# Philip (Phil) Arevalo

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Room 206

Chicago, Illinois 60637

EDUCATION

Ph.D., MIT

2012 - 2017

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Microbiology

Advisor: Martin Polz

Thesis: Horizontal gene transfer as a cohesive force in microbial populations.

Sc.B., Brown University, with honors

2007 - 2011

Applied Mathematics-Biology

Advisor: Jeremy Rich

Thesis: Diversity of anammox bactera in ocean sediments.

RESEARCH EXPERIENCE University of Chicago, Department of Ecology and Evolution

PI: Sarah Cobey

Ruth L. Kirschstein NRSA Postdoctoral Fellow Postdoctoral Scholar Starting September 2019

2017 - Present

- Modeling the effects infection history on susceptibility to influenza.
- Developing a computational method to identify influenza epitopes.
- Inferring factors responsible for the specificity of the human antibody response against influenza.

MIT, Microbiology Graduate Program

PI: Martin Polz

# NSF Graduate Research Fellow

2012 - 2017

- Developed a method to identify microbial populations in networks of horizontal gene transfer.
- Inferred the history of a polysaccharide degradation pathway in a group of marine bacteria.
- Assembled and analyzed a collection of over 700 marine bacterial isolates.

Honors and Awards

#### Ruth L. Kirschstein NRSA Postdoctoral Fellow

Starting September 2019

**NSF** Graduate Research Fellow

2012 - 2017

MIT Energy Initiative BP Energy Fellow

2012 - 2013

Povar Prize in Zoology and Physiology (awarded in recognition for outstanding academic achievements in physiology or

zoology at Brown University)

2011

Society of Sigma Xi

2011

Preprints in Review

- 1. Arevalo P., McLean H.Q., Belongia E.A., Cobey S. (2019). Earliest infections predict the age distribution of seasonal influenza A cases. medRxiv. doi:10.1101/19001875
- 2. Chase A.B., **Arevalo P.**, Brodie E.L., Polz M.F., Karaoz U., Martiny J.B.H. (2019). Sympatric and allopatric differentiation delineates population structure in free-living terrestrial bacteria. bioRxiv. doi:10.1101/644468

- Arevalo P.\*, VanInsberghe D.\*, Elsherbini J., Gore J., Polz M.F. (2019). A reverse ecology approach based on a biological definition of microbial populations. Cell, 178(4). doi:10.1016/j.cell.2019.06.033
- 2. Polzin J., **Arevalo P.**, Nussbaumer T., Polz M.F, Bright M. (2019). *Polyclonal symbiont populations in hydrothermal vent tubeworms and the environment*. Proceedings of the Royal Society B, 286(1896). doi:10.1098/rspb.2018.1281
- 3. Rich J.J., **Arevalo P.**, Chang B.X., Devol A.H., Ward B.B. (2018). Anaerobic ammonium oxidation (anammox) and denitrification in Peru margin sediments. Journal of Marine Systems. doi:10.1016/j.jmarsys.2018.09.007
- Arevalo P., VanInsberghe D., Polz M.F. (2018) A Reverse Ecology Framework for Bacteria and Archaea. In: Population Genomics: Microorganisms. Springer, Cham. doi: 10.1007/13836\_2018\_46.
- 5. Kauffman K., Hussain F., Yang J., **Arevalo P.**, Brown J., Cutler M., Kelly L., Polz M.F. (2018). A major lineage of non-tailed dsDNA viruses as unrecognized killers of marine bacteria. Nature, 554:118-122. doi:10.1038/nature25474
- Burks D.J., Norris S., Kauffman K.M., Joy A., Arevalo P., Azad R.K., Wildschutte H. (2017). Environmental vibrios represent a source of antagonistic compounds that inhibit pathogenic Vibrio cholerae and Vibrio parahaemolyticus strains. MicrobiologyOpen, 6(5). doi:10.1002/mbo3.504.
- Takemura A., Corzett C., Hussain F., Arevalo P., Datta M., Yu X., Le Roux F., Polz M.F. (2017). Natural resource landscapes of a marine bacterium reveal distinct fitness-determining genes across the genome. Environmental Microbiology, 19:2422–2433. doi:10.1111/1462-2920.13765.
- 8. Chase A.B., Arevalo P., Polz M.F., Berlemont R., Martiny J.B.H. (2016). Evidence for ecological flexibility in the cosmopolitan genus Curtobacterium. Frontiers in Microbiology, 7:1874. doi: 10.3389/fmicb.2016.01874
- 9. Hehemann J.-H.\*, **Arevalo P.\***, Datta M.S.\*, Yu X., Corzett C., Preheim S.P., Henschel A., Timberlake S., Alm E.J., Polz M.F. (2016). *Adaptive radiation by waves of gene transfer leads to fine-scale resource partitioning in marine microbes*. Nature Communications, 7. doi:10.1038/ncomms12860.

