Sarah L. Lebeis

Michigan State University Department of Plant, Soil, and Microbial Sciences Department of Microbiology and Molecular Genetics Plant Resilience Institute lebeissa@msu.edu

Education:

Emory University, Atlanta, GA, 2003-2008 <u>Doctor of Philosophy</u>, Microbiology and Molecular Genetics, Graduate Division and Biology and Biological Sciences

Michigan State University, East Lansing, MI, 1999-2002

<u>Bachelor of Science</u>, Double major: Lyman Briggs School of Science Residential Program (concentration: Biology) and Microbiology & Molecular Genetics (concentration: Genomics and Molecular Genetics)

Professional Experience:

Assistant Professor: Department of Plant, Soil, and Microbial Sciences, Michigan State University, October 2020 – present

Assistant Professor: Department of Microbiology and Molecular Genetics, Michigan State University, October 2020 - present

Assistant Professor: Plant Resilience Institute, Michigan State University, October 2020 - present

Adjunct Professor: Department of Microbiology, The University of Tennessee, September 2020present

Assistant Professor: Department of Microbiology, The University of Tennessee, February 2014-September 2020

Adjunct Professor: Genome Science and Technology program, The University of Tennessee, March 2014-September 2020

Adjunct Professor: Department of Biology, North Carolina Central University, January 2013 – May 2013

Postdoctoral Research Associate: Department of Biology, The University of North Carolina at Chapel Hill, January 2009 - January 2014, (research advisor: Dr. Jeffery Dangl)

Undergraduate Research/Laboratory Assistant: Department of Microbiology and Molecular Genetics, Michigan State University, September 2001 – July 2003, (research advisor: Dr. Karen Friderici)

Undergraduate Research/Laboratory Assistant: Department of Plant Pathology, Michigan State University, September 2000 – July 2003, (supervisor: Dr. Mary Hausbeck)

Industry Intern: Ancile Pharmaceutical Inc., June 2001 – July 2001, La Jolla, CA

Funding and Research Awards:

2022:

• *Project GREEEN:* Evaluating a Role for the Soybean Root Microbiome in Nutrient Uptake. July 2022-June 2023.

2020:

• USDA-grant: Agricultural Microbiomes in Plant Systems and Natural Resources: Identifying plant genes associated with beneficial foliar yeast in *Populus trichocarpa*. September 2020-August 2024.

2019:

• Science Alliance JDRD: Harvesting plant and bacterial genetic determinants of microbiome structure and function from GWAVA. March 2019 – July 2020

2018:

- *NSF-grant:* CAREER: Defining colonization mechanisms and functions of *Streptomyces* strains in root microbiome. August 2018-July 2023
- *Science Alliance JDRD:* Mining GWAVA for *Streptomyces* interactions in the plant microbiome. March 2018 July 2019

2017:

• Institute for a Secure and Sustainable Environment grant: Nutrient and microbial community implications associated with the addition of duckweed to wastewater remediation.

2016:

- *JGI-Small scale Community Science Project:* Characterizing functional chemotaxis receptors in different root zones. November 2016-October 2018.
- *NSF grant*: Collaborative Proposal: Dimensions of Biodiversity: The evolution of novel interactions within a network of plant, insect, and microbial biodiversity. September 2016-August 2021.

2015:

- *Quest scholar of the week:* Office of Research and Engagement, Distinction to showcase outstanding faculty, staff, and students at the University of Tennessee, September 2015
- *EMSL-JGI-Community Science Project:* Providing sequencing and facility services for "Uncovering the composition and function of the aquatic microbiome for duckweeds", October 2014-December 2016.

Prior to 2014:

• *Postdoctoral fellowship*: SPIRE (Seeding Postdoctoral Innovators in Research and Education) postdoctoral training program at The University of North Carolina, Chapel Hill, funded by the NIGMS, 2009-2012.

