bacterial 16S rRNA genes using short sequencing reads: Evaluation of effective study designs.* PLoS One. 2013;8(1):e53608 (epub 2013 Jan 7)

## Publications – non-peer reviewed

- 2. **Davenport ER**. Genetic variation shapes murine gut microbiota via immunity. Trends in Immunology. 2020;41(1)
- 1. **Davenport ER**. Tooth be told, genetics influence oral microbiome. Cell Host & Microbe. 2017;22(3)

## **Presentations**

**Invited Plenary Platform Presentations** 

Davenport, ER. Using human genetics to understand the physiological basis of host

#### **Invited Platform Presentations**

2020

2020	microbiome interactions. The Vanderbilt Institute for Infection, Immunology, and Inflammation (VI4) Annual Symposium [*virtual due to COVID-19]
2020	<b>Davenport, ER</b> . Using human genetics to understand the physiological basis of host microbiome interactions. Workshop: Cultivating communities: Making sense of host-microbiome interactions through the lens of genetics. The Allied Genetics Conference (TAGC). Washington DC [*virtual due to COVID-19]
2019	<b>Davenport, ER</b> . Causes and consequences of human gut microbiome variation. One Health Workshop. Pennsylvania State University. State College, PA
2019	<b>Davenport, ER</b> , Spector TD, Ley RE, and Clark AG. Simultaneously modeling host genetics and microbiome impact on immune-related traits. Pennsylvania State University Bioinformatics and Genomics Annual Retreat. State College, PA
2018	<b>Davenport, ER</b> , Spector TD, Ley RE, and Clark AG. Simultaneously modeling host genetics and microbiome composition reveals the heritability and proportion of variance explained due to the microbiome for immune-related traits. Probabilistic Modeling in Genomics (ProbGen). Cold Spring Harbor Labs, NY  * Co-chair of "Cancer, the microbiome, and beyond" session

- Davenport ER. Role of host genetics in shaping the gut microbiota. The Human Capital and Economic Opportunity (HCEO) Working Group Conference on The Gut Microbiome in Human Biology and Health: New Opportunities for the Study of Health Disparities. Chicago, IL
- Davenport ER. The role of host genetics in determining human gut microbiome composition. The American Association of Physical Anthropologists Annual Meeting. Wiley Invited Podium Symposium Humans as Holobionts: The Microbiome as a Biological System in Human Evolution. New Orleans, LA.
- Davenport ER. The role of host genetics in determining human gut microbiome composition. The 2016 Nordic-North American Symposium on Antimicrobial Resistance and Molecular Population Genomics in Houston, TX.

#### Platform Presentations

- Davenport ER, Spector TD, Ley RE, and Clark AG. Co-occurrence network modeling reveals disease-specific configurations of microbiome community structure across 2,500 twins. American Society of Human Genetics Annual Meeting (ASHG) in Orlando, FL.
- Davenport ER, Spector TD, Ley RE, and Clark AG. Modeling human gut microbiome community structure across healthy and diseased states in 2,500 twins. Society of Molecular Biology and Evolution Annual Meeting (SMBE) in Austin, TX.
- 2017 **Davenport ER**, Spector TD, Ley RE, and Clark AG. Modeling human gut microbiome community structure across healthy and diseased states in 2,500 twins. Biology of Genomes (BoG) in Cold Spring Harbor, NY.
- Davenport ER, Mizrahi-Man O, Barreiro LB, Ober C, and Gilad Y. Examining the roles of diet, age, and sex on the composition of the human fecal microbiome. University of Chicago Molecular Biosciences Cluster Retreat in Galena, IL.

