

Haikuo Li

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EDUCATION

PhD Student, Program in Molecular Genetics and Genomics

8/2019 – present

Washington University in St. Louis, MO, United States

- Thesis mentor: Benjamin D. Humphreys, M.D., Ph.D.

Bachelor of Science, Bioscience (Zhiyuan Honors Program)

9/2015 – 6/2019

Shanghai Jiao Tong University, Shanghai, China

- 2019 Top 0.2% Bachelor Thesis: Rank #1 in Bioscience
- 2019 Outstanding Graduate in Bachelor Degree, Shanghai

Visiting Student, Immunobiology

6/2018 – 4/2019

Yale University, New Haven, CT, United States

- Research supervisor: Aaron M. Ring, M.D., Ph.D.

PUBLICATIONS

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1. **Li, H.**, Dixon, E. E., Wu, H., & Humphreys, B. D. (2022). Comprehensive single-cell transcriptional profiling defines shared and unique epithelial injury responses during kidney fibrosis. **Cell Metabolism**. (PDF)
 2. **Li, H.**, & Humphreys, B. D. (2022). Mouse kidney nuclear isolation and transcriptional profiling with single-cell combinatorial indexing RNA sequencing. **STAR Protocols**. (in revision)
 3. **Li, H.**, & Humphreys, B. D. (2022). New functions for basophils identified in kidney fibrosis. **Nature Immunology**, 23(6), 824-825. (PDF)
 4. Muto, Y*, **Li, H.*** (equal contribution), & Humphreys, B. D. (2022). Single-cell transcriptomics. Innovations in Nephrology: Breakthrough Technologies in Kidney Disease Care (**Book Chapter**). (in press)
 5. **Li, H.**, & Humphreys, B. D. (2021). Single cell technologies: Beyond microfluidics. **Kidney360**, 2(7), 1196. (PDF)
 6. **Li, H.**, & Humphreys, B. D. (2020). Surveying the human single-cell landscape. **Kidney International**, 98(6), 1385-1387. (PDF)
 7. Ku, X.*, **Li, H.*** (equal contribution), Wang, J.*, et al. Jin, J.#, Yan, W#. Proteomic portrait of human lymphoma revealed protein molecular fingerprint for disease specific subtypes and progression. **Phenomics**. (under review)

RESEARCH EXPERIENCE

PhD Student, Benjamin Humphreys Lab

4/2020 – present

Division of Nephrology, Washington University in St. Louis

- Developing a single-cell atlas of kidney fibrosis with single-cell multimodal profiling

- Studying metabolic mechanisms that drive kidney fibrosis
- Developing single-cell combinatorial indexing (split-pool barcoding) platforms

PhD Rotation Student, Tim Peterson, Sidharth Puram, Benjamin Humphreys Labs 8/2019 – 4/2020

Washington University in St. Louis

- Peterson Lab: Understanding the intracellular effects of Cationic Amphipathic Drugs on organelles
- Puram Lab: Studying head and neck cancer by CITE-seq
- Humphreys Lab: Characterizing kidney injury and repair markers by RNAscope

Visiting Student, Aaron Ring Lab 6/2018 – 4/2019

Department of Immunobiology, Yale University

- Modulating immune cytokines by protein engineering

Undergraduate Researcher, Wei Yan Lab 9/2016 – 7/2018

Shanghai Center for Systems Biomedicine, Shanghai Jiao Tong University

- Identification of biomarkers of lymphoma with mass spectrometry; clinic proteomics

