

## Luke R. Thompson, Ph.D.

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Research Professor & NGI Program Coordinator  
Northern Gulf Institute (NGI), Mississippi State University  
NOAA Atlantic Oceanographic and Meteorological Laboratory (AOML)  
4301 Rickenbacker Causeway, Miami, FL 33149

## Research Interests

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Microbial ecology and biogeography | eDNA | Biodiversity monitoring | Biological big data | Environmental metadata

## Education

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Massachusetts Institute of Technology, *Cambridge, Massachusetts*

*Ph.D.* Biology 2003–2010  
Microbiology *Specialization*  
Thesis: Auxiliary metabolic genes in viruses infecting marine cyanobacteria ([link to full text](#))  
Advisors: Sallie W. Chisholm and JoAnne Stubbe

Stanford University, *Stanford, California*

*B.S.* Biological Sciences *with Honors* 1998–2002  
Chemistry *Minor*

## Previous Positions

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Scripps Institution of Oceanography, *La Jolla, California*

*Lecturer (part-time)* Graduate-level Python course 2015–2018

University of California, *San Diego, California* & University of Colorado, *Boulder, Colorado*

*Postdoctoral Scholar* Rob Knight Lab 2013–2016

King Abdullah University of Science and Technology, *Thuwal, Saudi Arabia*

*Postdoctoral Fellow* Red Sea Research Center 2010–2013

## Publications

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Google Scholar Profile: [goo.gl/Te4va1](https://scholar.google.com/citations?user=Te4va1)

ORCID ID: [orcid.org/0000-0002-3911-1280](https://orcid.org/0000-0002-3911-1280)

60. Anderson, S. R., Silliman, K., Barbero, L., Gomez, F. A., Stauffer, B. A., Schnetzer, A., Kelble, C. R., **Thompson, L. R.** (2026). Microbial community dynamics over large spatial and environmental gradients in a subtropical ocean basin. *Applied and Environmental Microbiology*, 92(2), e0188925. <https://doi.org/10.1128/aem.01889-25>
59. Silliman, K., Wilcox Talbot, L. A., Applegate, M., Aichinger Dias, L., Garrison, L. P., Grace, M., Paterson, C. N., **Thompson, L. R.**, Vollmer, N. L., Rosel, P. E. (2026). Evaluating prey availability for the Rice's whale (*Balaenoptera ricei*) based on environmental DNA. *Ecology and Evolution*, 16(1), e72789. <https://doi.org/10.1002/ece3.72789>
58. Takahashi, M., Paupério, J., Woollard, P., Frøslev, T., **Thompson, L.**, Silliman, K., Formel, S., Hunter, C., Babo, C., Deiner, K., Schriml, L., Berry, O. (2026). A TDWG and GSC collaborative initiative to develop an

environmental DNA (eDNA) metadata checklist. *Biodiversity Information Science and Standards*, 10(e185867), e185867. <https://doi.org/10.3897/biss.10.185867>