Publications: (* designates graduate student, ** designates undergraduate student, ^ designates equal contribution, # designates corresponding author)

- O'Banion, B.S.*, Kelley, B.R., and <u>Lebeis, S. L.</u>[#] Arabidopsis RBOHD differentially impacts microbially-induced root phenotypes and tissue-specific competitive colonization patterns (In preparation)
- O'Banion, B.S.*, Jones, P.*, Demetros, A.A.*, Kelley, B.R., Wagner, A.S.*, Chen, J.G., Michero, W., Reynolds, T.B., Jacobson, D., and <u>Lebeis, S.L.</u>[#] Plant inositol transport influences bacterial colonization phenotypes (Under Review)
- Gates, A.D.*, French, A.M.**, Demetros, A.A.*, Kelley, B.R., and <u>Lebeis, S.L.</u>[#] A *Streptomyces* consortium contributes distinct microbial interactions during *Arabidopsis thaliana* microbiome assembly (Under Review)
- 4) Moccia, K. and <u>Lebeis, S.L.</u>[#] *Pantoea* spp. genomic landscapes reveal genetic factors important for interactions with plants (Under Review)
- 5) Acosta, K.*, Sorrels, S., Chrisler, W., Huang, W., Gilbert, S., Brinkman, T., Michael, T.P., <u>Lebeis, S.L.</u>, and Lam, E. Optimization of molecular methods for detection and quantification of specific duckweed-bacteria interactions. (Accepted)
- 6) Shastry, V.*, Bell, K.L., Buerkle, C.A., Fordyce, J.A., Forister, M.L., Gompert, Z., <u>Lebeis</u>, <u>S.L.</u>, Lucas, L.K., Marion, Z.H., and Nice, C.C. (2022) A continental-scale survey of *Wolbachia* infections in blue butterflies reveals evidence of interspecific transfer and invasion dynamics. *G3.* 12(10): jkac213.
- 7) Gompert, Z., Dodson, C., <u>Lebeis, S</u>., Fordyce, J., Lucas, L., Buerkle, A., Forister, M., Saley, T.*, Philbin, C., Yoon, S., Perry, E., Sneck, M., and Harrison, J. (2022) Additive genetic effects in interacting species jointly determine the outcome of caterpillar herbivory. *PNAS.* 119(36): e2206052119.
- 8) Beals, K.K.*, <u>Lebeis, S.L.</u>, Bailey, J.K., and Schweitzer, J.A. (2022) Conditionality of soil microbial mediation of Solidago plant phenotype: indicator taxa within complex microbiomes influence some, but not all Solidago traits. *Plant and Soil.* 1-18
- Gilbert, S., Poulev, A., Chrisler, W., Acosta, K.*, Scott, K., Orr, G., <u>Lebeis, S.L.</u>, and Lam, E. (2022) Auxin-producing bacteria from duckweeds have different colonization patterns and effects on plant morphology. *Plants* 11(6): 721.
- 10) Forister, M.L., Philbin, C.S., Marion, Z.H., Buerkle, C.A., Dodson, C.D., Fordyce, J.A., Forister, G.W., <u>Lebeis, S.L.</u>, Lucas, L.K, Nice, C.C., and Gompert, Z. (2020) Leveraging biological complexity to predict patch occupancy in a recent host range expansion. *Science Advances*. 6(48): eabc6852
- 11) Dickey, J.*, Fordyce, J., and <u>Lebeis, S.L.</u>[#] (2020) Bacterial communities of the rhizosphere explained by spatial structure and sampling grain. *Molecular Ecology*. 80(4): 846-858.
- 12) Moccia, K.*, Papoulis, S.*, Willems, A.**, Fordyce, J.A., and <u>Lebeis, S.L.</u> (2020) Using 18S rRNA amplicon sequencing to explore the eukaryotic members of the *Medicago sativa* microiome. *Phytobiomes*. PBIOMES-02-20-0022R
- 13) Huang, W., Gilbert, S., Poulev, A., Acosta, K., <u>Lebeis, S.,</u> Long, C., and Lam, E. (2020) Host-specific and tissue-dependent orchestration of microbiome community structure in traditional rice paddy ecosystem. *Plant & Soil.* 452: 379-395.