### Poster presentations

- Davenport ER, Goodrich JK, Bell JT, Spector TD, Ley RE, and Clark AG. ABO antigen and secretor status are not associated with gut microbiota composition. American Society of Human Genetics Annual Meeting (ASHG) in Baltimore, MD.
- Davenport ER, Mizrahi-Man O, Michelini K, Barreiro LB, Ober C, and Gilad Y. poopQTLs: Genome-wide associations of the human gut microbiota. Society for Molecular Biology and Evolution Annual Meeting (SMBE) in San Juan, PR.
- Davenport ER, Mizrahi-Man O, Michelini K, Barreiro LB, Ober C, and Gilad Y. *Temporal variation in human gut microbiome composition in the Hutterites*. American Society of Human Genetics Annual Meeting (ASHG) in Boston, MA.
- Davenport ER, Mizrahi-Man O, Michelini K, Barreiro LB, Ober C, and Gilad Y. Examining the temporal stability of the fecal microbiome in an isolated, founder population. Cell Symposium: the Microbiome and Host Health in Lisbon, Portugal.
- Davenport ER, Mizrahi-Man O, Barreiro LB, Ober C, and Gilad Y. Examining the roles of diet, age, and sex on the composition of the human fecal microbiome. American Society of Human Genetics Annual Meeting (ASHG) in San Francisco, CA.
- Davenport ER, Mizrahi-Man O, Barreiro LB, Ober C, and Gilad Y. Examining the genetic basis of interindividual variation in the human fecal microbiome. International Human Microbiome Conference in Paris, France.
- Yao T, **Davenport ER**, Poroyko V, Liu D, Lemanske R, Gern J, Ober C, Jackson D, Gilad Y, Pinto J. *The nasal microbiome and development of asthma in a birth cohort*. Biology of Genomes (BoG) in Cold Spring Harbor, NY.

## **Invited Seminars**

2021	UGA Interdepartmental Health Disparities Working Group, University of Georgia Department of Biology, Trinity University Department of Human Genetics, University of Chicago Department of Cellular and Molecular Medicine, University of Arizona Gut Interest Group, Penn State University College of Medicine
2020	Department of Public Health Sciences, Penn State College of Medicine Department of Biology, Vanderbilt University [cancelled because of COVID-19] Department of Human Genetics, UCLA [cancelled because of COVID-19] Department of Human Genetics, University of Chicago [cancelled because of COVID-19]
2019	Penn State Microbiome Center Exchange, Penn State College of Medicine Department of Biology, Duke University Department of Nutrition, Pennsylvania State University Department of Biology, University of New Mexico Department of Biology, Marquette University Department of Biology, Auburn University Department of Biology, Pennsylvania State University
2018	Max Planck Institute for Developmental Biology Department of Biology, Bucknell University Department of Biology, Barnard College Department of Biology, Smith College
2017	Department of Biology, Wake Forest University
2016	The Huck Institutes of the Life Sciences, Pennsylvania State University
2013	Department of Biology, Chicago State University
2013	Emory University and Yerkes National Primate Research Center

# Teaching Experience

University Courses – Instructor of Record

- 2020 Penn State University, Department of Biology, *BIOL439: Practical Bioinformatics* (~25 students)
- 2020 Penn State University, Department of Biology, *PSU16: First Year Seminar* (~30 students)

## **Teaching Assistantships**

2014	University of Chicago, Biological Sciences Division: HGEN 47300: Genomics and Systems Biology (Taught 6 lectures)
2011	University of Chicago, Biological Sciences Division: HGEN 47000: Human Genetics I
2011	University of Chicago, Biological Sciences Division: HGEN 47300: Genomics and Systems Biology
2010	University of Chicago, Biological Sciences Division: MGCB 31400: Genetic Analysis of Model Organisms

