Lawrence A. David

CONTACT INFORMATION

Durham, NC 27708 Web: http://www.ladlab.org

EDUCATION

Massachusetts Institute of Technology 2005-2010

Ph.D., Computational & Systems Biology

Columbia University 2001-2005

B.S., Biomedical Engineering, summa cum laude

PROFESSIONAL EXPERIENCE

Assistant Professor, Duke University	2013-
Junior Fellow, Harvard Society of Fellows	2010-2013
Ph.D. student, Massachusetts Institute of Technology - PI: Eric Alm	2005-2010
Undergraduate research assistant, Columbia University - PI: Chris Wiggins	2002-2005

PUBLICATIONS

[* DENOTES EQUAL CONTRIBUTION]

- 11. **LA David**, AC Materna, J Friedman, I Baptista, MC Blackburn, A Perrotta, SE Erdman, EJ Alm. Host lifestyle affects human microbiota on daily timescales. *Genome Biol*, **15**:R89, Jul 2014.
- 10. LA David, CF Maurice, RN Carmody, DB Gootenberg, JE Button, BE Wolfe, AV Ling, AS Devlin, Y Varma, MA Fischbach, SB Biddinger, RJ Dutton, PJ Turnbaugh. Diet rapidly and reproducibly alters the gut microbiome. <u>Nature</u>, 505(7484):559-563, Jan 2014.
- 9. G Szabo, SP Preheim, KM Kauffman, **LA David**, J Shapiro, EJ Alm, MF Polz. Reproducibility of *Vibrionaceae* Population Structure in Coastal Bacterioplankton. *ISME J*, **7**:509-519, Mar 2013.
- 8. CS Smillie*, MB Smith*, J Friedman, OX Cordero, **LA David**, EJ Alm. Ecology drives a global network of gene exchange connecting the human microbiome. *Nature*, **480**(7376):241-244, Dec 2011.
- 7. **LA David**, EJ Alm. Rapid evolutionary innovation during an Archean Genetic Expansion. *Nature*, **469**(7328):93-96, Jan 2011.
- 6. SP Preheim, Y Boucher, H Wildschutte, **LA David**, D Veneziano, EJ Alm and MF Polz. Metapopulation structure of Vibrionaceae among coastal marine invertebrates. *Environ Microbiol*, **13**:265-275, Jan 2011.
- 5. BJ Shapiro, **LA David**, J Friedman, EJ Alm. Looking for Darwin's footprints in the microbial world. *Trends Microbiol* **17**(5):196-204, May 2009.

- 4. BJ Dubin-Thaler, JM Hofman, Y Cai, H Xenias, I Spielman, AV Shneidman, **LA David**, HG Dobereiner, CH Wiggins, MP Sheetz. Quantification of Cell Edge Velocities and Traction Forces Reveals Distinct Motility Modules during Cell Spreading. *PLoS ONE*, **3**(11), Nov 2008.
- 3. DE Hunt*, **LA David***, D Gevers, SP Preheim, EJ Alm, MF Polz. Resource Partitioning and Sympatric Differentiation Among Closely Related Bacterioplankton. <u>Science</u>, **320**(5879):1081-85, May 2008.
- 2. **LA David**, CH Wiggins. Benchmarking of dynamic Bayesian Networks from stochastic time-series data. *Ann NY Acad Sci.*, **1115** 90-101, Dec 2007.
- CF Glenn, DK Chow, L David, C Cooke, M Gami, W Iser, K Hanselman, I Goldberg and CA Wolkow. Behavioral deficits during early states of aging in *Caenorhabditis elegans* result from locomotory deficits possibly linked to muscle frailty. <u>J Gerontol A Biol Sci Med Sci.</u>, 59(12):1251-60, Dec 2004.

