



Paula V. Welander

Associate Professor of Environmental Earth System Science

Bio

BIO

Paula Welander is a microbiologist who received her undergraduate degree from Occidental College in Los Angeles. She pursued her PhD studies in microbiology at the University of Illinois at Urbana-Champaign and completed her postdoctoral studies at MIT in the Departments of Biology and of Earth, Atmospheric, and Planetary Sciences. Paula joined the Stanford faculty in 2013 where her current research program is focused on understanding the biosynthesis and physiological function of “molecular fossils” or biomarkers in extant bacteria.

ACADEMIC APPOINTMENTS

- Associate Professor, Earth System Science
- Member, Bio-X

ADMINISTRATIVE APPOINTMENTS

- NASA Astrobiology Postdoctoral Fellow, Massachusetts Institute of Technology, (2012-2012)
- Research Scientist, Massachusetts Institute of Technology, (2011-2012)
- NSF Minority Postdoctoral Fellow, Massachusetts Institute of Technology, (2008-2011)
- National Science Foundation Graduate Research Fellow, University of Illinois at Urbana-Champaign, (2002-2005)
- Graduate College Fellow, University of Illinois at Urbana-Champaign, (2001-2007)
- Research Associate I, Beckman Research Institute, City of Hope, (1999-2001)
- Research Assistant I, California Institute of Technology, (1998-1999)

HONORS AND AWARDS

- Excellence in Teaching Award, Stanford University (2019)
- Faculty Early Career Development (CAREER) Award, National Science Foundation, Division of Earth Sciences (2018)
- Geobiology and Geomicrobiology Division Award for Outstanding Research (pre-tenure), Geological Society of America (2018)
- Hoagland Award Fund for Innovations in Undergraduate Teaching, Stanford University (2017)
- Terman Fellow, Stanford University (2014)
- Gabilan Faculty Fellow, Stanford University (2013)
- NASA Astrobiology Postdoctoral Fellowship, NASA Postdoctoral Program (2012)
- OGD 2010 Best Paper Award, Geochemical Society (2011)
- NSF Minority Postdoctoral Research Fellowship, National Science Foundation (2008)
- Mame Shiao Debbie Award, University of Illinois at Urbana-Champaign (2006)

- Outstanding Teaching Assistant, University of Illinois at Urbana-Champaign (2005-2006)
- Recipient, Gordon Conference Minority Student Travel Grant (2003)
- NSF Graduate Research Fellowship, National Science Foundation (2002)
- Graduate College Fellowship, University of Illinois at Urbana-Champaign (2001)
- Recipient, Declined, NIH MCB Training Grant Fellowship, University of Illinois at Urbana-Champaign (2001)
- Recipient, Howard Hughes Medical Institute Undergraduate Research Grant (1998)
- Fellow, Richter Fellowship (1997)
- Undergraduate Research Academic Support Program Grant, Occidental College (1997)
- Recipient, Cal Grant (1995 – 1998)

BOARDS, ADVISORY COMMITTEES, PROFESSIONAL ORGANIZATIONS

- Member, American Society of Microbiology (2002 - present)
- Member, European Association of Geochemistry (2013 - present)
- Member, American Geological Union (2013 - present)
- Editorial Board Member, Geobiology Journal (2014 - present)
- Member, Society for Advancement of Chicanos and Native Americans in Science (SACNAS) (2016 - present)

PROFESSIONAL EDUCATION

- Ph.D., University of Illinois at Urbana-Champaign, Urbana, IL , Microbiology (2007)
- M.S., University of Illinois at Urbana-Champaign, Urbana, IL , Microbiology (2003)
- B.A., Occidental College, Los Angeles, CA , Kinesiology (1998)

Research & Scholarship

CURRENT RESEARCH AND SCHOLARLY INTERESTS

Earth's history is marked with periods of dramatic atmospheric and climate fluctuations that have greatly affected life and its evolution. Much of our knowledge of how the biosphere responded to these ancient events stems from studies of faunal and floral fossils revealing profound changes in the abundance and diversity of macroscopic organisms. However, much less is known about microbial community responses to such dramatic environmental changes. This is one of the great challenges of geomicrobiology - how do we study microbial communities in the context of Earth's distant past?