- 14) Liu, F.*, Rice, J., Lopes, V., Grewal, P., <u>Lebeis, S.L.,</u> Hewezi, T., and Staton, M. (2020)
 Overexpression of strigolactone-associated genes exerts fine-tuning selection on soybean rhizosphere bacterial and fungal microbiomes. *Phytobiomes.* 4(3): 239-251.
- 15) Acosta, K., Xu, J., Gilbert, S., Brinkman, T., Denison, E.**, <u>Lebeis, S.L</u>. and Lam, E. (2020) Duckweed hosts a taxonomically similar bacterial assemblage as the terrestrial leaf microbiome. *PLoS One.* 15(2): e0228560.
- 16) <u>Lebeis, S.L.</u>[#] (2020) mSphere of Influence: Peering through a keyhole into the unseen world. *mSphere*. 5(1): e00980-19. (Invited)
- 17) Moccia, K.*, Willems, A.**, Papoulis, S.*, Flores, A., Forister, M.L., Fordyce, J.A., and <u>Lebeis, S.L.</u>[#] (2020) Distinguishing Nutrient-dependent plant driven colonization patterns in alfalfa. *Environmental Microbiology Reports*. 12(1): 70-77.
- 18) O'Banion, B.S.*, O'Neal, L.*, Alexandre, G., and <u>Lebeis, S.L.</u> (2020) Bridging the gap between single-strain and community-level plant-microbe chemical interactions. *Molecular Plant-Microbe Interactions* 33(2): 124-134. (Invited)
- 19) Moccia, K.* and <u>Lebeis, S.L.*</u>(2019) Microbial Ecology: How to Fight the Establishment. *Current Biology*. 29(24): R1320-R1323. (Invited)
- 20) Li, Z., Yao, Q., Guo, X., Crits-Christoph, A., Mayes, M.A., Hervey, W.J., Lebeis, S.L., Banfield, J.F., Hurst, G.B., Hettich, R.L., and Pan, C. (2019) Genome-resolved proteomic stable isotope probing of soil microbial communities using 13CO₂ AND 13C-methanol. *Frontiers in Microbiology*. 10: 2706.
- 21) Liu, F.*, Hewezi, T., <u>Lebeis, S.L.</u>, Pantalone, V., Grewal, P.S., and Staton, M.E. (2019) Soil indigenous microbiome and plant genotypes cooperatively modify soybean rhizosphere microbiome assembly. *BMC microbiology*. 19(1): 1-19.
- 22) Chewning, S.S.^{*}, Grant, D.L.^{*}, O'Banion, B.S.^{*}, Kennedy, B.J.^{*}, Campagna, S.R., and <u>Lebeis, S.L.[#]</u> (2019) Root-associated *Streptomyces* harboring *mel* genes demonstrate enhanced plant colonization. *Phytobiomes Journal*. 3(3): 165-176.
- 23) Gilbert, S., Xu, J., Acosta, K., Poulev, A., <u>Lebeis, S.L.,</u> and Lam, E. (2018) Bacterial production of indole related compounds reveal their role in association between duckweeds and endophytes. *Frontiers in Chemistry.* 6: 265. doi: 10.3389/fchem.2018.00265.
- 24) Levy, A, Clingenpeel, S., Gonzalez, I., Herrera Paredes, S., Stillman, K., Monteiro, F., Alvares, B., Lundberg, D., Lu, T.-Y., <u>Lebeis, S.L</u>., Jin, Z., McDonald, M., Feltcher, M., del Rio, T., Grant, S., Doty, S., Ley, R., Pelletier, D., Tringe, S., and Woyke, T. (2017) Genetic determinants of bacterial adaptation to plants. *Nature Genetics.* 50: 138-150. doi: 10.1038/s41588-017-0012-9.
- 25) <u>Lebeis, S.L.</u> and Robatzek, S. (2017) Editorial overview: Biotic interactions: Inferring global implications for the molecular interface between plants and their biotic interactions across scales. *Current Opinion in Plant Biology: Biotic Interactions*. 38: v-vii. doi: 10.1016/j.pbi.2017.06001. (Invited)
- 26) <u>Lebeis, S.L.</u>[#] (2017) Plant Microbiome Identification and Characterization. *Current Protocols in Plant Biology*. 2: 135-146. doi: 10.1002/cppb/20048. (Invited)
- 27) Herrera Paredes, S and <u>Lebeis, S. L.[#]</u> (2016) Giving back to the community: Microbial Mechanisms of plant-soil interactions. *Functional Ecology*. 30: 1043-1052. doi: 10.1111/1365-2435.12684. (Invited)

