

Myoung-Hwan Chi, Ph. D.

Research Lab Facility Scientist
Noble Research Institute, LLC.
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EDUCATION and TRAINING

<u>Institution</u>	<u>Major</u>	<u>Degree</u>	<u>Year</u>
Seoul National University, Korea	Plant Microbiology	Ph. D.	2009
Seoul National University, Korea	Agricultural Biology	B.S.	2002

PROFESSIONAL EXPERIENCE

<u>Institution</u>	<u>Job title</u>	<u>Year</u>
Noble Research Institute, LLC	Research Lab Facility Scientist	2021.07 - Present

Responsible for scientific and administrative supports for lab members and collaborators

- Managing a group collaboration for EPSCoR project in Red River Farm and Coffey Ranch (2021-)

<u>Institution</u>	<u>Job title</u>	<u>Year</u>
Noble Research Institute, LLC	Senior Research Associate	2011.10 – 2021.07
<ul style="list-style-type: none">- Managed an internal group collaboration for ‘microbial population and grazing animal impact under no-till and cover crop’ project in Pasture Demonstration Facility (2016-2017)- Managed an external group collaboration for ‘switchgrass sustainability project’ in Red River Farm and HQ3 (2017-2021)- Managed an external group collaboration for ‘deployment and metagenomics analysis of beneficial switchgrass microbes in a field environment’ in Red River Farm and HQ3 (2018-2021)- Managed multi-years of field applications on forage species by using beneficial microbes isolated from tall grass prairie in Oklahoma (2009-2021)- Led and guided new team members, visiting scientists, summer scholars, and SOTC interns		

<u>Institution</u>	<u>Job title</u>	<u>Year</u>
Noble Research Institute, LLC	Postdoc Fellow	2009.09 – 2011.09
Responsible for performing scientific research involving diverse microbial species.		
<ul style="list-style-type: none">- Identified new hypothesis on dissemination mechanism of model and human pathogenic		

fungi

- Involved in isolation and identification of root endophytes and generated a collection of 1500 fungal isolates and 550 bacterial isolates.

<u>Institution</u>	<u>Job title</u>	<u>Year</u>
Seoul National University, Korea	Graduate Student	2002.03 – 2009.08
<ul style="list-style-type: none"> - Created a large insertional mutant library of rice blast pathogen - Involved in development of database cataloging phenotypes/genotypes of the library and a portal site providing tools for various analyses and data curation. - Contributed to writing grant proposals - Led an interdisciplinary project (Biogreen21), managed budget and orchestrated collaborations 		

CERTIFICATIONS

National certificate for plant protection specialist (2002-)

Certificate of completion for Microbiome Bioinformatics with QIIME2 (2021-)

PUBLICATIONS

Wang, N. R., Melnyk, R. A., Wiesmann, C. L., Hossain, S. S., Chi, M. H., Martens, K. & Haney C. H. 2021. Commensal *Pseudomonas fluorescens* protect Arabidopsis from closely-related *Pseudomonas* pathogens in a colonization-dependent manner. mBio. (Accepted)

Ray, P., Guo, Y., Chi, M. H., Krom, N., Boschiero, C., Watson, B., Huhman, D., Zhao, P., Singan, V. R., Lindquist, E. A., Yan, J., Adam, C., & Craven, K. D. 2021. *Serendipita* fungi modulate the switchgrass root transcriptome to circumvent host defenses and establish a symbiotic relationship. MPMI. Doi: 10.1094/MPMI-04-21-0084-R

Ray, P., Guo, Y., Chi, M. H., Krom, N., Saha, M. C. & Craven, K. D. 2020. *Serendipita bescii* promotes winter wheat growth and modulates the host root transcriptome under phosphorus and nitrogen starvation. Environmental Microbiology, doi: 10.1111/1462-2920.15242.

Ray, P., Chi, M. H., Guo, Y., Chen, C., Adam, C., Kuo, A., LaButti, K., Lipzen, A., Barry, K. W., Grigoriev, I. V., Tang, Y. & Craven, K. D. 2018. Genome sequence of the plant growth-promoting fungus *Serendipita vermicifera* subsp. *bescii*: The first native strain from North America. Phytobiomes Journal, 2 (2), 62-63 https://doi.org/10.1094/PBIOMES-04-17-0017-A.

Chi, M. H., and Craven, K. D. 2016. RacA-mediated ROS signaling is required for polarized cell differentiation in conidiogenesis of *Aspergillus fumigatus*. PloS one, 11(2), e0149548.

Chi, M. H. and Craven, K. D. 2013. Oxygen and an extracellular phase transition independently control central regulatory genes and conidiogenesis in *Aspergillus fumigatus*. PLoS ONE. 8: e74805.