Although microbes do not readily leave diagnostic morphological fossils, subtle microbial signatures are recorded in sedimentary rocks spanning billions of years. One such group of biosignatures are well-preserved lipid compounds that have specific biological origins and can function as biomarkers for the occurrence of specific microbes or environmental conditions at the time of deposition. However, many questions remain about the extant sources and the biosynthetic pathways of these lipids in modern organisms. In my research group, we focus on improving our understanding of the taxonomic distribution, the biosynthesis, and physiological function of biomarker lipids in extant microbes. Below are three biomarker areas that we are currently exploring in the Welander group:

1. Bacterial production of eukaryotic biomarkers. Eukaryotic biomarkers are specific lipid molecules that are considered diagnostic for certain eukaryotic organisms – from multicellular animals like sponges to unicellular eukaryotes such as protist. However, some bacterial species have been shown to produce these “eukaryotic” lipids and there are several open questions regarding how bacteria synthesize and utilize these lipids. We have identified unique bacterial proteins in aerobic methanotrophs for the synthesis of tetrahymanol, a cyclic triterpenoid primarily produced by ciliated protists, and sterols, essential lipids found in almost all eukaryotic cells. We are currently exploring the biochemical mechanisms behind these unique bacterial proteins and are also investigating the physiological function of these “eukaryotic” lipids in a variety of bacteria.

2. Identification of orphan biomarker sources. Orphan biomarkers are lipids identified in ancient or modern sediments for which there are no extant sources or the extant sources are not consistent with their occurrence in a specific environment or time period. One example is isoarborinol, an unusual pentacyclic triterpenol whose only known extant sources are certain flowering plants. Through our work, we have identified two novel arborinol lipids structurally similar to isoarborinol, which we named eudoraenol and adriaticol, in the marine bacterium *Eudoraea adriatica*. We are currently investigating the phylogenetic distribution of eudoraenol cyclase homologs in environmental metagenomes and are utilizing an *E. coli* heterologous expression system developed in our lab to express these cyclases in the lab.

3. Lipid biosynthesis in archaea. Glycerol dialkyl glycerol tetraethers (GDGTs) are unique archaeal membrane lipids that can function not just as biomarkers for archaea but can also be used as paleotemperature proxies. However, the pathway for GDGT synthesis has not been fully characterized and we have been attempting to identify the missing synthesis proteins in the thermoacidophile *Sulfolobus acidocaldarius*. Thus far, we have identified a novel protein, calditol synthase (Cds), necessary for modifying the glycosylated membrane head groups of *Sulfolobus*. We have also identified two novel proteins, GDGT-ring synthases, required for introducing the cyclopentane rings within the core GDGT structure. We are currently characterizing these GDGT biosynthesis proteins and continue to search for other unknown tetraether lipid biosynthesis proteins in archaea and bacteria.

Teaching

COURSES

2019-20

- Microbial Physiology: BIO 180, EARTHSYS 255, ESS 255, GEOLSCI 233A (Win)
- Topics in Geobiology: ESS 208, GEOLSCI 208 (Win)

2018-19

- Diversity and Inclusion in the Geosciences: EARTH 203 (Win)
- Microbial Physiology: BIO 180, EARTHSYS 255, ESS 255, GEOLSCI 233A (Win)
- Molecular Geomicrobiology Laboratory: BIO 142, EARTHSYS 143, ESS 143, ESS 243 (Spr)
- Topics in Geobiology: ESS 208, GEOLSCI 208 (Win)

2017-18

- Microbial Physiology: BIO 180, EARTHSYS 255, ESS 255, GS 233A (Aut)
- Molecular Geomicrobiology Laboratory: BIO 142, EARTHSYS 143, ESS 143, ESS 243 (Spr)
- Topics in Earth System Science: ESS 301 (Spr)
- Topics in Geobiology: ESS 208, GS 208 (Win)

2016-17

- The Invisible Majority: The Microbial World That Sustains Our Planet: EARTHSYS 44N (Spr)

STANFORD ADVISEES

Postdoctoral Faculty Sponsor

Amber Bonds

Doctoral (Program)