- 28) <u>Lebeis, S.L.</u>^{*}, Herrera Paredes, S.[^], Lundberg, D.S.[^], Breakfield, N., Gehring, J., McDonald, M., Malfatti, S., Glavina del Rio, T., Jones, C.D., Tringe, S.G., and Dangl, J.L. (2015) A plant defense hormone modulates colonization of the root microbiome by specific bacterial taxa. *Science.* 349: 860-864. PMID 26184915. doi: 10.1126/science.aaa8764.
- 29) Sloan, S.S.* and <u>Lebeis, S.L.</u>[#] (2015) Exercising influence: distinct biotic interactions shape root microbiomes. *Current Opinions in Plant Biology*. 26: 32-36. doi: 10.1016/j.pbi.2015.05.026.
- 30) Macquard, S., Garrido-Oter, R., Gonzalez, A., Spaepen, S., Ackermann, G., <u>Lebeis, S.</u>, McHardy, A.C., Dangl, J.L., Knight, R., Ley, R., and Schulze-Lefert, P. (2015) Microbiota and Host Nutrition across Plant and Animal Kingdoms. *Cell Host and Microbe*. 17(5): 603-616. doi: 10.1016/j.chom.2015.04.009.
- 31) <u>Lebeis, S.L.</u>[#] (2014) Greater than the sum of their parts: characterizing root microbiomes on the community-level. *Current Opinions in Plant Biology*. 24: 82-86. doi: 10.1016/j.pbi.2015.02.004.
- 32) <u>Lebeis, S.L.</u>[#] (2014) The potential for give and take in the plant microbiome. *Frontiers in Plant Science*. 5: 287. doi: 10.3389/fpls/2014.00287. (Invited)
- 33) Lebeis, S.L., Rott, M., Dangl, J.D., and Schulze-Lefert, P. (2012) Culturing a plant microbiome community at the cross-Rhodes. *New Phytologist*. 196(2): 341-344. doi: 10.1111/j.1469-8137.2012.04336.x.
- 34) Lundberg, D.S.[^], <u>Lebeis, S.L.[^]</u>, Herrera Paredes, S.[^], Yourstone, S.[^], Gehring, J., Malfatti, S., Tremblay, J., Engelbrekston, A., Kunin, V., Glavina del Rio, R., Edgar, R.C., Eickhorst, T., Ley, R.E., Hugenholtz, P., Tringe, S., and Dangl, J. (2012) Defining the core *Arabidopsis thaliana* root microbiome. *Nature*. 488: 86-90. doi: 10.1038/nature11237. (Cover story)
- 35) <u>Lebeis, S.</u> and Kalman, D. (2009) Aligning antimicrobial drug discovery with complex and redundant host-pathogen interactions. *Cell Host and Microbe*. 5(2): 114-122. doi: 10.1016/j.chom.2009.01.008.
- 36) <u>Lebeis, S.L.</u>, Powell, K.R., Merlin, D., Sherman, M.A., and Kalman, D. (2009) IL-1 receptor signaling protects mice from lethal intestinal damage caused by the attaching and effacing pathogen *C. rodentium. Infection and Immunity*. 77(2): 604-614. doi: 10.1128/IAI.00907-08.
- 37) <u>Lebeis, S.L.</u>, Sherman, M.A., and Kalman, D. (2008) Protective and destructive effects of the innate immune response to enteropathogenic *Escherichia coli* and related A/E pathogens. *Future Microbiology*. 3: 315-328. doi: 10.2217/17460913.3.3.315.
- 38) <u>Lebeis, S.L.</u>, Bommarius, B., Parkos, C.A., Sherman, M.A., and Kalman, D. (2007) TLR signaling mediated by MyD88 is required for a protective innate immune response by neutrophils to *Citrobacter rodentium*. *Journal of Immunology*. 179(1): 566-577. PMID: 17579078.
- 39) Reeves, P.M., Bommarius, B., <u>Lebeis, S.</u>, McNulty, S., Christensen, J., Swimm, A., Chahroudi, A., Chavan, R., Feinberg, M.B., Veach, D., Bornmann, W., Sherman, M., and Kalman, D. (2005) Disabling poxvirus pathogenesis by inhibition of Abl-family tyrosine kinases. *Nature Medicine*. 11(7): 731-739. PMID: 15980865.

40) Rothrock, C.R., Murgia, A., Sartorato, E.L., Leonardi, E., Wei, S., <u>Lebeis, S.L.</u>, Yu, L.E., Elfenbein, J.L., Fisher, R.A., and Friderici, K.H. (2003) Connexin 26 35delG does not represent a mutational hotspot. *Human Genetics*. 113(1): 18-23. PMID: 1268473.