GRANTS AWARDED

- Young Investigator Grant, Global Probiotics Council "A high-throughput pipeline to find probiotics that inhibit select gut microbes" (\$50K) PI: Lawrence David; 07/2014-06/2015.
- Research Fellowship, Alfred P. Sloan Foundation (\$50K) PI: Lawrence David; 02/2014-02/2016.
- Pilot Grant, Duke Center for the Genomics of Microbial Systems "Measurement and manipulation of environmental oxygen and the human gut microbiome" (\$20K) PIs: Lawrence David & Marc Deshusses; 10/2013-10/2014.
- Feasibility Grant, Harvard Center for AIDS Research. "The gut microbiota and microbial translocation in early HIV-1 infection." (\$78K) PIs: Lawrence David & Douglas Kwon; 07/2011-07/2012.
- Milton Fund, Harvard University. "The human gut microbiota and *Vibrio cholerae* infection." (\$39K) PI: Lawrence David: 01/2011-12/2012.

Fellowships & Awards

- 2014 Alfred P. Sloan Research Fellow
- 2013 Whitehead Scholar, Duke University
- 2010 Elected Junior Fellow, Harvard Society of Fellows
- 2009 MIT Whitaker Health Sciences Fund Fellowship
- 2005 National Defense Science and Engineering Graduate Fellowship
- 2005 Dept. of Energy Computational Science Graduate Fellowship [declined]
- 2005 Ford Foundation Diversity Predoctoral Fellowship [declined]
- 2005 Columbia Thomas "Pop" Harrington Award
- 2005 Columbia Robert E. and Claire S. Reiss Graduate Prize
- 2004 Barry M. Goldwater Scholarship
- 2002 Westchester Filipino-American Scholarship
- 2001 Columbia C.P. Davis Scholar

SERVICE

- Instructor (1 of 4): ideSHi Genomics Workshop, International Centre for Diarrhoeal Disease Research, Bangladesh, March 3-5, 2014
- Committee member: Duke CBB Graduate Admissions 2013-2014.
- Co-organizer: Duke GeMS Symposium, 2014.
- Co-organizer: Duke GeMS Microbiome Service, 2013-.
- Advisor: Boston Museum of Science, 2013.
- Founding member: ASM General Meeting Junior Advisory Group, 2011-2013.
- Referee: Biology Letters, BMC Bioinformatics, Clinical Infectious Diseases, Genome Research, InnoCentive, mBio, PLoS One, PLoS Pathogens, PNAS.

TEACHING & MENTORSHIP

- (Duke) Rachael Bloom, University Program in Genetics & Genomics, PhD student, 2014-
- (Duke) Aspen Reese, University Program in Ecology, PhD candidate, 2013-
- (Duke) Eric Mastrolonardo, undergrad research, 2013-
- (Duke) Sai Nimmagadda, undergrad research, 2013-
- (Duke) Calvin Lau, undergrad research, 2013-
- (Duke) Olivia Lin, undergrad research, 2013-2014
- (Harvard) Noah Apthorpe, undergrad research, under supervision of Peter Turnbaugh, 2012.
- (MIT) Shirley Galbiati, undergrad research, under supervision of Eric Alm, 2010.
- (MIT) Albert Wang, undergrad research, under supervision of Eric Alm, 2008-2010.
- (MIT) Alexandra Konings, undergrad research, under supervision of Eric Alm, 2008.
- (MIT) Graduate Resident Tutor, 2008-2010.
- (Cambridge Rindge & Latin) Science Olympiad and Biology tutor, 2007-2008.
- (MIT) Teaching assistant "Computation for Biological Engineers," 2006.

MEDIA

Print and online media coverage of research includes: the BBC, Bloomberg, Boston Globe, the Daily Mail, GenomeWeb, HealthDay, LiveScience, MSNBC, National Geographic: Phenomena, the New Scientist, NPR, Prevention Magazine, Radio In Vivo (WCOM), Science Daily, Science News, The Scientist, Scientific American, StoryCollider, USA Today, and Wired.