57. Bravo, G., Bigatti, G., Lozada, M., **Thompson, L.**, Livore, J., Mendez, M., Arribas, L., Bigatti, L., Christian, T., Macaya, E., Londoño-Cruz, E., Moity, N., Cruz-Motta, J., Flores, A., Vélez-Rubio, G., Palomo, M., Cordeiro, C., Pellizzari, F., Cárdenas-Calle, M., Kanhai, L. D., Vivar Linares, I., Gil-Kodaka, P., Martinez, L., Sugliano, P., Trigo, A., Zottola, J., Blanco, D., Tricase, M., Bravo, N., Degrati, M., Tavano Formigo, C., Muller-Karger, F., Montes, E. (2025). Expanding the scale and scope of the Marine Biodiversity Observation Network Pole to Pole of the Americas: Merging rocky intertidal biodiversity surveys with environmental DNA and plankton imaging applications. *Research Ideas and Outcomes*, 11(e163815), e163815. <https://doi.org/10.3897/rio.11.e163815>
56. Kelliher, J. M., Aljumaah, M., Bordenstein, S. R., Brister, J. R., Chain, P. S. G., Dundore-Arias, J. P., Emerson, J. B., Fernandes, V. M. C., Flores, R., Gonzalez, A., Hansen, Z. A., Hatcher, E. L., Jackson, S. A., Kellogg, C. A., Madupu, R., Miller, C. M. L., Mirzayi, C., Moustafa, A. M., Mungall, C., Oliver, A., Pariente, N., Pett-Ridge, J., Record, S., Reji, L., Reysenbach, A.-L., Rich, V. I., Richardson, L., Schriml, L. M., Shabman, R. S., Sierra, M. A., Sullivan, M. B., Sundaramurthy, P., Thibault, K. M., **Thompson, L. R.**, Tighe, S., Vereen, E., Eloë-Fadrosh, E. A. (2025). Microbiome data management in action workshop: Atlanta, GA, USA, June 12-13, 2024. *Environmental Microbiome*, 20(1), 40. <https://doi.org/10.1186/s40793-025-00702-9>
55. Kelliher, J. M., Mirzayi, C., Bordenstein, S. R., Oliver, A., Kellogg, C. A., Hatcher, E. L., Berg, M., Baldrian, P., Aljumaah, M., Miller, C. M. L., Mungall, C., Novak, V., Palucki, A., Smith, E., Tabassum, N., Bonito, G., Brister, J. R., Chain, P. S. G., Chen, M., Degregori, S., Dundore-Arias, J. P., Emerson, J. B., Moreira C Fernandes, V., Flores, R., Gonzalez, A., Hansen, Z. A., Jackson, S. A., Moustafa, A. M., Northen, T. R., Pariente, N., Pett-Ridge, J., Record, S., Reji, L., Reysenbach, A.-L., Rich, V. I., Richardson, L., Roux, S., Schriml, L. M., Shabman, R. S., Sierra, M. A., Sullivan, M. B., Sundaramurthy, P., Thibault, K. M., **Thompson, L. R.**, Tighe, S., Vereen, E., STREAMS Consortium, Eloë-Fadrosh, E. A. (2025). STREAMS guidelines: standards for technical reporting in environmental and host-associated microbiome studies. *Nature Microbiology*, 10(12), 3059–3068. <https://doi.org/10.1038/s41564-025-02186-2>
54. Muller-Karger, F. E., Canonico, G., VanSumeren, H., Myers, C., Ogburn, M. B., Hammerschlag, N., Pendleton, L., Motyka, J., **Thompson, L. R.**, Montes, E., McEachron, L., Pearlman, J., Brenner, J., Williams, B., Kavanaugh, M. T., Biddle, M. (2025). A prosperous economy requires ocean biodiversity information, technology, and innovation. *Marine Technology Society Journal*, 59(3), 7–8. <https://doi.org/10.4031/mts.59.3.7>
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52. Takahashi, M., Frøsvlev, T. G., Paupério, J., Thalinger, B., Klymus, K., Helbing, C. C., Villacorta-Rath, C., Silliman, K., **Thompson, L. R.**, Jungbluth, S. P., Yong, S. Y., Formel, S., Jenkins, G., Laporte, M., Deagle, B., Rajbhandari, S., Jeppesen, T. S., Bissett, A., Jerde, C., Hahn, E. E., Schriml, L. M., Hunter, C., Newman, P., Woollard, P., Harper, L. R., Dunn, N., West, K., Haderlé, R., Wilkinson, S., Acharya-Patel, N., Lopez, M. L. D., Cochrane, G., Berry, O. (2025). A Metadata Checklist and data formatting guidelines to make eDNA FAIR (Findable, Accessible, Interoperable, and Reusable). *Environmental DNA (Hoboken, N.J.)*, 7(3), e70100. <https://doi.org/10.1002/edn3.70100>
51. Gold, Z., Kelly, R. P., Shelton, A. O., Thompson, A. R., Goodwin, K. D., Gallego, R., Parsons, K. M., **Thompson, L. R.**, Kacev, D., Barber, P. H. (2024). Archived DNA reveals marine heatwave-associated shifts in fish assemblages. *Environmental DNA (Hoboken, N.J.)*, 6(1). <https://doi.org/10.1002/edn3.400>

