# Lin Gui

5747 S. Ellis Avenue, Jones 203/204, Chicago, IL, 60637 Email: glin6@uchicago.edu; Website: https://gl-ybnbxb.github.io/

### **EDUCATION**

Ph. D. in Statistics, The University of Chicago, Chicago, USA
M.S. in Statistics, The University of Chicago, Chicago, USA
B.S. in Statistics, University of Science and Technology of China, Hefei, China
2018-2020
2014-2018

### RESEARCH INTERESTS

- Statistical Inference and Multiple Testing
- Generative Models
- Causal Inference and Machine Learning
- Biostatistics

### **PUBLICATIONS**

• Concept Algebra for Score-Based Conditional Models

Zihao Wang, Lin Gui, Jeffrey Negrea, Victor Veitch *NeurIPS* 2023

• Causal Estimation for Text Data with (Apparent) Overlap Violations

Lin Gui, Victor Veitch *ICLR* 2023

• Detecting Multiple Replicating Signals using Adaptive Filtering Procedures

Jingshu Wang, Lin Gui, Weijie J. Su, Chiara Sabatti, Art B. Owen *The Annals of Statistics* 50 (4), 1890-1909

#### RESEARCH

- A Theoretical and Practical Analysis of the Heavy-Tailed Combination Test for Global Test With Correlated Hypotheses
- Undertook comprehensive theoretical evaluations to decipher the intricacies of the state-of-theart Cauchy combination test and its expansion, termed the heavy-tailed combination test, tailored for the global test with correlated hypotheses.
- Conducted empirical studies, offering a general practical guideline for the Cauchy/heavy-tailed combination test
- Enhanced the heavy-tailed combination test into a multiple testing procedure adept at controlling the family-wise error rate (FWER) and introduced a shortcut for this closed testing procedure.
- Implemented the method on genetic data to address real-world challenges.
- Gene Transcription Mechanism in 3D Genome
- Designed a statistical tool aimed at identifying gene pairs with a high likelihood of cotranscription due to shared enhancers (close to each other). A core challenge is accommodating the 3D dynamic structures of chromosomes. Thus, spatial details of gene pairs must be analyzed using 3D genome data, such as Hi-C data.

# **CONFERENCES AND PRESENTATIONS**

Jul. 2023
May. 2023
Aug. 2021

# **CODING SKILLS**

- R, Python, MATLAB, SQL
- PyTorch, Numpy, Pandas

# **HONORS & AWARDS**

• Nominee, The 3/th. Guo Moruo Scholarship (The highest honor at USTC)	2017
Winner, Outstanding Student Scholarship, USTC	2016-2017
Winner, China National Scholarship, USTC	2015