Park, S. Y., Choi, J., Lim, S. E., Lee, G. W., Park, J., Kim, Y., Kong, S., Kim, S. Rho, H. S., Jeon, J. **Chi, M. H.**, Kim, S., Khang, C. H., Kang, S. and Lee Y. H. 2013. Global expression profiling of transcription factor genes provides new insights into pathogenicity and stress responses in the rice blast fungus. PLoS Pathogens. 9: e1003350.

Park, S. Y., **Chi, M. H.**, Milgroom, G. M., Kim, H., Han, S. S., Kang, S and Lee, Y. H. 2010. Genetic stability of *Magnaporthe oryzae* during successive passages through rice plants and on artificial medium. Plant Pathol. J. 26(4): 313-320

Chi, M. H., Park, S. Y., Kim, S. and Lee, Y. H. 2009. A novel pathogenicity gene is required in the rice blast fungus to suppress the basal defenses of the host. PLoS Pathogens. 5: e1000401

Chi, M. H., Park, S. Y., Kim, S., Lee, Y. H. 2009. A quick and safe method for fungal DNA extraction. Plant Pathol. J. 25: 108-111.

Yi, M., **Chi, M. H.**, Khang, C. H., Park, S. Y., Kang, S. Valent, B., and Lee, Y. H. 2009. The ER chaperone LHS1 is involved in asexual development and rice infection by the blast fungus *Magnaporthe oryzae*. Plant Cell. 21: 681-695.

Park, S.Y., Jwa, N. S., **Chi, M. H.**, Lee, Y. H. 2009. A fluorescence-based cDNA-AFLP method for identification of differentially expressed genes. Plant Pathol. J. 25: 184-188.

Kim, S., Park, S. Y., Kim, K. S., Rho, H.S., **Chi, M. H.** Choi, J., Park, J., Kong, S., Park, J., Goh, J., and Lee, Y. H. 2009. Homeobox transcription factors are required for conidiation and appressorium development in the rice blast fungus *Magnaporthe oryzae*. PLoS Genetics 5: e1000757.

Jeon, J. Goh, J. Yoo, S. **Chi, M. H.** Choi, J. Rho, H. S. Park, J. Han, S. S. Kim, B. R. Park, S. Y. Kim, S. and Lee, Y. H. 2008. A putative MAP kinase kinase kinase, MCK1, is required for cell wall integrity and pathogenicity of the rice blast fungus, *Magnaporthe oryzae*. Mol. Plant Microbe Interact. 21: 525-534.

Choi, J. Park, J. Jeon, J. **Chi, M. H.** Goh, J. Yoo, S. Y. Park, J. Jung, K. Kim, H. Park, S. Y. Rho, H. S. Kim, S. Kim, B. R. Han, S. S. Kang, S. and Lee, Y. H. 2007. Genome-wide analysis of T-

DNA integration into the chromosomes of *Magnaporthe oryzae*. Mol. Microbiol. 66: 371-382.

Jeon, J. Park, S. Y. **Chi, M. H.** Choi, J. Park, J. Rho, H. S. Kim, S. Goh, J. Yoo, S. Choi, J. Park, J. Y. Yi, M. Yang, S. Kwon, M. J. Han, S. S. Kim, B. R. Khang, C. H. Park, B. Lim, S. E. Jung, K. Kong, S. Karunakaran, M. Oh, H. S. Kim, H. Kim, S. Park, J. Kang, S. Choi, W. B. Kang, S. and Lee, Y. H. 2007. Genome-wide functional analysis of pathogenicity genes in the rice blast fungus. Nat. Genet. 39: 561-565.

SCIENTIFIC COMMUNITY SERVICES

- 2013-Present Manuscript reviewer for The Plant Pathology Journal, PLOS ONE, Research in Plant Disease and Mycobiology
2016-2018 Editorial Board of Mycobiology and Research in Plant Disease

INVITED SEMINARS

2019. 10. “Biology of a Plant-beneficial Mycorrhizae, *Serendipita bescii*”
 at Seoul National University, Suncheon National University, and Korean Institute of Science and Technology

EXTRACURRICULAR ACTIVITIES

- 1996.12- 1999. 02 Military Service as an infantry in Korean Army
2000.03 - 2001.02 Head of classical music club in Seoul National University
2004.03 - 2005.02 President of student association for Division of Plant Microbiology
2000.09 - 2000.12 President of student association for Division of Agricultural Biology
2014.08 – 2019.09 Leader of Noble tennis team and organizer of annual tournament event

References

Dr. Yong-Hwan Lee

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Field Plot Operations Manager

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