Invited Talks and Seminars:

| 2022 | November 7 th | - Division of Biology seminar, Kansas State University, Manhattan, Kansas | |
|------|--|---|--|
| | September 30 th – Kellogg Biological Station seminar, Hickory Corners, Michigan | | |
| | September 16 | th – Plant Sciences Society Meeting, National Institute for Biology, Ljubljana, | |
| | | Slovenia(Online) | |
| | July 9 th | ASPB, Annual Plant Biology Meeting, Portland, Oregon | |
| | May 16 th | - Great Lakes Bioenergy Research Center, Lake Geneva, Wisconsin | |
| | February 1 st | Society for Industrial Microbiology and Biotechnology, San Diego | |
| | | (Canceled) | |
| | January 14 th | Plant Nutrition and Development Symposium, MSU (Online) | |
| 2021 | December 14 ^t | ^h - John Lawrence Seminar, Lawrence Berkeley National Laboratory (Online) | |
| 2020 | November 16 ^t | ^h – Interdisciplinary Plant Group seminar, University of Missouri (Online) | |
| | October 27 th | - Plant Pathology Department seminar, University of Wisconsin (Online) | |
| | October 1 st | - Plant and Microbial Biology Department seminar, University of Zurich | |
| | | (Online) (International) | |
| | July 28 th | - ASM Micro 2020 (Online) | |
| 2019 | July 8 th | FEMS Microbiology, Glasgow, Scotland (International) | |
| | April 17 th | - Plant Microbiome Workshop, Banbury Center, Cold Spring Harbor, NY | |
| | April 8 th | - Biology Department seminar, Vanderbilt University, Nashville, TN | |
| 2018 | December 17 ^t | ^h - Evolutionary Biology group seminar, University of Kiel, Kiel, Germany | |
| | | (International) | |
| | November 8 th | - Bacteriology Department, University of Wisconsin, Madison, WI | |
| | October 3 rd | - Biochemistry and Cellular and Molecular Biology Department seminar, | |
| | | University of Tennessee, Knoxville, TN | |
| | June 21 st | 21st Penn State Plant Biology Symposium: Wild and Tamed | |
| | | Phytobiomes, University Park, PA | |
| | March 29 th | - Plant Sciences Department seminar, University of Arizona, Tucson, AZ | |
| 2017 | December 4 th | Microbiology and Molecular Genetics Department seminar, Emory University, Atlanta, GA | |
| | June 27 th | - American Society of Plant Biologists annual conference, Honolulu, HI | |
| | January 8 th | - Microbiology Department seminar, University of Georgia, Athens, GA | |
| 2016 | September 10 | th - The 6 th ASM conference on Beneficial Microbes, Seattle, WA | |
| | August 24 th | The International Society for Microbial Ecology, Montreal, Canada | |
| | | (International) | |
| | July 19 th | The International Society for Molecular Plant and Microbe Interactions conference, Portland, OR (International) | |
| | May 10 th | - New Model Systems for Linking Evolution and Ecology. EMBL | |
| | - / - | Heidelberg, Germany (International) | |
| | March 30 th | - MicroSeminar, Web-based Microbiology Seminar Series | |
| | | | |

| 2015 | March 23 rd January 31 st November 6 th October 26 th | (online) (microseminar.wordpress.com) Joint Genome Institute's Annual User's meeting, Walnut Creek, CA Donald Danforth Center seminar, St. Louis, MO Root Biology Workshop, Noble Foundation, Ardmore, OK Plant Biology Department seminar, Michigan State University, East Lansing, MI |
|------|--|---|
| | August 3 rd | - American Phytopathology Society. Pasadena, CA |
| | June 30 th | - Phytobiomes 2015: Designing a New Paradigm for Crop Improvement conference. Washington, D.C. |
| | March 18 th | Department of Horticulture seminar, University of Kentucky, Lexington, KY |
| 2014 | November 12 th | - Plant-Microbe Interaction group seminar, Oak Ridge National Lab, Oak Ridge, TN |
| | September 8 th | - Entomology and Plant Pathology Department seminar, University of Tennessee, Institute of Agriculture, Knoxville, TN |
| | August 28 th | - Plant Research Center colloquium. Knoxville, TN |
| | July 8 th | - The International Society for Molecular Plant and Microbe Interactions conference, Rhodes, Greece (International) |
| | January 10 th | - USDA/DOE Plant Feedstocks Genomics PI/PD meeting, San Diego, CA |
| 2013 | December 4 th | Plant Genomes and Biotechnology: From Genes to Networks, Cold Spring Harbor, NY |
| 2012 | November 29 th | Evolution and Development seminar series, Duke University, Durham, NC |
| | November 6 th · | - Plant Biology Department seminar, North Carolina State University, Raleigh, NC |