Summer Intern, Manyuan Long Lab 6/2017 – 8/2017

Department of Ecology and Evolution, The University of Chicago

- Identification of mammalian positively selected genes by polygenic analysis

Science Olympiad (Mathematics), Shandong Province Team, China 7/2014 – 2/2015

- Top10 students selected to participate in the Chinese Mathematics Olympics

SKILLS

Wet lab experiment

- Extensive experience in single-cell and single-nucleus library generation from diverse technologies, including 10X Genomics, sci-RNA-seq, SHARE-seq and INTACT, as well as multimodal profiling including RNA-seq, ATAC-seq, CARLIN and CITE-seq
- Extensive experience in molecular biology technologies such as cloning, vector construction, qPCR, immunohistology, and in-situ hybridization
- Extensive experience in tissue culture including primary cell isolation, immunocytochemistry and Seahorse metabolic measurement
- Strong experience in animal work such as mouse kidney disease surgery (UUO/IRI) and tumor implantation
- Strong experience in clinical sample management and processing such as human kidney dissection
- Strong background in protein chemistry including mass spectrometry sample preparation & recombinant protein preparation and protein liquid chromatography

Computational workflow

- Extensive experience in using Python, R, Shell and Jupyter
- Extensive experience in single-cell sequencing data preprocessing and analysis including UMAP visualization, data integration, sample demultiplexing, cell trajectory inference, fate mapping, gene activity prediction and multimodal analysis at the million-cell level
- Strong experience in analysis of bulk RNA-seq, proteomics and metabolomics data
- Extensive experience in data mining and discovering biological insights
- Strong training background in mathematics

TEACHING EXPERIENCE

Assistant Instructor, Washington University in St. Louis	8/2022 – present
• Structural bioinformatics of proteins (Bio4525)	
Peer Study Mentor, Washington University in St. Louis	1/2022 – 6/2022
• Genomics (Bio5488) and Python-based coding tutoring	
Assistant Instructor, Washington University in St. Louis	1/2021 – 6/2021
• Genomics (Bio5488)	
Undergraduate Teaching Assistant, Shanghai Jiao Tong University	
• College Genetics Course (2/2019–6/2019)	
• College Macrobiology Course (2/2018–6/2018)	
• College Biochemistry Course (9/2017–1/2018)	

PRESENTATIONS & POSTERS

Speaker, Nephrology Division Research Seminars, Washington University in St. Louis	9/2022
Retreat Talk (MGG/CSB/HSG/IMSD programs), Washington University in St. Louis	9/2022
POSTER (PDF) PhD program retreat, Washington University in St. Louis	9/2022
• Cell profiling defines metabolic dysregulation in kidney fibrosis	
DBBS Friday Talks (MGG/CSB/HSG/IMSD programs), Washington University in St. Louis	5/2022
REBUILDING A KIDNEY Spring Meeting lighting talk	4/2022
PhD Program Thesis Committee Meeting, Washington University in St. Louis	8/2021, 4/2022
REBUILDING A KIDNEY Work in Progress small group meeting	12/2021
PhD Program Qualifying Examination Committee Meeting	9/2020
POSTER (PDF) SJTU Academic Festival (Best Poster Award)	2/2019
• Modulating the Tumor-Targeting Specificity of “Decoy-Resistant” Interleukin-18 by Protein Engineering	
POSTER (PDF) Human Proteome Organization World Congress	10/2018
• Clinical Proteomics Analysis using Data Independent Acquisition-Mass Spectrometry (DIA-MS) Identified Classifiers for Molecular Characterization of Lymphoma	
POSTER (PDF) SJTU Academic Festival (Best Poster Award)	12/2017
• Detecting Positively Selected Genes among Mammalian Species Using Phylogenetic Analysis of Maximum Likelihood	

HONORS & ACTIVITIES

Member, ASN (American Society of Nephrology)	2020 - present
Top 0.2% Bachelor Thesis of Shanghai Jiao Tong University (URL) (Thesis PDF)	2019
Outstanding Graduate in Bachelor Degree, Shanghai	2019
Academic Excellence Scholarship (First-class), Shanghai Jiao Tong University	2016, 2017, 2018
Rank #1 Student Presentation, National Biology Education Conference of Chinese Ministry of Education	2018
Vice President, Students' Union of Zhiyuan Honors Program, Shanghai Jiao Tong University	2017 - 2018
Top 0.1% in Chinese University Entrance Examination (681 points)	2015
Bronze medal, Chinese Mathematical Olympiad (CMO)	2014