### **Guest Lectures**

2020	Penn State University, Anthropology ANTH497: Microbial Insights into Anthropology – "Microbiome studies in the Hutterites"
2016	Cornell University, Molecular Biology and Genetics: BIOMG 4870: Human Genomics – "Cystic Fibrosis and PKU"
2016	Cornell University, Biological Sciences: BIOMI 3210: Human Microbes and Health – "Microbiome studies in the Hutterites"
2016	Pennsylvania State University, Biochemistry and Molecular Biology: BMB 484: Functional Genomics – "Introduction to Population Genetics"
2015	Cornell University, Molecular Biology and Genetics: BIOMG 4870: Human Genomics – "Linkage disequilibrium mapping, or Genome-wide Association Studies (GWAS)"
2011	University of Chicago, Biological Sciences Division: <i>HGEN 47000: Human Genetics I – "Human genome structure and variation"</i>

## Workshops

2021	<b>Instructor</b> – "Git and GitHub", Penn State, Huck Institutes of the Life Sciences, Microbiome Center
2020	<b>Instructor</b> – "Microbiome Center Kick-Start Workshop", Penn State, Huck Institutes of the Life Sciences, Microbiome Center
2019	$\label{local-control} \textbf{Instructor} - "Introduction to Statistics with R", Max Planck Institute for Developmental Biology; Tübingen, Germany$
2016	Instructor – "Learn about Git and Github", Cornell University, CPGSA
2016	Instructor – "Introduction to R", University of Chicago, Biological Sciences Division

## Software Carpentry Workshops [content I taught]

2017	Lead Instructor, TGen, Phoenix, AZ (June) [R and version control with Git]
2016	<b>Instructor</b> , University of Chicago, Biological Sciences Division (September) [review of shell and R, writing reproducible reports, and version control with Git]
2016	<b>Lead Instructor</b> , Cornell University, Department of Molecular Biology and Genetics (August) [version control with Git]
2015	<b>Instructor</b> , University of Chicago, Biological Sciences Division (September) [review of shell and R, writing reproducible reports, and version control with Git]
2015	Instructor, Pennsylvania State University (June) [shell and version control with Git]
2014	<b>Instructor</b> , University of Chicago, Biological Sciences Division (September) [version control with Git]
2014	Lead Instructor, University of Toronto (July) [version control with Git]
2013	Instructor, University of Chicago, Biological Sciences Division (September) [shell]
2013	Instructor, University of Chicago. (June) [shell]

## Data Carpentry Workshops [content I taught]

2016	Instructor, Cornell University (June) [reproducible reports with Rmarkdown and R
	programming]
2015	<b>Instructor</b> , Cornell University (January) [automating repetitive tasks with command line
	shell

# Mentorship and Advising

## Student and Postdoc Advising (Primary Advisor)

Years	Name	Туре	Institution	Program
2020 –	Nur Shahir	Postdoc	Penn State	Biology
2020 –	Erica Ryu	Ph.D.	Penn State	Biology
2020 –	Robert Nichols	Postdoc	Penn State	Biology
2018	Adon Chowdhury	Undergrad	Cornell	Biometry and Statistics
2017 – 2018	Trang Dau	Undergrad	Cornell	Human Bio, Health, & Society
2016 – 2017	Xiaoling Gong	Scientist	Cornell	Molecular Bio and Genetics
2015 – 2017	Monica Guardado <sup>1</sup>	Undergrad	Penn State	Biology

#### Notes:

### Dissertation Committee Member

<sup>&</sup>lt;sup>1</sup>Awarded an American Society of Microbiology (ASM) Research Capstone Fellowship in 2017

Defended	Deg.	Name	University	Department/Program
In progress	Ph.D.	Miranda DePriest	Penn State	Plant Path. and Env. Micro.
In progress	M.S.	Nellie Heitzman	Bucknell University	Biology
In progress	Ph.D.	Shreya Ramachandran	University of Chicago	Human Genetics
In progress	Ph.D.	Scott Eckert	Penn State	Bioinfo. and Genomics
In progress	Ph.D.	Jeremy Sutherland	Penn State	Bioinfo. and Genomics
In progress	Ph.D.	Julia Stewart	Penn State	Biology
In progress	Ph.D.	Sterling Wright	Penn State	Anthropology
In progress	Ph.D.	Lixiang Zhang	Penn State	Statistics
In progress	MPS	Veronica Cappas	Penn State	Forensic Science