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48. Silliman, K., Anderson, S., Storo, R., **Thompson, L.** (2023). A case study in sharing marine eDNA metabarcoding data to OBIS. *Biodiversity Information Science and Standards*, 7(e111048), e111048. <https://doi.org/10.3897/biss.7.111048>
47. **Thompson, L. R.**, Thielen, P. (2023). Decoding dissolved information: environmental DNA sequencing at global scale to monitor a changing ocean. *Current Opinion in Biotechnology*, 81(102936), 102936. <https://doi.org/10.1016/j.copbio.2023.102936>
46. Anderson, S. R., **Thompson, L. R.** (2022). Optimizing an enclosed bead beating extraction method for microbial and fish environmental DNA. *Environmental DNA (Hoboken, N.J.)*, 4(2), 291–303. <https://doi.org/10.1002/edn3.251>
45. Clayton, S., Alexander, H., Graff, J. R., Poulton, N. J., **Thompson, L. R.**, Benway, H., Boss, E., Martiny, A. (2022). Bio-GO-SHIP: The time is right to establish global repeat sections of ocean biology. *Frontiers in Marine Science*, 8, 767443. <https://doi.org/10.3389/fmars.2021.767443>
44. Den Uyl, P. A., **Thompson, L. R.**, Errera, R. M., Birch, J. M., Preston, C. M., Ussler, W., III, Yancey, C. E., Chaganti, S. R., Ruberg, S. A., Doucette, G. J., Dick, G. J., Scholin, C. A., Goodwin, K. D. (2022). Lake Erie field trials to advance autonomous monitoring of cyanobacterial harmful algal blooms. *Frontiers in Marine Science*, 9(1021952). <https://doi.org/10.3389/fmars.2022.1021952>
43. Jorge, F., Brealey, J. C., Brindley, P. J., Buysse, M., Cantacessi, C., Duron, O., Fichorova, R., Fitzpatrick, C. R., Hahn, M., Hunter, C., Hervé, V., Knoll, L. J., Kohl, K. D., Lalle, M., Lukeš, J., Martínez, J. M., Perkins, S. L., Poulin, R., Rosario, K., Schneider, A. C., Schriml, L. M., **Thompson, L. R.**, Walls, R. L., Dheilly, N. M. (2022). MlXS-SA: a MlXS extension defining the minimum information standard for sequence data from symbiont-associated micro-organisms. *ISME Communications*, 2(1), 9. <https://doi.org/10.1038/s43705-022-00092-w>
42. Lim, S. J., **Thompson, L. R.**, Young, C. M., Gaasterland, T., Goodwin, K. D. (2022). Dominance of *Sulfurospirillum* in metagenomes associated with the methane ice worm (*Sirsoe methanicola*). *Applied and Environmental Microbiology*, 88(15), e0029022. <https://doi.org/10.1128/aem.00290-22>
41. Shaffer, J. P., Nothias, L.-F., **Thompson, L. R.**, Sanders, J. G., Salido, R. A., Couvillion, S. P., Brejnrod, A. D., Lejzerowicz, F., Haiminen, N., Huang, S., Lutz, H. L., Zhu, Q., Martino, C., Morton, J. T., Karthikeyan, S., Nothias-Esposito, M., Dührkop, K., Böcker, S., Kim, H. W., Aksenov, A. A., Bittremieux, W., Minich, J. J., Marotz, C., Bryant, M. M., Sanders, K., Schwartz, T., Humphrey, G., Vásquez-Baeza, Y., Tripathi, A., Parida, L., Carrieri, A. P., Beck, K. L., Das, P., González, A., McDonald, D., Ladau, J., Karst, S. M., Albertsen, M., Ackermann, G., DeReus, J., Thomas, T., Petras, D., Shade, A., Stegen, J., Song, S. J., Metz, T. O., Swafford, A. D., Dorrestein, P. C., Jansson, J. K., Gilbert, J. A., Knight, R., Earth Microbiome Project 500 (EMP500) Consortium. (2022). Standardized multi-omics of Earth's microbiomes reveals microbial and metabolite diversity. *Nature Microbiology*, 7(12), 2128–2150. <https://doi.org/10.1038/s41564-022-01266-x>
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38. Formel, N., Enochs, I. C., Sinigalliano, C., Anderson, S. R., **Thompson, L. R.** (2021). Subsurface automated samplers for eDNA (SASe) for biological monitoring and research. *HardwareX*, 10(e00239), e00239. <https://doi.org/10.1016/j.ohx.2021.e00239>