Malory Brown, Alysha Lee, Marisa Mayer, Hanon McShea

GRADUATE AND FELLOWSHIP PROGRAM AFFILIATIONS

- Biology (School of Humanities and Sciences) (Phd Program)

Publications

PUBLICATIONS

- **Deciphering the evolutionary history of microbial cyclic triterpenoids** *FREE RADICAL BIOLOGY AND MEDICINE*
Welander, P. V.
2019; 140: 270–78
- **Soil exchange rates of COS and (COO)-O-18 differ with the diversity of microbial communities and their carbonic anhydrase enzymes** *ISME JOURNAL*
Meredith, L. K., Ogee, J., Boye, K., Singer, E., Wingate, L., von Sperber, C., Sengupta, A., Whelan, M., Pang, E., Keiluweit, M., Brueggemann, N., Berry, J. A., Welander, et al
2019; 13 (2): 290–300
- **GDGT cyclization proteins identify the dominant archaeal sources of tetraether lipids in the ocean.** *Proceedings of the National Academy of Sciences of the United States of America*
Zeng, Z., Liu, X. L., Farley, K. R., Wei, J. H., Metcalf, W. W., Summons, R. E., Welander, P. V.
2019
- **Calditol-linked membrane lipids are required for acid tolerance in *Sulfolobus acidocaldarius*.** *Proceedings of the National Academy of Sciences of the United States of America*
Zeng, Z., Liu, X., Wei, J. H., Summons, R. E., Welander, P. V.
2018
- **C-4 sterol demethylation enzymes distinguish bacterial and eukaryotic sterol synthesis.** *Proceedings of the National Academy of Sciences of the United States of America*
Lee, A. K., Banta, A. B., Wei, J. H., Kiemle, D. J., Feng, J., Giner, J. L., Welander, P. V.
2018; 115 (23): 5884–89
- **Synthesis of arborane triterpenols by a bacterial oxidosqualene cyclase** *PROCEEDINGS OF THE NATIONAL ACADEMY OF SCIENCES OF THE UNITED STATES OF AMERICA*
Banta, A. B., Wei, J. H., Gill, C. C., Giner, J., Welander, P. V.
2017; 114 (2): 245-250
- **Microbial communities and organic biomarkers in a Proterozoic-analog sinkhole.** *Geobiology*
Hamilton, T. L., Welander, P. V., Albrecht, H. L., Fulton, J. M., Schaperdoth, I., Bird, L. R., Summons, R. E., Freeman, K. H., Macalady, J. L.
2017; 15 (6): 784–97
- **Fractionation of the methane isotopologues (CH₄)-C-13, (CH₃D)-C-12, and (CH₃D)-C-13 during aerobic oxidation of methane by *Methylococcus capsulatus* (Bath)** *GEOCHIMICA ET COSMOCHIMICA ACTA*
Wang, D. T., Welander, P. V., Ono, S.
2016; 192: 186-202
- **Sterol Synthesis in Diverse Bacteria** *FRONTIERS IN MICROBIOLOGY*
Wei, J. H., Yin, X., Welander, P. V.
2016; 7
- **A distinct pathway for tetrahymanol synthesis in bacteria** *PROCEEDINGS OF THE NATIONAL ACADEMY OF SCIENCES OF THE UNITED STATES OF AMERICA*
Banta, A. B., Wei, J. H., Welander, P. V.
2015; 112 (44): 13478-13483
- **Methane Oxidation and Molecular Characterization of Methanotrophs from a Former Mercury Mine Impoundment.** *Microorganisms*
Baesman, S. M., Miller, L. G., Wei, J. H., Cho, Y., Matys, E. D., Summons, R. E., Welander, P. V., Oremland, R. S.
2015; 3 (2): 290-309
- **Elucidation of the *Burkholderia cenocepacia* hopanoid biosynthesis pathway uncovers functions for conserved proteins in hopanoid-producing bacteria** *ENVIRONMENTAL MICROBIOLOGY*
Schmerk, C. L., Welander, P. V., Hamad, M. A., Bain, K. L., Bernards, M. A., Summons, R. E., Valvano, M. A.
2015; 17 (3): 735-750