Student Presentations:

(* indicates graduate student, ** indicates undergraduate student)

2022 - O'Banion, B.S.* and <u>Lebeis, S.L.</u> Plant and bacterial inositol exchange influences root colonization outcomes. International Society for Microbial Ecology, Lausanne, Switzerland (**Oral presentation**)

- O'Banion, B.S.* and <u>Lebeis, S.L.</u> Plant and bacterial inositol exchange influences root colonization outcomes. International Phytobiomes Conference, Denver, CO (**Oral presentation**)

Gates, A.*, Hawkins, A.**, Demetros, A.A.*, Kelley, B., and Lebeis, S.L.
Untangling salicylic acid driven microbial ineractions during root microbiome assembly. American Society for Plant Biology, Portland, OR (Poster presentation)
Hawkins, A.**, Gates, A.*, and Lebeis, S.L. Salicylic acid influence on Arabidopsis thaliana root microbiomes. University Undergraduate Research and Arts Forum, Michigan State University, East Lansing, MI (Poster presentation)

 DeClaire, M.**, Kelley, B.R., and <u>Lebeis, S.L.</u> Influence of biological products on soybean drought performance. University Undergraduate Research and Arts Forum, Michigan State University, East Lansing, MI (Poster presentation) 2021 - Gates, A.* and <u>Lebeis, S.L.</u> Generation of enriched bacterial communities to define salicylic acid influence in the root endosphere. IS-MPMI Congress: eSymposia Series, Plant-microbe interactions in the environment. (Online) (Oral presentation)

- O'Banion, B.* and <u>Lebeis, S.L.</u> Plant and bacterial inositol exchange influences root colonization outcomes. IS-MPMI Congress: eSymposia Series, Plant-microbe interactions in the environment. **(Online)** (**Oral presentation**)

- 2019 Hyde, T.** and Lebeis, S.L. 2019. Plant-associated bacteria compete differently on diverse media sources. Poster presented at the Office of Research Undergraduate Research Symposium, University of Tennessee, Knoxville, TN (Poster presentation)
 - O'Banion, B.*, O'Neal, L.*, Alexandre, G., and <u>Lebeis, S.L.</u> Spatially discrete micro-niches govern root microbiome assembly. Molecular Plant-Microbe Interaction International Symposium, Glasgow, Scotland. (**Oral presentation**)
 - O'Banion, B.*, Hyde, T.,**, O'Neal, L.*, Alexandre, G., and <u>Lebeis, S.L.</u> Spatially discrete micro-niches govern root microbiome assembly. FEMS Microbiology Symposium, Glasgow, Scotland. (Oral presentation)
 - Moccia, K.*, Willems, A.*, Papoulis, S.*, Fordyce, J., and <u>Lebeis, S.L.</u>
 Distinguishing nutrient-dependent plant driven bacterial colonization patterns in alfalfa. Molecular Plant-Microbe Interactions International Symposium, Glasgow, Scotland (Poster presentation)
 - O'Banion, B.,*, Jones, P.*, Grant, D.*, Chewning, S.S.*, Jacobson, D., and <u>Lebeis</u>,
 <u>S.L.</u> Mining GWAVA for key factors shaping microbiome structure. Science Alliance JDRD Symposium, Knoxville, TN (Poster presentation)
- 2018 O'Banion, B.*, O'Neal, L.*, Alexandre, G., and Lebeis, S.L. 2018. Assessing impact of spatially distinct chemotaxis on overall root-microbiome assembly. The KY and TN Regional ASM Meeting, Lexington, KY. (Poster presentation)
 Moccia, K.*, Willems, A.**, & Lebeis, S. From sporadic to consistent: Colonization of microbes can depend on nutrient availability. International Society for Microbial Ecology, Leipzig, Germany. (Poster presentation)
 Chewning, S. S.*, Grant, D.**, O'Banion, B.*, & Lebeis, S. Streptomyces employ melanin production to compete and succeed in the root microbiome. Joint Genome Institute Annual User's meeting, Walnut Creek, CA. (Poster presentation)
- 2017 Sloan, S. S.*, Grant, D.**, Whitley, C.**, &. Lebeis, S. L. Unique metabolic potential of root-associated Streptomyces strains predicts their induced plant phenotypes. Microbial and Plant Systems Modulated by Secondary Metabolites Meeting, July 24-26, Walnut Creek, CA. (Poster presentation)