37. Lim, S. J., **Thompson, L. R.** (2021). Mitohelper: A mitochondrial reference sequence analysis tool for fish eDNA studies. *Environmental DNA* (Hoboken, N.J.), 3(4), 706–715. <https://doi.org/10.1002/edn3.187>
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## Invited & Contributed Talks

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54. 2026 Ocean Sciences Meeting, Glasgow, Scotland, February 22-27, 2026, “Supporting NOAA Ocean Exploration with a Comprehensive Environmental DNA Data Analysis and Sharing Platform” (contributed).
53. 2025 NOPP Ocean Life Forum 2.0: Improving Observations of Marine Life through Investment in Innovation, Johns Hopkins University Applied Physics Laboratory, Laurel, MD, September 15th-16th, 2025 (invited).
52. National Academies of Sciences, Engineering, and Medicine, Gulf Research Program, Meeting of Experts: Genomics Approaches to Understanding, Managing, and Monitoring Biodiversity in the U.S. Gulf, New Orleans, June 11, 2025, “Federal Genomics Perspectives” (invited).
51. 2025 NOAA Enterprise Data Management Workshop, May 20, 2025, “The NOAA Ocean DNA Explorer (NODE) for Environmental DNA Data” (contributed).
50. 5th Marine Biodiversity Workshop: From the Sea to the Cloud, Puerto Madryn and Puerto Pirámides, Patagonia, Argentina, April 1, 2025, “Environmental DNA for Biodiversity Monitoring” (invited).
49. 2nd Australian and New Zealand Environmental DNA Conference, Wellington, New Zealand, February 19, 2025, “Demonstrating marine eDNA technology for monitoring biodiversity, characterizing ocean biogeochemistry, and enhancing other monitoring methods” (invited).
48. 2024 GOMO Arctic Research Program PI Meeting, Washington, DC, December 11, 2024, “Integrated Arctic Ecosystem Toolkit to Enhance Forecasting for Climate Resilience” (invited).
47. Rosenberg Institute Seminar Series, Estuary & Ocean Science Center, San Francisco State University, September 11, 2024, “Plankton community dynamics in the northern Gulf of Mexico” (invited).
46. Microbial Data and Tools without Borders: Advancing an Open Science Ecosystem (ASM Microbe 2024 mini-conference), Atlanta, GA, June 13, 2024, “Panel discussion: Advancing a vibrant research ecosystem across stakeholders” (invited).
45. Gulf of Mexico Coastal Acidification Network (GCAN), April 30, 2024, “Current and projected basin-wide plankton responses to increasing temperature and acidification in the Gulf of Mexico” (invited).
44. NOAA Living Marine Resources Cooperative Science Center (LMRCSC) Science Meeting, Miami, FL, April 23, 2024, “Methods and Applications of Environmental DNA Technology to Support NOAA’s Mission” (invited).
43. Nova Southeastern University, Department of Biological Sciences, Dania Beach, FL, September 22, 2023, “Plankton biodiversity patterns in the Gulf of Mexico and beyond” (invited).
42. Northeast Coastal Acidification Network (NECAN) Webinar on Current and New Technology, Sensors, and Methods, July 11, 2023, “Environmental DNA methods for assessing ecosystem responses of Gulf of Mexico prokaryotic and eukaryotic communities to ocean acidification” (invited).
41. Ocean Carbon and Biogeochemistry 2023 Summer Workshop, Woods Hole, Massachusetts, June 15, 2023, “Sustained observations of global ocean biology – session introduction” (invited).
40. NOAA Ocean Acidification Working Group, May 18, 2023, on the responses of marine protist and bacterial communities to OA in the Gulf of Mexico (data from GOMECC-4) (invited).
39. Metabolomics Association of North America Microbiome Interest Group, February 13, 2023, “Behind the Paper from the Knight Lab and EMP Group at UC San Diego” (invited).
38. Bio-GO-SHIP eDNA preservation/extraction intercalibration exercise on Wednesday, January 25, for a BioGeoSCAPES webinar hosted by Woods Hole Oceanographic Institution.
37. NOAA Ocean Acidification Community Meeting & Mini Symposium, La Jolla, California, January 4-6, 2023, “Patterns of microbial and plankton diversity from eDNA collected on GOMECC-4” (poster).
36. Atlantic International Research Centre (AIR Centre) Marine Biodiversity Networking Fridays, December 9, 2022, “Bio-GO-SHIP: expanding biological ocean observations for plankton ecosystem science and monitoring to the global scale” (invited).
35. ICES Annual Science Conference, Dublin, Ireland, September 19, 2022, “Machine learning-based taxonomic classification of DNA sequences in marine metagenomes” (contributed).
34. World Aquatic Veterinary Medical Association (WAVMA), August 26, 2022, “Environmental DNA for the study of ocean biodiversity from microbes to vertebrates” (invited).
33. NOAA 'Omics Seminar Series, July 20, 2022, “Global marine biodiversity monitoring through partnership and innovation” (contributed).

