



# PETER(YUNLIANG) LI

Dr. Peter (Yunliang) Li is a seasoned researcher specializing in molecular biology and environmental microbiome. His early work focused on gene function and regulatory networks. In subsequent research, he applied amplicon and shotgun metagenomic sequencing methods to investigate the role of microbiomes in farming systems and petroleum-contaminated soil remediation.

## EDUCATION

- Jul. 2014  
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Sept. 2009 • **Ph.D. in Developmental Biology**  
University of Chinese Academy of Sciences  Beijing, China
- Jul. 2009  
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Sept. 2005 • **M.S. in Developmental Biology**  
University of Chinese Academy of Sciences  Beijing, China
- Jul. 2005  
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Sept. 2001 • **B.S. in Biotechnology**  
Shandong Agricultural University  Shandong, China

## RESEARCH EXPERIENCE

- Current  
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Jun. 2019 • **Postdoctoral Researcher & Research Associate**  
University of Saskatchewan  Saskatoon, SK
  - Proficiently prepared DNA libraries for amplicon metagenomic sequencing
  - Developed efficient pipelines for processing shotgun and amplicon metagenomic sequencing data.
  - Skillfully processed and analyzed microbiome sequencing data utilizing various bioinformatic and biostatistical tools.
  - Optimized the qPCR reaction system for amplifying genes associated with nitrogen metabolism in soil.
- Apr. 2018  
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Apr. 2015 • **Postdoctoral Researcher**  
Agriculture and Agri-Food Canada  Swift Current, SK & Quebec, QC
  - Conducted next-generation sequencing of the root-associated microbiome using the Illumina sequencing platform.
  - Expertly processed and analyzed amplicon and shotgun metagenomic sequencing data.
- Mar. 2015  
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Nov. 2014 • **Postdoctoral Researcher**  
McGill University  Montreal, QC
  - Modified potato disease-related gene (WRKY1) using CRISPR/Cas9 system
  - Optimized transient transformation systems of potato and wheat leaf protoplasts

## CONTACT

 [liyunliang81@gmail.com](mailto:liyunliang81@gmail.com)

 [github](#)

 [LinkedIn](#)

## SKILLS

Molecular biology  
Bioinformatics  
Statistics

## LANGUAGES

R  
Python  
Bash  
C++

Made with the R package  
[pagedown](#).

The source code is available on  
[github.com/Yunliang/cv](https://github.com/Yunliang/cv).

Last updated on 2024-01-14.

Jul. 2014  
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Sept. 2005

● **M.S. and Ph.D. student**

Institute of Botany, Chinese Academy of Sciences

📍 Beijing, China

- Conducted gene function studies and deciphered gene regulatory network using molecular biological techniques, including gene cloning, vector construction, qPCR, protein-DNA interaction assays, protein-protein interaction analysis, and sub-cell localization using protein-GFP fusion



## SELECTED PUBLICATIONS

2024  
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2024

● **Characteristics of root-associated microbiome across the development of wheat cultivars over 100 years using a simple and effective sampling method**

📍 University of Saskatchewan

- Yunliang Li, Steven D. Mamet, Bobbi L. Helgason, Curtis J. Pozniak, Andrew G. Sharpe, Steven D Siciliano
- Ready for submission

2023  
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2023

● **Contrasting nitrogen fertilization and Brassica napus (canola) variety development impact recruitment of the root-associated microbiome**

📍 University of Saskatchewan

- Yunliang Li, Sally Lynne Vail, Melissa M. Arcand, and Bobbi Helgason
- Journal: Phytobiomes Journal 7(1): 125-137
- <https://doi.org/10.1094/PBIOMES-07-22-0045-R>

2023  
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2023

● **Root and rhizosphere fungi associated with the yield of diverse Brassica napus genotypes**

📍 University of Saskatchewan

- Yunliang Li, Navid Bazghaleh, Sally Vail, Steven D. Mamet, Steven D. Siciliano, and Bobbi Helgason
- Journal: Rhizosphere p.100677
- <https://doi.org/10.1016/j.rhisph.2023.100677>

2021  
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2021

● **Effect of arbuscular mycorrhizal inoculation and crop sequence on root associated microbiome, crop productivity and nutrient uptake in wheat-based and flax-based cropping systems**

📍 University of Saskatchewan

- Yunliang Li, Mario Laterriere, Chih-Ying Lay, Rim Klabi, Jacynthe Masse, Marc St-Arnaud, Etienne Yergeau, Newton Z. Lupwayi, Yantai Gan, and Chantal Hamel
- Journal: Applied Soil Ecology 168: 104136
- <https://doi.org/10.1016/j.apsoil.2021.104136>

2020  
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2020

- **Long-term effects of nitrogen and phosphorus fertilization on soil microbial community structure and function under continuous wheat production**  
📍 University of Saskatchewan
  - Yunliang Li, Julien Tremblay, Luke Bainard, Barbara Cade-Menun, Chantal Hamel
  - Journal: Environmental microbiology 22(3): 1066-1088
  - <https://doi.org/10.1111/1462-2920.14824>

2019  
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2019

- **Influence of introduced arbuscular mycorrhizal fungi and phosphorus sources on plant traits, soil properties, and rhizosphere microbial communities in organic legume-flax rotation**  
📍 Agriculture and Agri-Food Canada
  - Yunliang Li, Yantai Gan, Newton Lupwayi, and Chantal Hamel
  - Journal: Plant and Soil 443(1-2):87-106
  - <https://doi.org/10.1007/s11104-019-04213-8>

2019  
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2019

- **Site specificity in establishment of a commercial arbuscular mycorrhizal fungal inoculant**  
📍 Agriculture and Agri-Food Canada
  - Vasilis Kokkorisa, Yunliang Li, Chantal Hamel, Keith Hanson, Miranda Harta
  - Journal: Science of The Total Environment 660:1135-1143
  - <https://doi.org/10.1016/j.scitotenv.2019.01.100>

2015  
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2015

- **Transcription factor StWRKY1 regulates phenylpropanoid metabolites conferring late blight resistance in potato**  
📍 McGill University
  - Yogendra, Kalenahalli N., Arun Kumar, Kobir Sarkar, Yunliang Li, Doddaraju Pushpa, Kareem A.Mosa, Raj Duggavathi, and Ajjamada C. Kushalappa
  - Journal: Journal of Experimental Botany 660:1135-1143
  - <https://doi.org/10.1093/jxb/erv434>

2014  
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2014

- **Arabidopsis VQ-motif- containing protein 29 represses seedling de-etiolation by interacting with PIF1**  
📍 University of Chinese Academy of Sciences
  - Yunliang Li, Yanjun Jing, Junjiao Li, Gang Xu and Rongcheng Lin
  - Journal: Plant Physiology 164: 2068- 2080
  - <https://doi.org/10.1104/pp.113.234492>

2013  
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2013

- **Arabidopsis Chromatin Remodeling Factor PICKLE Interacts with Transcription Factor HY5 to Regulate Hypocotyl Cell Elongation**  
📍 University of Chinese Academy of Sciences
  - Yanjun Jing, Dong Zhang, Xin Wang, Weijiang Tang, Wanqing Wang, Junling Huai, Gang Xu, Dongqin Chen, Yunliang Li and Rongcheng Lin
  - Journal: Plant Cell 25: 242-256
  - <https://doi.org/10.1105/tpc.112.105742>