# Presentations and posters

- 1. **Arevalo P.**, VanInsberghe D., Elsherbini J., Gore J., Polz M.F. A reverse ecology approach based on a biological definition of microbial populations. July 2019. Microbiome. Cold Spring Harbor Laboratory. Oral presentation.
- 2. Arevalo P., McLean H.Q., Belongia E.A., Cobey S. Earliest infections predict the age distribution of seasonal influenza A cases. May 2019. Models of Infectious Disease Agent Study (MIDAS) Annual Meeting. Oral presentation.
- 3. Arevalo P., Vieira M., Cobey S. Original antigenic sin and antigenic maps. April 2019. Santa Fe Institute Working Group on Aging and Adaptation in Infectious Diseases. Oral presentation.
- 4. **Arevalo P.**, McLean H.Q., Belongia E.A., Cobey S. *The role of immune history in population-level influenza dynamics*. July 2018. Centers of Excellence for Influenza Research and Surveillance (CEIRS) Annual Network Meeting. Oral presentation.
- 5. **Arevalo P**. The role of immune history in population-level influenza dynamics. April 2018. University of Chicago Ecology & Evolution Darwin's Weekly. Oral presentation.
- 6. **Arevalo P**. Horizontal gene transfer as a cohesive force in microbial populations. April 2017. University of Chicago Ecology & Evolution Darwin's Weekly. Oral presentation.

- 7. **Arevalo P**. & Polz M.F. A biological definition for microbial populations and its application to a reverse ecology approach. August 2016. 16th International Symposium for Microbial Ecology. Evolution. Oral presentation.
- 8. **Arevalo P.** A biological definition for microbial populations and its application to a reverse ecology approach. Center for Microbiome Informatics and Therapeutics. June 2016. Work-in-progress meeting. Oral presentation.
- 9. Arevalo P., Wuchter C., Yang T-H., Coolen M., Sievert S. Stratified bacterial and archaeal communities across the oxygen minimum zone of the Eastern Tropical North Pacific. August 2012. 14th International Symposium for Microbial Ecology. Microbes in a changing ocean. Oral presentation.
- 10. Arevalo P., Sylva S., Toney C., Le Bris N., Seewald J., & Sievert S. Bacterial and archaeal community structure along a geochemical gradient in hydrothermally influenced sediments of Guaymas Basin, Gulf of California. August 2012. 14th International Symposium for Microbial Ecology. Microbial life in extreme environments. Poster session.
- 11. **Arevalo P.** & Rich J. Diversity and abundance of anammox bacteria along environmental gradients in the Peru margin. February 2011. American Society for Limnology and Oceanography Aquatic Sciences Meeting. Poster session.

TEACHING AND MENTORSHIP EXPERIENCE

## University of Chicago

Guest lecturer, Evolutionary and Genomic Medicine Winter 2018 and Winter 2019

#### **MIT**

Kaufman Teaching Certificate ProgramSummer 2017Rotation student advisor, Jai PadmakumarSpring 2017Teaching assistant, Microbial Population GenomicsFall 2016Rotation student advisor, Joseph ElsherbiniWinter 2015Teaching assistant, Microbial Genetics and EvolutionFall 2014

Centro de Biología Molecular, Managua, Nicaragua

Lecturer, bioinformatics workshop March 2014

SERVICE

#### University of Chicago

Organizer, Ecology and Evolution Theory Group Fall 2018 and Winter 2019

#### MIT

#### Graduate Resident Tutor

2014 - 2017

Ran regular study breaks, assisted in conflict resolution, and insured general well-being of undergraduate students in a residential setting.

Co-organizer, Microbial Engineering and Science Semi- 2014 - 2015 nar

#### **SPLASH** instructor

2013 and 2015

Taught high school students at annual event organized by members of the MIT community.

### Cambridge Rindge and Latin School, a local public high school

Guest lecturer, Biotechnology III November 2016

# **Brown University**

Writing Fellow 2008 - 2011

Provided writing support for ten students per semester focused on argumentation, style, and clarity.

OTHER RESEARCH EXPERIENCE

Woods Hole Oceanographic Institution

PI: Stefan Sievert

Research assistant 2011 - 2012

• Analyzed microbial community composition in oxygen minimum zones and hydrothermal vents.

Brown University, Department of Ecology and Evolutionary Biology

PI: Jeremy Rich

Research assistant

2009 - 2011

• Assessed the diversity of anammox bacteria in ocean sediments.