32. American Society for Microbiology General Meeting (ASM Microbe 2022), Washington, DC, June 12, 2022, “Crowdsourcing the microbiome: no guts, no glory” (invited).
31. Michigan State University, East Lansing, Michigan, April 21, 2022, “Environmental DNA for the study of ocean biodiversity from microbes to vertebrates” (invited).
30. Introduction to Environmental DNA for Applications in Florida, Webinar, January 28 and February 4, 2021, “NOAA eDNA Research in South Florida and the Gulf of Mexico” (invited).
29. Institute of Food and Agricultural Sciences, University of Florida, Davie, March 4, 2020, “Environmental DNA tools and applications for oceans and the Great Lakes” (invited).
28. Scripps Institution of Oceanography, November 1, 2019, “Environmental DNA tools and applications for oceans and the Great Lakes” (invited).
27. University of Southern Mississippi, Hattiesburg, October 18, 2019, “Applications and tools for environmental DNA” (invited).
26. OceanObs’19, Honolulu, Hawaii, September 18, 2019, Panelist for Special Session “Incorporating environmental DNA into global ocean observing systems: opportunities and challenges” (invited).
25. 1st Parasite Microbiome Project Workshop, Clearwater, Florida, January 11, 2019, “Standards-enabled large-scale surveys of Earth’s microbial communities” (invited).
24. International Microbiome and Metagenomics Standards Alliance, November 8, 2018, “Standards-enabled metagenomics and metabolomics of Earth’s microbial communities” (invited).
23. San Diego State University, October 18, 2018, “Multi-omics of Earth’s microbial communities” (invited).
22. 148th Annual Meeting of the American Fisheries Society, Atlantic City, August 20, 2018, “Computational workflows for rapid and customizable analysis of amplicon sequencing data from environmental DNA” (contributed).
21. 17th International Symposium on Microbial Ecology, Leipzig, Germany, August 16, 2018, “EMP500: multi-omics of diverse microbial environments in the Earth Microbiome Project” (invited).
20. Department of Earth System Science, Stanford University, May 30, 2018, “The Earth Microbiome Project: investigating the structure of microbial diversity on Earth” (invited).
19. DOE Joint Genome Institute, Walnut Creek, May 29, 2018, “The Earth Microbiome Project: lessons from a scientifically crowdsourced microbiome survey” (invited).
18. University of Oulu, Finland, March 22, 2018, “The Earth Microbiome Project: revealing global patterns of microbial distribution” (invited).
17. ‘Omic Biomonitoring Workshop, Max-Planck-Institute for Marine Microbiology, Bremen, Germany, February 21, 2018, “Global characterization of microbial taxonomic and functional diversity in the Earth Microbiome Project” (invited).
16. San Diego Microbiology Group, January 11, 2018, “Mapping the microbes of Earth” (invited).
15. Scripps Institution of Oceanography, December 1, 2017, “A communal catalogue of Earth’s microbes” (invited).
14. TDWG 2017: Biodiversity Information Standards, Ottawa, Canada, October 2, 2017 (contributed).
  - Sachs, J., **L. Thompson**, N. El-Kayssi & S. Bilku. (2017). Using MlxS: an implementation report from two metagenomic information systems. *Biodivers Inform Sci Stand* 1:e20637, <https://doi.org/10.3897/tdwgproceedings.1.20637>.
13. ICES Annual Science Conference, Fort Lauderdale, September 18, 2017, “The Earth Microbiome Project: lessons from a massive metagenetic survey” (contributed).
12. GRDI-Ecobiomics: Workshop on Environmental and Protocol Metadata, Ottawa, Canada, May 31, 2016, “Environmental metadata and the Earth Microbiome Project” (invited/webinar).
11. Sharp HealthCare Obesity Crisis Conference, San Diego, May 13, 2016, “Healthy gut and microbiomes” (invited).
10. Microbiome Connections to the Environment and Health, University of California, Irvine, September 25, 2015, “Computational tools for environmental genomics and the Earth Microbiome Project” (invited).
9. Scripps Institution of Oceanography, February 13, 2015, “Metagenomic analysis and the Earth Microbiome Project” (invited).

