Haikuo Li

haikuo.li@wustl.edu | Washington University in St. Louis
https://haikuoli.github.io/ | Twitter @HaikuoLi

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.
- Thesis committee: Ting Wang, Samantha Morris, Jeffrey Millman, Allegra Petti, Michael Meers

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

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**. (accepted)
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**). (**PDF**)
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 polygenetic 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 interference, 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