- **Diverse capacity for 2-methylhopanoid production correlates with a specific ecological niche** *ISME JOURNAL*
Ricci, J. N., Coleman, M. L., Welander, P. V., Sessions, A. L., Summons, R. E., Spear, J. R., Newman, D. K.
2014; 8 (3): 675-684
- **Molecular indicators of microbial diversity in oolitic sands of Highborn Cay, Bahamas** *Geobiology*
Edgecomb, V. P., Bernhard, J. M., Beaudoin, D., Pruss, S., Welander, P. V., Shubotz, F., Mehay, S., Gillespie, A. L., Summons, R. E.
2013; 11: 234-251
- **Identification and quantification of polyfunctionalized hopanoids by high temperature gas chromatography-mass spectrometry.** *Organic geochemistry*
Sessions, A. L., Zhang, L., Welander, P. V., Doughty, D., Summons, R. E., Newman, D. K.
2013; 56: 120-30
- **Identification and characterization of Rhodopseudomonas palustris hopanoid biosynthesis mutants** *Geobiology*
Welander, P. V., Doughty, D. M., Wu, C. H., Mehay, S., Summons, R. E., Newman, D. K.
2012; 10: 163-177
- **Discovery, taxonomic distribution, and phenotypic characterization of a gene required for 3-methylhopanoid production.** *Proceedings of the National Academy of Sciences of the United States of America*
Welander, P. V., Summons, R. E.
2012; 109 (32): 12905-10
- **Identification of the bacteriochlorophylls, carotenoids, quinones, lipids, and hopanoids of Candidatus Chloracidobacterium thermophilum** *Journal of Bacteriology*
Garcia Costas, A. M., Tsukatani, Y., Rijpstra, W. I., Schouten, S., Welander, P. V., Summons, R. E., Bryant, D. A.
2012; 194: 1158-68
- **Using Taguchi-based statistics to produce robust PCR results** *Bioprocessing*
Welander, P., Cantin, E., Lundberg, P.
2011; 10: 22-26
- **Identification of a methylase required for 2-methylhopanoid production and implications for the interpretation of sedimentary hopanes.** *Proceedings of the National Academy of Sciences of the United States of America*
Welander, P. V., Coleman, M. L., Sessions, A. L., Summons, R. E., Newman, D. K.
2010; 107 (19): 8537-42
- **The continuing puzzle of the great oxidation event** *Current Biology*
Sessions, A. L., Doughty, D. M., Welander, P. V., Summons, R. E., Newman, D. K.
2009; 19: R567-R574
- **Hopanoids play a role in membrane integrity and pH homeostasis in Rhodopseudomonas palustris TIE-1.** *Journal of bacteriology*
Welander, P. V., Hunter, R. C., Zhang, L., Sessions, A. L., Summons, R. E., Newman, D. K.
2009; 191 (19): 6145-56
- **Mutagenesis of the C1 oxidation pathway in Methanosarcina barkeri: new insights into the Mtr/Mer bypass pathway** *Journal of Bacteriology*
Welander, P. V., Metcalf, W. W.
2008; 190: 1928-1936
- **Tumor necrosis factor (TNF) protects resistant C57BL/6 mice against herpes simplex virus-induced encephalitis independently of signaling via TNF receptor 1 or 2** *Journal of Virology*
Lundberg, P., Welander, P. V., Edwards, 3rd, C. K., van Rooijen, N., Cantin, E.
2007; 81: 1451-1460
- **Loss of the mtr operon in Methanosarcina blocks growth on methanol, but not methanogenesis, and reveals an unknown methanogenic pathway** *Proceedings of the National Academy of Sciences*
Welander, P. V., Metcalf, W. W.
2005; 102: 10664-10669
- **Herpes simplex virus type 1 DNA is immunostimulatory in vitro and in vivo** *Journal of Virology*
Lundberg, P., Welander, P., Han, X., Cantin, E.
2003; 77: 11158-11169

- **A locus on mouse chromosome 6 that determines resistance to herpes simplex virus also influences reactivation, while an unlinked locus augments resistance of female mice** *Journal of Virology*

Lundberg, P., Welander, P., Openshaw, H., Nalbandian, C., Edwards, C., Moldawer, L., Cantin, E.
2003; 77: 11661-11673