8. 2nd International Symposium on Sponge Microbiology, Baltimore, October 27, 2014, “The Earth Microbiome Project: early successes and challenges” and panel member of “Group discussion: Where to from here?” (invited).
7. 15th International Symposium on Microbial Ecology, Seoul, South Korea, August 29, 2014, “Mapping the genetic diversity and physicochemical space of the Red Sea basin” (contributed).
6. Pacific Northwest National Laboratory, November 1, 2013, “Marine metagenomics: the Red Sea and the Earth Microbiome Project” (invited).
5. Woods Hole Oceanographic Institution, July 30, 2012, “Let my microbes go! The Red Sea and stress tolerance strategies of *Prochlorococcus*, cyanophage, and SAR11” (invited).
4. Winter Enrichment Program, KAUST, January 23, 2012, “The coming big data singularity” (invited).
3. Integrative Microbial Ecology Workshop, KAUST, June 26, 2010, “Auxiliary metabolic genes in viruses infecting marine cyanobacteria” (invited).
2. DOE Genomics:GTL (Genomes-To-Life) Contractor–Grantee Workshop VII, Bethesda, February 9, 2009, “Viruses hijacking cyanobacterial carbon metabolism” (contributed).
1. Boston Bacterial Meeting, June 22, 2007, “Transaldolase in viruses infecting *Prochlorococcus*: hijacking host carbon metabolism with a non-cyanobacterial enzyme” (contributed).