## Qualifying Exam Committee Member

Year	Program	Students
2020	Bioinformatics and Genomics	Jeremy Sutherland, Shaopeng Liu

### **Rotation Advisor**

Year	Program	Name(s)
2020	Bioinfo. & Genomics	Kyle McGovern
2019	Biology	Erica Ryu

# Academic Honors and Funding

2017	Genetics Society of America (GSA) DeLill Nasser Travel Award for Professional Development in Genetics (\$1000)
2016 - 2019	NIH Ruth L. Kirschstein National Research Service Award (NRSA) – F32DK109595 (\$173,079)
2014	University of Chicago Biological Sciences Division Travel Award (\$500)
2011	University of Chicago Digestive Diseases Research Core Center (DDRCC) Pilot and Feasibility award (\$20,000, written by E.R. Davenport to support dissertation research, submitted by Y. Gilad)
2010 - 2012	NIH Genetics and Regulation Training Grant (University of Chicago – 2 years of stipend support and tuition)
2007	Graduated with comprehensive honors: honors in Bacteriology and the liberal arts (University of Wisconsin – Madison)
2004	Dean's List (University of Wisconsin – Madison)
2003	William F. Vilas Scholarship (University of Wisconsin – Madison)

# Professional Development

2021	Safer People Safer Places Transgender and Gender Inclusion 101 – Penn State University
2020	Becoming an Active Witness who Hears, Sees, Feels, and Does active bystander training – Penn State University
2019	Mentoring Matters Workshop – Eberly College of Science, Penn State University
2017 – 2018	Cornell Center for Teaching Excellence GET SET teaching and learning workshops completed:
	- An Integrated Course Design Approach to Planning Your Class
	- Tips on Writing a Strong Teaching Philosophy Statement
	- Integrating Technology into Your Classroom
	- Developing Service-Learning in the Disciplines
	- Flipping the Classroom as a TA
	- Using Your Research Experience to Improve Your Teaching
	- Holding Effective Office Hours
	- Teaching and Mentoring Across Differences
	- Leading a Discussion in an Online Classroom
2017	<b>Assessing Learning and Teaching certificate</b> – Cornell University Center for Teaching Innovation. Workshops included:
	- Designing Learning Outcomes for Your Course
	- Utilizing Classroom Assessment Techniques to Evaluate Student Learning
	- Developing Rubrics for Effective Grading
2017	<b>Creating an Engaging Classroom certificate</b> – Cornell University Center for Teaching Innovation
	- Building a Collaborative Learning Environment
	- Using Theater Techniques to Enhance Your Teaching
	- Engaging Students in Quantitative Courses
2016	<b>The Practice of Inclusive Teaching in STEM certificate</b> – Cornell University Center for Teaching Excellence
2016	Building Mentoring Skills for an Academic Career certificate program – Cornell University Center for the Integration of Teaching and Learning (CU-CIRTL)
2015 - 2016	Postdoc Leadership Certificate Program – Cornell University
2014	<b>Certificate in University Teaching</b> – University of Chicago Center for Teaching and Learning
2013	Software Carpentry Instructor training – Software Carpentry

# Professional Affiliations

2019 -	Society for Advancement of Chicanos/Hispanics and Native Americans in Science (SACNAS)
2017 -	Genetics Society of America (GSA)
2014 -	Society for Molecular Biology and Evolution (SMBE)
2012 -	American Society of Human Genetics (ASHG)
2017	American Society for Microbiology (ASM)
2017	American Association of Physical Anthropologists (AAPA)
2016 – 2019	National Postdoc Association (NPA)
2011 – 2018	American Association for the Advancement of Science (AAAS)

