Siyi Chen, Ph.D.

RESEARCH INTERESTS

My research addresses Mendelian randomization/TWAS, causal inference, structural equation modeling, stochastic modeling, species problem, approximate Bayesian computation, and Bayesian statistics.

EDUCATION

Rice University Doctor of Philosophy, Statistics	$\begin{array}{c} \text{Houston, TX} \\ \text{Dec 2020} \end{array}$
Rice University Master of Science, Statistics	Houston, TX May 2015
Peking University Bachelor of Science, Chemistry	Beijing, China July 2013

Positions and Employment

Biostatistics & Data Science (BSDS), LSU Health School of Public Health Assistant Professor	2024-present New Orleans, LA
RUSH Alzheimer's Disease Center, Rush University Biostatistician	2023-2024 Chicago, IL
$ {\bf Department \ of \ Neurological \ Sciences, \ Rush \ University \ Medical \ Center} \\ {\it Instructor} $	2023-2024 Chicago, IL
Division of Biostatistics, School of Public Health, University of Minnesota Post-Doctoral Associate	2021-2023 Minneapolis, MN
Department of Statistics, Rice University Graduate Research Assistant	$2016\text{-}2020$ $Houston,\ TX$

RESEARCH EXPERIENCE

Postdoctoral research associate

2021-2023

Department of Biostatistics, University of Minnesota

Minneapolis, MN

• Supervisor: Dr. Wei Pan

Research Assistant

2017-2020 Houston, TX

Department of Statistics, Rice University

• Supervisor: Dr. Marek Kimmel

• Developed a sequential truncated-population approximate Bayesian computation (ABC) algorithm and applied the ABC methods to symmetric Dirichlet-multinomial model and the proposed asymmetric Dirichlet-multinomial model to estimate hematopoietic stem cells count

Research Assistant 2016-2017

Department of Statistics, Rice University

Houston, TX

- Simulated barcoding experiments and empirical distributions of the estimators of barcodes number in mice blood cell samples
- Developed a multi-type Markov age-dependent branching process for modeling hematopoietic stem cell expansion

Siyi Chen MR-SuSiE: a robust Mendelian randomization framework for identifying causal proteins in Alzheimer's disease with correlated and invalid instruments, *Submitted*

Siyi Chen Two-sample bi-directional causality between two traits with some invalid IVs in both directions using GWAS summary statistics, *Human Genetics and Genomics Advances*, *July 10*, 2025

Katie A Matatall, Mira Jeong, **Siyi Chen**, Deqiang Sun, Fengju Chen, Qianxing Mo, Marek Kimmel, Katherine King (2016) "Chronic infection depletes hematopoietic stem cells through stress-induced terminal differentiation", *Cell Reports*, 17, 2584–2595.

Siyi Chen, Zhaotong Lin, Xiaotong Shen, Ling Li, and Wei Pan "Inference of causal metabolite networks in the presence of invalid instrumental variables with GWAS summary data," *Genetic Epidemiology (2023)*

Siyi Chen "Statistical modeling for species count data with heterogeneity," https://scholarship.rice.edu/handle/1911/109642

Siyi Chen, Katherine King, Marek Kimmel "Statistical inference from stem cell barcoding data using adaptive approximate Bayesian computation", *Submitted*.

https://assets.researchsquare.com/files/rs-187743/v1_covered.pdf?c=1631853488

Presentations

ENAR 2025 Presentation	New Orleans, LA 2025
STATGEN 2024 Presentation	Pittsburgh, PA 2024
2023 IGES Annual Meeting Presentation	Nashville, TN 2023

TEACHING EXPERIENCE

Teaching Assistant

Department of Statistics

Rice University

Houston, TX

- Mathematical Probability I, Fall 2015
- Probability, Fall 2015 & Fall 2016
- Probability in Bioinformatics and Genetics, Spring 2016

Instructor

LSU Health
School of Public Health

New Orleans, LA

- Introduction to Biostatistics, Fall 2025
- Statistical Inference I, Fall 2025
- Applied Linear Models, Spring 2025
- Nonparametric Statistics, Spring 2025
- Introduction to Biostatistics, Fall 2024
- Statistical Inference I, Fall 2024