## News Features

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- Interviewed for article in The Guardian, “How eDNA technology is changing the game for protecting ocean species”, June 2023
- Interviewed for American Blue Economy Podcast episode “Exploring the Wonders of Biotechnology in the American Blue Economy”, February 2023
- UC San Diego News Center on 2018 *Nature Communications* article “Taxon-specific aerosolization of bacteria and viruses in an experimental ocean–atmosphere mesocosm”
- News features on 2017 *Nature* article “A communal catalogue reveals Earth’s multiscale microbial diversity”:
  - *Nature News & Views*: J. Raes. (2017). Microbiology: Crowdsourcing Earth’s microbes. *Nature* 551:446–447, <https://doi.org/10.1038/nature24756>.
  - *WIRED*, *The Scientist*, *Science Friday*, *Ars Technica*, and others.
- PNAS Plus Author Summary of 2011 *PNAS* article “Phage auxiliary metabolic genes and the redirection of cyanobacterial host carbon metabolism”
- Faculty of 1000 recommendation for 2011 *PNAS* article “Phage auxiliary metabolic genes and the redirection of cyanobacterial host carbon metabolism”

## Awards & Honors

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- National Oceanographic Partnership Program 2024 Excellence in Partnering Award, “Piloting Biological Global Ocean Ship-based Hydrographic Investigations Program (Bio-GO-SHIP)”
- Finalist, Mid-Career Research Scholar Award, Mississippi State University, 2023
- Graduate Teaching Award, Scripps Institution of Oceanography, 2019
- NOAA/OAR Outstanding Scientific Paper Award for Oceans and Great Lakes, 2018
- Ranked First in Selection Process for Visiting Professor in Environmental Science, University of Brasília, 2018 (declined position)
- SABIC Postdoctoral Fellowship, King Abdullah University of Science and Technology, 2012
- SABIC Postdoctoral Fellowship, King Abdullah University of Science and Technology, 2011
- 2011 *PNAS* article selected by Faculty of 1000 as a “must read” article (FFa score of 8)
- Poster Award, Cells, Circuits, and Computation Conference, Harvard University, 2009

- Student Travel Grant, Genomics: GTL Conference, U.S. Department of Energy, 2009
- Praecis Presidential Fellowship, Massachusetts Institute of Technology, 2003
- Howard Hughes Summer Research Fellowship, Stanford University Department of Biological Sciences, 2001
- Semifinalist, Westinghouse Science Talent Search, 1998

## Research Cruises

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- NOAA Gulf of Mexico Trap cruise number 5 (NOAA-GMT-5), “Characterizing the Biodiversity and Variability of the Biological Carbon Pump in the Northern Gulf of Mexico,” R/V *Pelican*, Northern Gulf of Mexico, Nov 29–Dec 1, 2023
- Institute for Marine Research (Norway) cruise number 2019703, “Characterizing mesopelagic fish populations using environmental DNA (eDNA),” R/V *Kronprins Haakon*, Cape Verde to Oslo, May 1–27, 2019
- California Cooperative Oceanic Fisheries Investigations (CalCOFI) 1704SH, “eAUV mobile omics platform to reduce ship time,” NOAA Ship *Bell M. Shimada*, Monterey to San Francisco, Apr 15–20, 2017
- KAUST Red Sea Expedition 2013, “Gene expression of marine bacterial communities in the Eastern Mediterranean and Northern Red Sea,” R/V *Aegaeo*, Athens to Thuwal, Feb 14–Mar 6 and Mar 9–17, 2013
- KAUST Red Sea Expedition 2011, “Population genomics of marine bacteria along physicochemical gradients in the Eastern Red Sea,” R/V *Aegaeo*, Southern to Northern Red Sea, Sep 15–Oct 10, 2011
- Hawaii Ocean Time-series HOT-181, “Cultivation of *Prochlorococcus* from the North Pacific Subtropical Gyre,” R/V *Kilo Moana*, Station ALOHA, May 24–29, 2006

## Chaired Sessions

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- OCB2023 Summer Workshop, Woods Hole Oceanographic Institution, Woods Hole, Massachusetts, June 12–15, 2023, “Sustained observations of global ocean biology”
- ASLO Aquatic Sciences Meeting 2023, Palma de Mallorca, Spain, June 4–9, 2023, “Linking Ocean Microbiomes and Ecosystem Functions”
- TDWG 2017: Biodiversity Information Standards, Ottawa, Canada, October 2017, “Towards robust interoperability in multi-omic approaches to biodiversity monitoring”

