# **ZIQI ZHANG**

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# **EDUCATION**

#### Georgia Intitute of Technology, GA, USA

Sep 2020 - present

Ph.D. in Computational Science and Engineering

• Advisor: Prof. Xiuwei Zhang

# Georgia Intitute of Technology, GA, USA

Sep 2019 - present

Master of Electrical and Computer Engineering

• Advisor: Prof. Xiuwei Zhang

# Beihang University, Beijing, China

Sep 2015 - July 2019

Bachelor of Electronic and Information Engineering

• Beidou Honor Class

• Thesis advisor: Prof. Huaping Xu

# **PUBLICATION**

# **Journal & Conference papers**

- 1. **Ziqi Zhang**, Haoran Sun, Ragunathan Mariappan, Xinyu Chen, Xi Chen, Mika S Jain, Mirjana Efremova, Sarah A Teichmann, Vaibhav Rajan, Xiuwei Zhang. "scMoMaT jointly performs single cell mosaic integration and multimodal bio-marker detection", *Nature Communication*, 2023
- 2. **Ziqi Zhang**, Chengkai Yang, and Xiuwei Zhang. "scDART: integrating unmatched scRNA-seq and scATAC-seq data and learning cross-modality relationship simultaneously" *Genome Biology*, 2022
- 3. **Ziqi Zhang** and Xiuwei Zhang. "Inference of high-resolution trajectories in single-cell RNA-seq data by using RNA velocity", *Cell Reports Methods*, 2021
- 4. Vaibhav Rajan, Ziqi Zhang, Carl Kingsford, and Xiuwei Zhang. "Maximum likelihood reconstruction of ancestral-networks by integer linear programming", *Bioinformatics*, 2021

# **Preprints & Working papers**

- 1. **Ziqi Zhang**, Xinye Zhao, Peng Qiu, and Xiuwei Zhang. "scDisInFact: disentangled learning for integration and prediction of multi-batch multi-condition single cell RNA-sequencing data", *submitted to RECOMB*, 2022
- 2. **Ziqi Zhang**, Jongseok Han, Le Song, and Xiuwei Zhang. "Inferring cell-specific gene regulatory networks from single cell gene expression data", *BioRxiv*, 2022
- 3. Hechen Li, **Ziqi Zhang**, Michael Squires, Xi Chen, Xiuwei Zhang, "scMultiSim: simulation of multi-modality single cell data guided by cell-cell interactions and gene regulatory networks", *BioRxiv*, 2022
- 4. **Ziqi Zhang** and Xiuwei Zhang. "Velosim: Simulating single cell gene-expression and RNA velocity", *BioRxiv*, 2021

#### Workshop

- 1. **Ziqi Zhang** and Xiuwei Zhang. "Integrating unmatched scRNA-seq and scATAC-seq data and learning cross-modality relationship simultaneously" **Spotlight**, *16th Machine Learning in Computational Biology (MLCB)*, 2021
- 2. **Ziqi Zhang**, Haoran Sun, Ragunathan Mariappan, Xi Chen, Mika S Jain, Mirjana Efremova, Sarah A Teichmann, Vaibhav Rajan, Xiuwei Zhang. "Integrating unpaired scRNA-seq and scATAC-seq with unequal cell type compositions" *ICML worshop on Computational Biology*, 2021

#### **Presentations**

- 1. Talk: scDisInFact: the disentangled learning for integration and prediction of multi-batch multi-condition single-cell RNA sequencing data, HotCSE, School of Computational Science and Engineering, Georgia Institute of Technology, 2022
- 2. Talk: scMoMaT: Multimodal Data Integration Software, AWSOM-Atlanta Workshop on Single-cell Omics, 2022
- 3. Spotlight presentation: *Integrating unmatched scRNA-seq and scATAC-seq data and learning cross-modality relationship simultaneously*, MLCB, 2021
- 4. Flash talk: Learning latent embedding of multi-modal single cell data and cross-modality relationship simultaneously, CSHL, 2021
- 5. Poster presentation: CeSpGRN: Inferring cell-specific gene regulatory networks from single cell gene expression data, RECOMB, 2022
- 6. Poster presentation: scMoMaT: Integrating arbitrary number of single-cell multi-omics matrices using matrix-trifactorization, USC-QCB, 2022
- 7. Poster presentation: Inference of multiple trajectories in single-cell RNA-Seq data from RNA-velocity, VIB, 2020
- 8. Poster presentation: *Inference of multiple trajectories in single-cell RNA-Seq data from RNA-velocity*, RSGDREAM, 2020
- 9. Poster presentation: *Inference of multiple trajectories in single-cell RNA-Seq data from RNA-velocity*, CSHL-Asia, 2020
- 10. Oral presentation: Computational methods to study cell mechanisms from single cell data, Georgia Scientific Computing Symposium (GSCS), 2020

#### **TEACHING**

#### **Courses**

1. Head Teaching Assistant: CSE 6140 Algorithm 2021Fall

2. Teaching Assistant: CSE 8803 Machine learning in Computational Biology 2021Spring

3. Teaching Assistant: CSE 6140 Algorithm 2020Fall

#### Advisee

1. **Xinye Zhao**: graduate student in Electrical and Computer Engineering

Nov 2021-Oct 2022 **Project**: scDisInFact: disentangled learning for integration and prediction of multi-batch multi-condition single cell RNA-sequencing data

2. **Xinyu Chen**: graduate student in Bioengineering

August 2021-Nov 2021 **Project**: scMoMaT: Mosaic integration of single cell multi-omics data using matrix tri-factorization

3. **Jongseok Han**: graduate student in Computational Science and Engineering

\*\*Project: Inferring cell-specific gene regulatory networks from single cell gene expression data\*

\*\*Jun 2021-Nov 2021\*\*

\*\*Project: Inferring cell-specific gene regulatory networks from single cell gene expression data\*

4. **Chengkai Yang**: undergraduate student in Computer Science

Jun 2020-Jan 2021 **Project**: Learning latent embedding of multi-modal single cell data and cross-modality relationship simultaneously

# **AWARD**

#### Ph.D. in Georgia Tech

1. Thanks a teacher award as teaching assistant in CSE6140 Nov 2020

# **Undergraduate in Beihang University**

1. Outstanding Undergraduate in Beihang University 2019

2. Chinese Graduate Entrance Exam Waiver 2019

3. Learning Merit Scholarship in Beihang University	2018
4. Scholarship for student excellent in Science and Technology competition	2018
5. The Third Prize of National Undergraduate Electronic Design Contest	2017
6. The First Prize of BUAA Physics competition	2015

# **SERVICE**

Journal reviewer: Bioinformatics, Cell Reports Methods, BMC Bioinformatics, BMC Endocrine Disorders

Conference reviewer: RECOMB 2022, ISMB 2022

# **SKILLS**

Languages: Chinese, English

**Programming:** Python, Pytorch, R, LaTex, Linux, MATLAB, C++