## Outreach

Interview	(2020) Eberly College of Science "Hey! I Got a Question About That" podcast on the
	microbiome: https://www.youtube.com/watch?v=960zTLUD7VU&t=881s
Advocate	(2019) Rally for Medical Research Hill Day advocate, representing American Society of
	Human Genetics
Reviewer	(2015 - ) American Society of Human Genetics (ASHG) DNA Day Essay Judge
Member	(2015 - 2018) Genetics Education and Outreach Network (GEON)

# Leadership and Service

## Departmental

2020 -	Biology Seminar Series Organizing Committee, Department of Biology, PSU
2020 -	Biology Graduate Admissions Committee, Department of Biology, PSU
2010 - 2013	Molecular Biosciences organizational committee: student representative from the Department of Human Genetics on orientation week, annual molecular biosciences retreat, and recruitment organizing committees, Department of Human Genetics, University of Chicago

### Institutional

2021	Invited Panelist - "First Yea	rs as Faculty in STEM" GRAD	UCon, University of Chicago
2020	<b>Grant reviewer</b> - Penn State Pilot Award Program	•	nical and Translational Science
2018	Cornell University. So	cionships Policy Committee – lee http://theuniversityfaculty.committee/ for details.	·
2017 – 2018	•	ostdoc Fellow Invited Lecture . Department of Molecular Bio	
2016 – 2018	Member - Cornell Universit	y Postdoctoral Advisory Cour	ncil
2016 – 2017	Reviewer - Sigma Xi Grants-In-Aid of Research, Cornell University		
National			
2015 –	Reviewer -	Environmental	mSystems
	American Journal of	Microbiology	Nature Communications
	Primatology	Genes	Nature Microbiology
	Applied and	Genome Biology	npj Biofilms and
	Environmental Microbiology (AEM)	Genomics, Proteomics,	Microbiomes
	BMC Genomics	and Bioinformatics	PLoS Genetics
	Cell	Gut Microbes	PLoS ONE
		Journal of Allergy and	PNAS
	Cell Host & Microbe	Clinical Immunology (JACI)	Science
	Cell Reports	Journal of Evolutionary	Scientific Reports
	Communications Biology	Biology	Thorax
	Critical Reviews in Microbiology	Microbial Ecology	Trends in Genetics
	Diabetologia	Microbiome	Trends in Immunology
	eLife	Microorganisms	
2020 –	<b>Co-chair</b> - American Society Committee (CDC)	of Human Genetics (ASHG) (	Career Development
2019 –	Abstract Reviewer - SACN	AS research presentation abst	racts (Life Sciences)
2019 –	Abstract Reviewer - SACN	AS Travel Scholarships (Life Sc	ciences)
2019 –	<b>Board Member</b> - Cards Aga	ainst Humanity Science Amba	ssador Scholarship Program
2018 –		nerican Societies for Experime pportunities Subcommittee	ental Biology (FASEB)
2020 – 2021	Grant Reviewer - National S	•	
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2018, 2020 Grant Reviewer - Wellcome Trust Investigator Award in Science

2020	Moderator - The Allied Genetics Conference (TAGC) platform session: Disease Models and Aging (Mammal) *TAGC 2020 was held virtually due to COVID-19
2019	Incoming Chair - American Society of Human Genetics (ASHG) Training and Development Committee
2019	<b>Moderator</b> - American Society of Human Genetics (ASHG) Annual Meeting platform session: Gene Expression Variation Across Diverse Global Populations
2019	Abstract Reviewer - ASHG Annual Meeting abstracts (Bioinformatics and Computational Approaches)
2017 – 2019	<b>Member</b> - American Society of Human Genetics (ASHG) Training and Development (TDC) Committee
2017	<b>Moderator</b> - Academic Career Panel at the American Society of Human Genetics Annual Meeting (ASHG)
2011, 2012, 2	O14 Judge - Annual Chicago Public Schools Student Science fair
2011 - 2014	Judge - Annual Chicago Area Undergraduate Research Symposium (CAURS)