- Grant, D.**, Sloan, S. S.*, & <u>Lebeis, S. L.</u> Identification of Streptomyces isolatespecific secondary metabolites capable of influencing Arabidopsis thaliana root microbiome establishment. EUReCA event, April 25, Knoxville, TN. (Poster presentation)

2016 - Sloan, S. S.*, Massey, J.**, Kackos, C.**, & Lebeis, S. L. Streptomyces sculpt the

root microbiome of Arabidopsis thaliana. 16th International Symposium on Microbial Ecology, August 21-26, Montreal, Quebec, Canada. (**Poster presentation**)

- McGuire, K. G.**, Basso, J.*, Harvey, M. L.**, & <u>Lebeis, S. L.</u> Duckweed Dynasty: Characterization of microbes associated with duckweed. EUReCA event, April 26, Knoxville, TN. (**Poster presentation**)

2015 - Sloan, S. S.*, & <u>Lebeis, S. L.</u> Soil-resident Streptomycetaceae influence phenotypic characteristics of Arabidopsis thaliana. 27th Annual Plant Molecular Biology Consortium, September 18-20, Asheville, NC. (Poster presentation)

Teaching experience:

| Academic Session | Course Title | # Students |
|------------------|------------------------------------|------------|
| Spring 2023 | PLB 865: Plant Omics | 10 |
| Spring 2023 | MMG 499: Undergraduate Research | 1 |
| Spring 2023 | CSS460: Plant-Microbe Interactions | 14 |
| Spring 2022 | PLB 865: Plant Omics | 10 |
| Spring 2022 | MMG 499: Undergraduate Research | 1 |
| Spring 2022 | CSS491: Plant-Microbe Interactions | 14 |
| Fall 2021 | MMG 499: Undergraduate Research | 1 |

Courses taught at Michigan State University:

Courses taught at the University of Tennessee:

| Academic Session | Course Title | # Students |
|------------------|---|------------|
| Spring 2020 | MICR 606: Microbial Ecology Journal Club | 12 |
| Fall 2019 | MICR 330: Immunology | 120 |
| Spring 2019 | MICR 431: Advanced Immunology | 23 |
| Fall 2018 | BIOL 220: General Microbiology | 91 |
| Spring 2018 | MICR 650: Molecular Biology Topics | 7 |
| Fall 2017 | MICR 330: Immunology | 137 |
| Spring 2017 | MICR 431: Advanced Immunology | 22 |
| Fall 2016 | BIOL 220: General Microbiology | 101 |
| Spring 2016 | MICR 630: Immunology Topics | 7 |

| Fall 2015 | MICR 330: Immunology | 115 |
|-------------|--------------------------------|-----|
| Spring 2015 | MICR 310: General Microbiology | 83 |

Courses taught at North Carolina Central University:

| Academic Session | Course Title | # Students |
|------------------|------------------------------|------------|
| Spring 2013 | Introduction to Microbiology | 25 |
| Fall 2012 | Host-Microbe Relationships | 17 |
| Spring 2012 | Introduction to Microbiology | 25 |

Teaching Awards:

2019 College of Arts and Sciences, University of Tennessee: Undergraduate Teaching

- 2018 University of Tennessee, University of Tennessee: Chancellor's Excellence in Teaching
- 2017 Department of Microbiology, University of Tennessee: Outstanding Undergraduate Teaching

Academic Mentoring and Service:

Current MSU PhD Student:

Leah Knoor – Connecting plant root exudate components with root colonization outcomes

Expected graduation: Spring 2027

Imani Pascoe (co-mentored with Greg Bonito) – Defining metabolic exchange between switchgrass its fungal and bacterial microbiome members Expected graduation: Spring 2026

