

WENJING MA

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Ph.D. candidate, Computer Science and Informatics (BMI track), Emory University

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Personal website: <https://marvinquiet.github.io/>

Education

Ph.D. student in Computer Science (Biomedical Informatics Track)

expected 2023

Department of Computer Science, Emory University, Atlanta, GA, U.S.

GPA: 4.0/4.0

Advisor: Dr. Hao Wu

Visiting Scholar in Bioinformatics

08/2018 -- 06/2019

Center for Public Health Genomics, University of Virginia, Charlottesville, VA, U.S.

Advisor: Dr. Chongzhi Zang

M.S. and B.S. in Computer Science and Technology

09/2010 -- 06/2017

Beijing University of Posts and Telecommunications (BUPT), Beijing, China

Professional Experience

Database Administrator

08/2017 -- 06/2018

Investment Center, State Administration of Foreign Exchange, Beijing, China

Research Interests

- Develop machine learning and deep learning methods with applications in single-cell genomics data
- Integrate single-cell multi-omics data to reveal cell-type-specific gene regulatory network and explore epigenetic diseases such as neurodegenerative disease, obesity, cancer, etc.

Publications

(*: equal contribution)

Peer-Reviewed Journal

1. **Wenjing Ma**, Sumeet Sharma, Peng Jin, Shannon L. Gourley, Zhaohui S. Qin. *LRcell*: detecting the source of differential expression at the sub-cell-type level from bulk RNA-seq data. *Briefings in Bioinformatics*, (2022). [DOI](#) [[R package](#)]
 - 20-min talk @ [AWSOM Workshop 2022](#);
2. **Wenjing Ma**, Kenong Su, Hao Wu. Evaluation of some aspects in supervised cell type identification for single-cell RNA-seq: classifier, feature selection, and reference construction. *Genome Biology*, (2021). [DOI](#) [[project link](#)]
 - Honorable Mention Poster @ GSD 2021;
 - Best 5-minute presentation award @ Emerging Data Science Methods for Complex Biomedical and Cyber Data 2021

3. Shengen S. Hu, Lin Liu, Qi Li, **Wenjing Ma**, Michael J. Guertin, Clifford A. Meyer, Ke Deng, Tingting Zhang, Chongzhi Zhang. Intrinsic bias estimation for improved analysis of bulk and single-cell chromatin accessibility profiles using SELMA. *Nature Communications*, (2022). [DOI](#), [\[package\]](#)
4. **Wenjing Ma***, Zhenjia Wang*, Yifan Zhang, Neal E. Magee, Yayi Feng, Ruoyao Shi, Yang Chen, Chongzhi Zhang. BARTweb: a web server for transcriptional regulator association analysis. *NAR Genomics and Bioinformatics*, (2021). [DOI](#) [\[website\]](#) [\[RECOMB 2019 poster\]](#)

Book Chapters

5. Ronnie Y. Li, **Wenjing Ma**, Zhaohui S. Qin. Approaches to Marker Gene Identification from Single-Cell RNA-Sequencing Data. In *Handbook of Statistical Bioinformatics* (pp. 71-84). Springer, Berlin, Heidelberg. (2022) [DOI](#)

Collaborations

6. Yulin Jin, Kenong Su, Ha Eun Kong, **Wenjing Ma**, Zhiqin Wang, Yujing Li, Ronghua Li, Emily Allen, Hao Wu, Peng Jin. Cell-type specific DNA methylome signatures reveal epigenetic mechanisms for neuronal diversity and neurodevelopment disorder. *Human Molecular Genetics*, (2022) [DOI](#)

Under Review

7. **Wenjing Ma**, Jiaying Lu, Hao Wu. Cellcano: supervised cell type identification for single cell ATAC-seq data. *Under second-round revision at Nature Communications*, (2022). [\[Python package\]](#) [\[Documentation\]](#)
 - 3rd place (out of 47) Student Poster Presentation Award @ GSD 2022 [\[News\]](#)
 - 2023 ASA Section on Statistics in Genomics and Genetics (SGG) Student Paper Award [\[News\]](#)
 - Selected as one of four finalists to compete for the MCBIOS 2023 Young Scientist Excellence Award (YESA) in the Student category
8. Sumeet Sharma*, **Wenjing Ma***, Kerry J. Ressler, Thea Anderson, Dan. C. Li, Peng Jin, Shannon L. Gourley, Zhaohui Qin. Dysregulation of prefrontal oligodendrocyte lineage cells across mouse models of adversity and human major depressive disorder. *Under revision*, (2022).

Honors

Fellowship

- Women in Natural Sciences Fellowship (@ Emory University, GA, U.S., 2019 -- Present)

Travel Grants

- 2023 ABGOD (@ The University of Texas at Dallas)
- 2022 Grad Cohort for Women (@ New Orleans)

Professional Activities and Services

- Invited reviewer for International Journal of Computational Biology and Drug Design (IJCBD).
- Invited reviewer for IEEE BIBM 2022.
- Invited sub-reviewer for ISMB/ECCB 2023.

Teaching Experiences

BIOS 555: High-throughput data analysis using R and Bioconductor	<i>Fall 2022</i>
QTM 100: Intro to Statistical Inference - teaching weekly lab in R programming	<i>Fall 2021</i>
CS 584: Biomedical Image Analysis - teaching assistant	<i>Spring 2021</i>
CS 253: Data Structures and Algorithms - teaching assistant	<i>Fall 2020</i>
CS 224: Foundations of Computer Science - teaching assistant	<i>Spring 2020</i>