Lawrence A. David

CONTACT INFORMATION

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EDUCATION

Massachusetts Institute of Technology

2005-2010

Ph.D., Computational and Systems Biology

Thesis: "Novel phylogenetic approaches to problems in microbial genomics"

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. Rare events can trigger switching between alternative stable states of the human microbiome. *Submitted*.
- 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. *Nature*, accepted 2013.
- 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

- 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; Awarded 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; Awarded 07/2011-07/2012.
- Milton Fund, Harvard University. "The human gut microbiota and Vibrio cholerae infection." (\$39K) PI: Lawrence David; Awarded 01/2011-12/2012.

Fellowships & Awards

- 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

Teaching & Mentorship

- Graduate resident tutor: Live-in mentor for MIT's MacGregor undergraduate dormitory, 2008-2010.
- Biology tutor and Science Olympiad mentor (High School): Cambridge Rindge and Latin, 2007-2008.

• Teaching assistant (Undergraduate): "Computation for Biological Engineers," taught by Profs. Eric Alm & Drew Endy, MIT, Fall 2006. Average overall rating of 6.5/7.0.

Research project supervision

- Albert Wang, "Construction of an archaeal supertree." Undergraduate researcher, Alm Lab, MIT: 2008-present.
- Shirley Galbiati, "Inferring the origins of photosynthesis." Undergraduate researcher, Alm Lab, MIT: 2010.
- Alexandra Konings, "Reconstructing ancient biological networks." Undergraduate researcher, Alm Lab, MIT: 2008.

SERVICE

- Advisor: Boston Museum of Science, 2013-present.
- Founding member: ASM General Meeting Junior Advisory Group, 2011-2013.
- Referee: Biology Letters, BMC Bioinformatics, Genome Research, PLoS One, PLoS Pathogens, PNAS.

Media

Print and online media coverage of research includes: the BBC, Boston Globe, the Daily Mail, GenomeWeb, LiveScience, MSNBC, the New Scientist, Science Daily, The Scientist, StoryCollider, USA Today, and Wired.

Presentations

- 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, April 2013 (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, March 2013 (Talk)
- 17. "Dynamics and stability of human-associated microbial communities," National Institute of Health, Manila, Philippines, February 2013 (Talk)
- 16. "Resistance and response of the human gut microbiome to perturbation," Civil & Environmental Engineering Seminar, Duke University, Durham, NC, December 2012 (Talk)
- 15. "Succession dynamics of the human gut microbiota following Vibrio cholerae infection," icddr,b Seminar, Dhaka, Bangladesh, August 2012 (Talk)
- 14. "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, April 2012 (Talk)

- 12. "Daily tracking of human health, behavior, and gut microbiota over one year," Northeastern University, Boston, MA, March 2012 (Talk)
- 11. "Daily tracking of human health, behavior, and gut microbiota over one year," Memorial Sloan-Kettering Cancer Center, New York, NY, February 2012 (Talk)
- 10. "Daily tracking of human health, behavior, and gut microbiota over one year," Duke University, Durham, NC, February 2012 (Talk)
- 9. "Daily tracking of human health, behavior, and gut microbiota over one year," University of North Carolina, Chapel Hill, NC, February 2012 (Talk)
- 8. "Stability of the human gut microbiota," Microbiome Research in the Boston Area Seminar, Boston, MA, October 2011 (Talk)
- 7. "Stability of the human gut microbiota," Microbiome Seminar, New York University, New York, NY, October 2011 (Talk)
- 6. "Building models of Precambrian evolution from modern-day microbial gene families," Paleobiology in the Genomics Era Conference, JCVI, San Diego, CA, May 2011 (Talk)
- 5. "Of microbes and men: A year-long study of daily changes to the human microbiome," Quantitative Biology Workshop, MIT, January 2011 (Talk)
- 4. "Learning influential host factors and dynamical properties in human-associated microbial ecosystems," Human Microbiome Research Conference, St. Louis, MO, August 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, April 2010 (Poster; winner best poster)
- 1. "Extracting insight from increasingly large microbial genomic datasets," Ginkgo BioWorks, Boston, MA, March 2010 (Talk)