Kevin Santiago-Morales – Identification and characterization of phosphate nutrient status during plant microbiome assembly Expected graduation: Spring 2025

Moss Le – Influence of nutrition and foliar yeasts on plant disease resilience Expected graduation: Spring 2026

Current UTK PhD Students:

Bridget O'Banion – Defining the influence of chemotaxis to specific root niches in microbiome assembly and functioning Expected graduation: Spring 2023

Alexandra Gates – Distinguishing a role for *Streptomyces* strains during plant root microbiome assembly

Expected graduation: Fall 2023

Former UTK MS Student: David Grant – Investigating Streptomyces genes and secondary metabolites involved in plant root colonization Current position: Project Coordinator, Assured Bio Labs

Former PhD Student:

Katherine Moccia, PhD – More than the sum of their parts: Building a framework for understanding host-microbe interactions in *Medicago sativa* Graduation: Fall 2020 Current position: Assistant Professor, Biology Department, Wagner College

Sarah Stuart Chewning, PhD. – Microbial partners in health: Broadening our understanding of microbiome-host relationships Graduation: Fall 2018 Current position: Technical Sales Specialist, Thermo Fisher Scientific

Current graduate committees: 6 PhD students (4 at MSU and 2 at UTK)

University Service Award:

- 2017 College of Arts and Sciences: Undergraduate Academic Advising
- 2017 *College of Arts and Sciences:* Undergraduate Research Mentor of the Year, Natural Sciences division

Professional society memberships and contributions:

American Society for Microbiology (ASM) and American Phytopathological Society (APS)

Academic Service:

Co-editor: Current Opinions in Plant Biology: Biotic Interactions, volume 38, August 2017

Editorial board: Applied Environmental Microbiology

Reviewed manuscripts for: Applied Environmental Microbiology, Cell Host and Microbe, eLife, Frontiers in Microbiology, Frontiers in Plant Science, ISME Journal, Microbial Ecology, Molecular Plant-Microbe Interactions, Molecular Plant Pathology, New Phytologist, Plant Molecular Biology, PLoS One, Proceedings of the National Academy of Sciences USA, and Vaccine

Ad hoc grant reviewer and panelist for: Department of Energy, Joint Genome Institute, National Science and Engineering Research Council of Canada, National Science Foundation, Swiss National Science Foundation, and U.S. Department of Agriculture.

Conference session chair:

- 2017 American Society for Plant Biologists, Honolulu, HI
- 2016 The International Society for Molecular Plant and Microbe Interactions conference, Portland, OR
- 2016 New Model Systems for Linking Evolution and Ecology, EMBL, Heidelberg, Germany

Community outreach:

| October 2022 | EcoTek student mentor: Lab hosts EcoTek high school students |
|---------------|---|
| June 2019 | Designer/Instructor: Living soil, K-12 teacher workshop, Oak Ridge, TN |
| March 2017 | <i>Proctor:</i> Science Olympiad State Tournament competition, Science Olympiad, Knoxville, TN |
| February 2016 | <i>Judge:</i> Tennessee Junior Science and Humanities Symposium, Tennessee Junior Science and Humanities Symposium, Knoxville, TN. |

Popular press/Interviews:

Fang, Janet. "Plant Defense Hormones Build Healthy Microbiomes for Roots" <u>https://www.iflscience.com/plants-and-animals/plant-defense-hormones-build-healthy-microbiome-roots/</u>

(July 16, 2015)

Lawrence Berkeley National Laboratory. "Unearthing cornerstones in root microbiomes". ScienceDaily. www.sciencedaily.com/releases/2015/07/150716141939.htm

Manke, Kara. "Underground Immunity". The Scientist <u>http://www.the-scientist.com/?articles.view/articleNo/43530/title/Underground-Immunity/</u> (July 16, 2015)

Howard Hughes Medical Institute. "Defenses Up: Hormone Helps Plants Determine Friend from Foe"

http://www.sciencenewsline.com/articles/2015071714540027.html (July 16, 2015)

Ledford, Heidi. "Plant dwellers take the limelight: Researchers seek holistic view of botanic ecosystems". *Nature*. 523: 137-138. <u>http://www.nature.com/news/plant-denizens-get-the-big-science-treatment-1.17920</u> (July 9, 2015)