Jingqiu (Celia) Liao

324 Stocking Hall, Cornell University, Ithaca, NY 14850

Research Interests

Microbial evolution and ecology / Microbial biogeography / Environmental microbiology / Evolutionary and ecological genomics / Population genetics / Population genomics / Comparative genomics / Foodborne pathogens

Education

Ph.D. Candidate, Microbiology

2015-2020

CORNELL UNIVERSITY

Ithaca, NY, US

- Major advisor: Martin Wiedmann, Department of Food Science
- Minor advisors: Michael Stanhope (Genomics); Daniel Buckley (Soil Science)

Ph.D. exchange student, Microbiology

Fall 2019

MASSACHUSETTS INSTITUTE OF TECHNOLOGY

Cambridge, MA, US

· Advisor: Otto Cordero, Department of Civil and Environmental Engineering

M.S., Environmental Sciences

2011-2014

PEKING UNIVERSITY

Beijing, China

· Advisor: Yi Huang, Department of Environmental Sciences

B.S., Ecology

2007-2011

University of Science & Technology Beijing

Beijing, China

Research Experience _____

Graduate Research Associate

2016-Current

FIELD OF MICROBIOLOGY, CORNELL UNIVERSITY

Ithaca, NY, US

- Led projects (i) Listeria whole genome sequence reference sets for improved persistence assessment and source tracking, (ii) Microevolution of antimcrobial-resistant Salmonella enterica, and (iii) Influence of landscape on distribution of E. coli in watershed.
- Involved in projects (i) New Listeria species characterization, (ii) Development of mathematical models for developing environmental sampling programs for Listeria and assess L. monocytogenes-associated public health risks in the frozen food industry.

Research Associate 2014-2015

DEPARTMENT OF ENVIRONMENTAL SCIENCES, PEKING UNIVERSITY

Beijing, China

- Led project Biogeography and assembly of bacterial communities in lakes on Yungui Plateau, China.
- Involved in project Structure of bacterial community in lake sediment.

Graduate Research Assistant

2011-2014

DEPARTMENT OF ENVIRONMENTAL SCIENCES, PEKING UNIVERSITY

Beijing, China

• Involved in projects (i) Toxicity and ecology of oil contaminated soil in oil fields of China, (ii) Global Environment Outlook 5, and (iii) Regionalization of Dianchi Basin, China.

Publications

PREPRINT

[1] **Liao J**, Bergholz P, Wiedmann M. Terrestrial landscape impacts the biogeographic pattern of soil *Escherichia coli* via altering the strength of environmental selection and dispersal. bioRxiv doi:10.1101/2020.06.30.181495

PEER-REVIEWED

[1] Meeske AJ, Jia N, Cassel AK, Kozlova A, **Liao J**, Wiedmann M, Patel DJ, Marraffini LA. A phage-encoded anti-CRISPR enables complete evasion of type VI-A CRISPR-Cas immunity. Science 369:54-59

- [2] **Liao J**, Orsi RH, Carroll LM, Wiedmann M. Comparative genomics reveals different population structures associated with host and geographic origin in antimicrobial-resistant *Salmonella enterica*. Environ Microbiol 22:2811-2828
- [3] **Liao J**, Orsi RH, Carroll LM, Kovac J, Ou H, Zhang H, Wiedmann M. 2019. Serotype-specific evolutionary patterns of antimicrobial-resistant *Salmonella enterica*. BMC Evol Biol 19:132
- [4] Huang Y, Wang J, Chen H, **Liao J**, Huang M. Oil contamination and toxicity assessment of soil in main oil fields in China. Science Press, Beijing, China. June 2018 (Book in Chinese)
- [5] **Liao J**, Wiedmann M, Kovac J. 2017. Genetic stability and evolution of the *sigB* allele, used for *Listeria sensu stricto* subtyping and phylogenetic inference. Appl Environ Microbiol 83:e00306-17
- [6] **Liao J**, Cao X, Wang J, Zhao L, Sun J, Jiang D, Huang Y. 2017. Similar community assembly mechanisms underlie similar biogeography of rare and abundant bacteria in lakes on Yungui Plateau, China. Limnol Oceanogr 2:723-735
- [7] Cao X, Wang J, **Liao J**, Gao Z, Jiang D, Sun J, Zhao L, Huang Y, Luan S. 2016. Bacterioplankton communities response to key environmental variables in plateau freshwater lake ecosystems: a structural equation modeling analysis. Sci Total Environ 580:457-467
- [8] **Liao J**, Zhao L, Cao X, Sun J, Gao Z, Wang J, Jiang D, Fan H, Huang Y. 2016. *Cyanobacteria* in lakes on Yungui Plateau, China are assembled via niche processes driven by water physicochemical property, lake morphology and watershed land-use. Sci Rep 6:36357
- [9] **Liao J**, Cao X, Zhao L, Wang J, Gao Z, Wang MC, Huang Y. 2016. The importance of neutral and niche processes for bacterial community assembly differs between habitat generalists and specialists. FEMS Microbiol Ecol 92:fiw174
- [10] Wang J, Cao X, Chai L, **Liao J**, Huang Y, Tang X. 2016. Oxidative damage of naphthenic acids on the *Eisenia fetida* earthworm. Environ Toxico 31:1337-1343
- [11] **Liao J**, Wang J, Jiang D, Wang M, Huang Y. 2015. Long-term oil contamination causes similar changes in microbial communities of two distinct soils. Appl Microbiol Biotechnol 99:10299-10310
- [12] **Liao J**, Wang J, Huang Y. 2015. Bacterial community features are shaped by geographic location, physicochemical properties and oil contamination of soil in main oil fields of China. Microb Ecol 70:380-389
- [13] Cao X, Wang, J, **Liao J**, Huang Y, Sun J. 2015. The threshold responses of phytoplankton community to nutrient gradient in a shallow eutrophic lake. Ecol Indic 61:258-267
- [14] Wang J, Cao X, Chai L, **Liao J**, Huang Y, Tang X. 2015. Quantification and characterization of naphthenic acids from soil of oil exploring areas in China by GC/MS. Anal Methods 7:2149-2154
- [15] Wang J, Cao X, **Liao J**, Huang Y, Tang X. 2015. Carcinogenic potential of PAHs in oil-contaminated soils from the main oil fields across China. Environ Sci Pollut R 22:10902-10909
- [16] Yang Y, Wang J, **Liao J**, Xie S, Huang Y. 2015. Abundance and diversity of soil petroleum hydrocarbon egrading microbial communities in oil exploring areas. Appl Microbiol Biotechnol 99:1935-1946
- [17] Yang Y, Wang J, **Liao J**, Xie S, Huang Y. 2014. Distribution of naphthalene dioxygenase genes in crude oil contaminated soils. Microb Ecol 68:785-793
- [18] Liao J, Huang Y. 2013. Global trend in aquatic ecosystem research from 1992 to 2011. Scientometrics 98:1203-1219