## Committee Appointments, Working Groups & Consortia

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- U.S. Marine Biodiversity Observation Network (MBON) eDNA Working Group Leader, 2023–present
- NOAA OAR Omics Supergroup Leader, 2020–present
- National Academy of Sciences Expert Meeting on Biodiversity, New Orleans, LA, June 10–11, 2025
- NOS–OAR Summit Steering Committee, 2021–2022
- Earth HoloGenome Initiative (EHI) Scientific Committee, 2021–2022
- NOAA OAR Cloud Computing Strategy Document Writing Team, 2020
- National Microbiome Data Collaborative (NMDC) Champion, 2020–present
- National Microbiome Data Collaborative (NMDC) Research Coordination Network (RCN) steering committee, 2018–2020
- EMOSE Inter-Comparison of Marine Plankton Metagenomic Analysis Methods, 2017

## Peer Review

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Publons ID: [publons.com/a/1456009/](https://publons.com/a/1456009/)

- Associate Editor, *Proceedings of the Royal Society B*, 2020–2022
- Review Editor, *Frontiers in Marine Science*
- Reviewer for *Nature*, *Proceedings of the Royal Society B*, *Ecology Letters*, *The ISME Journal*, *Genes*, *Bacteriophage*, National Science Foundation, NOAA Office of Ocean Exploration and Research, and United States–Israel Binational Science Foundation

## Protocols

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- Earth Microbiome Project protocols on <https://www.protocols.io/workspaces/earth-microbiome-project> and <https://earthmicrobiome.org/protocols-and-standards/>.
- AOML Omics protocols on <https://github.com/NOAA-AOML/protocols>.

## Websites

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- **L. Thompson**, P. Vangay, K. Blumberg, D. Christianson, J.P. Dundore-Arias, B. Hu, R. Timme, and E. Wood-Charlson. (2020). Introduction to Metadata and Ontologies: Everything You Always Wanted to Know About Metadata and Ontologies (But Were Afraid to Ask). United States. Web. <https://doi.org/10.25979/1607365>.

## GitHub Repositories

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- Earth Microbiome Project: <https://github.com/biocore/emp>
- Tourmaline – Amplicon sequence processing workflow using QIIME 2 and Snakemake: <https://github.com/NOAA-AOML/tourmaline>
- Labelmaker – Printable QR-coded labels for samples: <https://github.com/NOAA-AOML/labelmaker>
- Python for Data Analysis – Online course for data science: <https://github.com/cuttlefishh/python-for-data-analysis>
- Papers – Code for manuscripts published by Thompson and colleagues: <https://github.com/cuttlefishh/papers>

## Teaching Experience

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- Lecturer and Course Developer, “Python for Data Analysis”, Scripps Institution of Oceanography, 2015–18
  - GitHub repository: <https://github.com/cuttlefishh/python-for-data-analysis>
  - YouTube channel: <https://www.youtube.com/channel/UCVZrIrWtcvTzYlRnX7RcDyG>
- Guest Lecturer, Gut Check: Exploring Your Microbiome, Coursera, 2018
- Instructor, Advanced Bioinformatics for Metagenomics and Population Genomics, University of Oulu, Finland, March 2018
- Instructor, SIO Transcriptomics Workshop, Scripps Institution of Oceanography, October 2017
- Instructor, QIIME Workshop, University of Costa Rica, January 2016
- Guest Lecturer, Gut Check: Exploring Your Microbiome, Coursera, 2014
- Co-Supervisor of Ph.D. Student, KAUST, 2011–13
- Mentor of Undergraduate Students, Chisholm Lab, MIT, 2005–10
- Head Teaching Assistant, General Biochemistry, MIT, 2007
- Teaching Assistant, Introductory Biology, MIT, 2005
- Biology Tutor for Undergraduate Students, Boston Area, 2005–08
- Science Explorers Program, Fletcher–Maynard Academy and Cambridge Community Center, 2005–08
- Radio Talk Show Co-Host, *Biologue*, WMBR-FM Cambridge, 2005–08