Emily R. Davenport

Curriculum Vitae – Updated July 2018 Postdoc - Cornell University Visiting Postdoc – Max Planck Institute for Developmental Biology

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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

2018 -	Visiting Post-doctoral Fellow. Supervisor: Ruth Ley Department of Microbiome Science, Max Planck for Developmental Biology; Tübingen, Germany.
2014 -	Post-doctoral Fellow . 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	Undergraduate Research Assistant/Independent Research . 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

- 15. 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* (submitted)
- 14. <u>Gong X</u>, **Davenport ER**, Wang D, Clark AG. Lack of spatial and temporal genetic structure of Anguilla japonica populations. (submitted)

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

- 10. **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

1. **Davenport ER**. Tooth be told, genetics influence oral microbiome. Cell Host & Microbe. 2017;22(3)

Presentations

Invited Platform Presentations

- 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.
- 2016 **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.
- 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. Society of Molecular Biology and Evolution Annual Meeting (SMBE) in Austin, TX.
- 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.
- 2012 **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

2015 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. 2014 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. 2013 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. 2013 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. 2012 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. 2012 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. 2011 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** 2016 **Davenport ER**. The role of host genetics in determining gut microbiome composition. The Huck Institutes of the Life Sciences, Pennsylvania State University 2013 Davenport ER. Seasonal variation in human gut microbiome composition. Chicago State University 2013 Davenport ER. Seasonal variation in human gut microbiome composition. Emory: Yerkes National Primate Research Center

Teaching Experience

Teaching Assistantships

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

Guest Lectures

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

2016	Instructor – "Learn about Git and Github", Cornell University, CPGSA
2016	Instructor – "Introduction to R", University of Chicago, Biological Sciences Division (September)

Software Carpentry Workshops [content I taught]

2017	Lead Instructor, TGen, Phoenix, AZ (June) [R and version control with Git]
2016	Instructor , University of Chicago, Biological Sciences Division (September) [review of shell and R, writing reproducible reports, and version control with Git]
2016	Lead Instructor , Cornell University, Department of Molecular Biology and Genetics (August) [version control with Git]

2015	Instructor , 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	Instructor , 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	Instructor, Cornell University (January) [automating repetitive tasks with command line
	shell

Mentorship

Adon Chowdhury - Undergraduate @ Cornell University, Lab of Andrew Clark (2018 -)

Adon is a Biometry and Statistics major who became interested in combining his interests in biology and data science by studying the microbiome. He is currently leading a project examining whether the dynamics of microbial community assembly in the gut are influenced by host genetics, using samples from the large TwinsUK cohort. Adon's background has proven perfect for this project, as it requires strong statistical and computational skills, along with broad biological understanding.

Trang Dau – Undergraduate @ Cornell University, Lab of Andrew Clark (2017 -)

Trang is a Human Biology, Health, and Society major who became interested in the two-way relationship between the microbiome and the human host after taking courses in microbiology and ecology at Cornell. She has contributed to a number of projects during her time in lab, including examining the role of host genetics in microbial community assembly and the role of microbial networks in health and disease.

Xiaoling Gong – Visiting scientist @ Cornell University, Lab of Andrew Clark (2016 - 2017)

Xiaoling is a mid-career investigator on a two-year fellowship from the Chinese Academy of Science visiting the Clark lab to expand the analysis of population structure of Japanese Eels, an important aquaculture species in Asia. During her time as a visiting scientist in the Clark Lab, I mentored her on the use of bioinformatics tools for analyzing RADseq data and the application of population genetic statistics to answer the open question of whether Japanese Eels are panmictic.

----> The manuscript describing this work, "Lack of spatial and temporal genetic structure of Anguilla japonica populations" is under review

Monica Guardado – Undergraduate @ Penn State University, Lab of George Perry (2015 - 2017)

Monica became interested in host-microbiome dynamics during coursework for her Biology major. During her time in the Perry Lab, she's been examining whether termite-eating behavior in chimpanzees results in the transfer of termite microbiota into the chimpanzee gut. I've mentored her on both the wet lab and computational aspects of analyzing 16S rRNA sequencing data.

> ----> Awarded an American Society of Microbiology (ASM) Research Capstone Fellowship (2017)

----> Now a Clinical Lab Associate at uBiome (a microbiome start-up)

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 (\$168,414)
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

- 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

- 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 Assessing Learning and Teaching certificate - 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 Creating an Engaging Classroom certificate - Cornell University Center for Teaching Innovation - Building a Collaborative Learning Environment - Using Theater Techniques to Enhance Your Teaching - Engaging Students in Quantitative Courses The Practice of Inclusive Teaching in STEM certificate – Cornell University Center for 2016 Teaching Excellence Building Mentoring Skills for an Academic Career certificate program - Cornell 2016 University Center for the Integration of Teaching and Learning (CU-CIRTL) Postdoc Leadership Certificate Program – Cornell University 2015 - 2016 2014 Certificate in University Teaching - University of Chicago Center for Teaching and Learning 2013 **Software Carpentry Instructor training** – Software Carpentry

- Developing Service-Learning in the Disciplines

Professional Affiliations

2017 -	Genetics Society of America (GSA)
2016 -	National Postdoc Association (NPA)
2014 -	Society for Molecular Biology and Evolution (SMBE)
2012 -	American Society of Human Genetics (ASHG)
2011 -	American Association for the Advancement of Science (AAAS)
2017	American Society for Microbiology (ASM)
2017	American Association of Physical Anthropologists (AAPA)

Leadership and Service

Reviewer Applied and Genes **PLoS Genetics**

> **Environmental** Gut Microbes PLoS ONE

Microbiology (AEM) Genomics, Proteomics, **PNAS**

BMC Genomics and Bioinformatics Scientific Reports Cell

Journal of Allergy and Trends in Genetics Cell Host & Microbe

Clinical Immunology

Trends in Immunology (JACI) Diabetologia

Environmental Microbiology Nature Communications

Reviewer Wellcome Trust Investigator Award in Science

Member (2017 -) American Society of Human Genetics (ASHG) Training and Development

Microbiome

Committee

Member (2015 -) Genetics Education and Outreach Network (GEON)

Representat- (2018) Consensual Relationships Policy Committee – Postdoc representative, Cornell ive University. See http://theuniversityfaculty.cornell.edu/news/consensual-relationships-

policy-committee/ for details.

Cofounder & (2017 - 2018) Postdoc Fellow Invited Lecture in Research and Career Development

Organizer series. Department of Molecular Biology, Cornell University (2016 - 2018) Cornell University Postdoctoral Advisory Council Member

Organizer & (2017) Academic Career Panel at the American Society of Human Genetics Annual

Moderator Meeting (ASHG)

Reviewer (2016 - 2017) Sigma Xi Grants-In-Aid of Research, Cornell University

(2011, 2012, 2014) Annual Chicago Public Schools Student Science fair (for district fair Judge

winners)

Judge (2011 - 2014) Annual Chicago Area Undergraduate Research Symposium (CAURS)

Member (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