PRESENTATIONS

- 28. "Manipulating the human microbiome." Bollum Symposium, University of Minnesota, Minneapolis, MN, May 2014 (Invited talk)
- 27. "Manipulating the human microbiome." DCEG Fellows Training Symposium, NIH/NCI, Bethesda, MD, Mar 2014 (Invited talk)
- 26. "Perturbing the human microbiome." LadHyX Seminar, Ecole Polytechnique, Palaiseau, France, Feb 2014 (Talk)
- 25. "Perturbing the human microbiome." Pasteur Fellowship Visiting Scholar, Institut Pasteur, Paris, France, Feb 2014 (Invited talk)
- 24. "Daily changes to human-associated microbial communities." Duke University Program in Ecology Seminar, Durham, NC, Nov 2013 (Talk)

- 23. "Daily changes to human-associated microbial communities." Duke Marine Lab Fall Seminar, Beaufort, NC, Sep 2013 (Talk)
- 22. "Daily changes to human-associated microbial communities." Duke Computational Biology & Bioinformatics Seminar, Durham, NC, Sep 2013 (Talk)
- 21. "Daily changes to human-associated microbial communities." Duke Genomic and Personalized Medicine Forum, Durham, NC, Sep 2013 (Talk)
- 20. "Daily changes to human-associated microbial communities." Duke University Program in Genetics & Genomics Seminar, Durham, NC, Sep 2013 (Talk)
- 19. "There's an app for that: Tracking the lives of humans and the microbes living inside of them." Genomics Standards Consortium 15, Bethesda, MD, Apr 2013 (Invited talk)
- 18. "There's an app for that: Tracking the lives of humans and the microbes living inside of them." Museum of Science, Boston, MA, Mar 2013 (Talk)
- 17. "Dynamics and stability of human-associated microbial communities," National Institute of Health, Manila, Philippines, Feb 2013 (Talk)
- 16. "Resistance and response of the human gut microbiome to perturbation," Civil & Environmental Engineering Seminar, Duke University, Durham, NC, Dec 2012 (Invited talk)
- 15. "Succession dynamics of the human gut microbiota following Vibrio cholerae infection," icddr,b Seminar, Dhaka, Bangladesh, Aug 2012 (Talk)
- "Reproducible succession dynamics of the human gut microbiota following Vibrio cholerae infection," International Research in Infectious Diseases Annual Meeting, Bethesda, MD, May 2012 (Talk)
- 13. "Daily tracking of human health, behavior, and gut microbiota over one year," MIT Synthetic Biology Lunch, Cambridge, MA, Apr 2012 (Talk)
- 12. "Daily tracking of human health, behavior, and gut microbiota over one year," Northeastern University, Boston, MA, Mar 2012 (Talk)
- 11. "Daily tracking of human health, behavior, and gut microbiota over one year," Memorial Sloan-Kettering Cancer Center, New York, NY, Feb 2012 (Talk)
- 10. "Daily tracking of human health, behavior, and gut microbiota over one year," Duke University, Durham, NC, Feb 2012 (Talk)
- 9. "Daily tracking of human health, behavior, and gut microbiota over one year," University of North Carolina, Chapel Hill, NC, Feb 2012 (Talk)
- 8. "Stability of the human gut microbiota," Microbiome Research in the Boston Area Seminar, Boston, MA, Oct 2011 (Talk)
- 7. "Stability of the human gut microbiota," Microbiome Seminar, New York University, New York, NY, Oct 2011 (Talk)
- "Building models of Precambrian evolution from modern-day microbial gene families," Paleobiology in the Genomics Era Conference, JCVI, San Diego, CA, May 2011 (Invited talk)
- 5. "Of microbes and men: A year-long study of daily changes to the human microbiome," Quantitative Biology Workshop, MIT, Jan 2011 (Talk)

- 4. "Learning influential host factors and dynamical properties in human-associated microbial ecosystems," Human Microbiome Research Conference, St. Louis, MO, Aug 2010 (Poster; travel grant recipient)
- 3. "Rapid evolutionary innovation during an Archean Genetic Expansion," Biology of Genomes Conference, Cold Spring, NY, May 2010 (Poster; travel grant recipient)
- 2. "The HuGE project: Daily Variation In Human Microbiome Composition Over the Course of One Year," Merck-MIT Symposium, Boston, MA, Apr 2010 (Poster; winner best poster)
- 1. "Extracting insight from increasingly large microbial genomic datasets," Ginkgo BioWorks, Boston, MA, Mar 2010 (Talk)