Presentations

Microbiology Graduate Seminar, Cornell University

Feb., 2020

ORAL PRESENTER

Ithaca, NY, US

• Biogeography of edaphic Listeria across the United States

Microbiome, Cold Spring Harbor Laboratory Meeting

Jul., 2019

POSTER PRESENTER

Cold Spring Harbor, NY, US

• Comparative genomics reveals different population structures associated with host and geographic origin in antimicrobialresistant Salmonella enterica

Microbiology Graduate Seminar, Cornell University

Mar., 2019

ORAL PRESENTER

Ithaca, NY, US

• Nationwide distribution of *Listeria* in soil from natural environment

Biology of Genomes, Cold Spring Harbor Laboratory Meeting

May, 2018

POSTER PRESENTER

Cold Spring Harbor, NY, US

• Serotype-specific evolutionary pattern of antimicrobial-resistant *Salmonella enterica*.

Microbiology Graduate Seminar, Cornell University

ORAL PRESENTER

• Serotype-specific evolutionary pattern of antimicrobial-resistant Salmonella enterica

Ithaca, NY, US

Apr., 2018

Oct., 2017

Cornell University - Nestle Whole Genome Sequencing Training, Cornell University

ORAL PRESENTER

Ithaca, NY, US

· Phylogenetic tree reconstruction.

Ecological & Evolutionary Genomics, Gordon Research Conference

Jul., 2017

POSTER PRESENTER

SENTER Biddeford, ME, US

• Genetic stability and evolution of the sigB allele used for Listeria subtyping and phylogenetic inference

Microbiology Graduate Seminar, Cornell University

Mar., 2017

ORAL PRESENTER

Ithaca, NY, US

• Genetic stability and evolution of sigB allele of Listeria sensu stricto

M.S. Dissertation Seminar, Peking University

Jun., 2014

ORAL PRESENTER

Beijing, China

The physicochemical property and microbial structure of oil contaminated soils

The 6th SNU-PKU-KU Environmental Workshop, Seoul National University

Jul., 2012

ORAL PRESENTER

Seoul, South Korea

• Conservation of Chinese ecosystems: strategy and challenge

The 6th TU-USTB Joint Meeting on Sciences & Engineering Design, Tohoku University

Jul., 2010

ORAL PRESENTER

Sendai, Japan

• The influence of climate change on human ecology

Professional Activities

Journal reviewer Int Biodeterior Biodegradation / Scientometrics / Fund Appl Limnol / Ecotox Environ Safe / Mar Pollut Bull **Topic editor** Frontiers in Water (in progress)

Teaching and Mentoring Experience

Teaching assistantBioMi2900 General Microbiology, Cornell UniversitySpring 2018Teaching assistantEnvironmental ecology, Peking UniversityFall 2011MentorYaoqian Hu, Peking University2014-2015MentorMuke Huang, Peking University2018-2019MentorCatharine Carlin; Cornell University2019-2020

Skills

Biology Microbial isolation and culture / Molecular biology (DNA) / High-throughput sequencing / Field sampling **Bioinformatics** Programming (Python, R) / Computational biology / Biostatistics **Languages** English, Chinese

Leadership & Outreach

Social chairField of Microbiology Students (FoMS), Cornell University2017-2018Volunteer3M environmental sampling workshop, Cornell University2016-2018Student representativeThe 13th Annual Retreat of the Center for Infection & Pathobiology, Cornell UniversityApr. 2017Vice-captainPKUI Team, Solar Decathlon China2012-2013Vice-presidentCollege Student Union, Peking University2012-2013Vice-presidentCollege Student Union, University of Science & Technology Beijing2009-2010VolunteerBeijing Olympic GamesSummer 2008

Honors & Awards

Graduate Research Assistantship Cornell University	2016-Current
Conference & Research Travel Grants Cornell University	2017-2019
Cornell Graduate Fellowship Cornell University	2015-2016
Graduate Research Assistantship Peking University	2011-2014
Sander Scholarship Peking University	2013
Academic Innovation Award Peking University	2013
Honor Student for Excellent Academic Records Peking University	2012
The 2 nd Prize of Academic Innovation Contest Peking University	2012
National Scholarship China	2010
National Inspirational Scholarship China	2009
Freshmen Scholarship University of Science & Technology Beijing	2008
Student Innovation Program Scholarship University of Science & Technology Beijing	2008