# Qimin Zhang

#### **EDUCATION**

Email: qqz5133@psu.edu Phone: +1(814)529-3984Google Scholar link

State College, USA

Aug. 2019 - May. 2024

Beijing, China

Sep. 2016 - June. 2019

Beijing, China

Sep. 2012 - June. 2016

• Beihang University

Bachelor of Engineering in Aircraft Airworthiness Bachelor of Science in Applied Mathematics (minor)

• University of Chinese Academy of Sciences

Master of Engineering in Computer Technology

### SKILLS

• Languages C++, C, Python, Shell

• The Pennsylvania State University

Ph.D in Computer Science and Engineering

• Technical Skills and Tools Machine Learning (scikit-learn, PyTorch), Bioinformatics, Cloud Computing (AWS), Web Development (HTML/CSS/JS), Linux, Git, Docker

### WORK EXPERIENCE

# • Laboratory Corporation of America Holdings (LabCorp) Senior Bioinformatics Specialist

Pittsburgh, PA

- July 2023 present
- Developed computational methods and pipelines for virus subtyping and strain-level identification from the metagenomics samples.
- Developed and re-engineering the data structure and algorithms of tools and software to provides verbose log, improve accuracy, and reduce computational cost.
- Developed pipeline for metabolomic pathway prediction from microbiome sequencing data- using statistical and machine learning methods, focusing on the discovery of biomarkers related to women's health diseases.
- Laboratory Corporation of America Holdings (LabCorp)

  Data Science / Bioinformatics Summer Intern

Pittsburgh, PA May 2022 - Aug 2022

- Developed a set of highly accurate data pipelines using machine learning to predict gut metabolites and metagenome functions from gut microbiome data
- Managed to work on a Colorectal Cancer dataset and achieved all metabolites well predicted

#### Research Experience

Developed open-source software and tools, such as Scallop2 and Aletsch, have **over 15,000 downloads** in Bioconda, and the number is actively increasing.

# • Bioinformatics Algorithms

Sep 2019 - Present

 $\circ$  Transcriptome assembly & bulk/single-cell RNA-seq data analysis

Developed a C++ software Scallop2, a transcript assembler specifically optimized for paired-/multi-end RNA-seq data. Designed and implemented a dynamic programming algorithm and an enhanced consensus algorithm to improve 85.9% and 46.6% in precision comparing with two leading tools at the same level of sensitivity.

# • Applied Machine Learning

Sep 2017 - present

• High-throughput Computing

Evaluated machine learning techniques for predicting resource usage in high-throughput computing. Applied the density-based Clustering methodologies to minimize resource waste and reduce time, cores and memory consumption by 13.82%, 16.62%, 49.15%, respectively.

#### o Healthcare

Explored machine learning techniques to biomedical problems. Extracted feature of sputum sound signals using wavelet transform algorithm. Implemented a BPNN model and improved the precision of detecting sputum to 84.53%.

## Publications

- 1. Hayden Brochu, Kuncheng Song, **Qimin Zhang**, Qiandong Zeng, Adib Shafi et al. A program for real-time surveillance of SARS-CoV-2 genetics. *medRxiv*, 2024. (Preprint)
- 2. Kathryn N Porter Starr, Marshall G Miller, Jessica T Wallis, **Qimin Zhang**, Kuncheng Song et al. Effects of Blueberry Consumption and Exercise on Gut Microbiome Composition in a Randomized Controlled Trial of Sedentary Older Adults With Overweight or Obesity. *Current Developments in Nutrition*, 8(Supplement\_2), 2024.
- 3. Qian Shi, **Qimin Zhang**, Mingfu Shao. Accurate assembly of multiple RNA-seq samples with Aletsch. *Bioinformatics*, 40(Supplement\_1): i307–i317, 2024.
- 4. Kuncheng Song, Hayden Brochu N, **Qimin Zhang**, Jonathan Williams D, Lakshmanan Iyer K. An in silico analysis of PCR-based monkeypox virus detection assays: a case study for ongoing clinical surveillance. *Viruses*, 15(12), 2327, 2023.
- 5. **Qimin Zhang**, Mingfu Shao. Transcript Assembly and Annotations: Bias and Adjustment. *PLoS Computational Biology*, 19(12): e1011734, 2023.
- 6. **Qimin Zhang**, Qian Shi, Mingfu Shao. Accurate assembly of multi-end RNA-seq data with Scallop2. *Nature Computational Science*, 2, 148-152, 2022.
- 7. **Qimin Zhang**, Nathaniel Kremer-Herman, Benjamin Tovar, Douglas Thain. Reduction of Workflow Resource Consumption Using a Density-based Clustering Model. 2018 IEEE/ACM Workflows in Support of Large-Scale Science (WORKS), pages. 1-9, Nov. 2018.
- 8. **Qimin Zhang**, Pei An, Shuquan Wang, Xiaoli Bai, Wei Zhang. Image-based Space Object Reconstruction and Relative Motion Estimation using Incremental Structure from Motion. 2018 IEEE CSAA Guidance, Navigation and Control Conference (CGNCC), Aug. 2018.
- 9. Yan Shi, Guoliang Wang, Jinglong Niu, **Qimin Zhang**, Maolin Cai et al. Classification of sputum sounds using artificial neural network and wavelet transform. *International Journal of Biological Sciences*, 14(8): 938–945, 2018.
- 10. **Qimin Zhang**, Jieru Zhao, Shuquan Wang. Design of Motion Control System for Frog-Inspired Bionic Hopping Robot. *International Conference on Mechatronics and Intelligent Robotics*, Pages: 502-509, Nov, 2017.
- 11. Jieru Zhao, Yang Li, **Qimin Zhang**, Zhongcai Pei. Research on Gait and Control of Bionic Hexapod Robot *Proceedings of the 2017 International Conference on Artificial Intelligence, Automation and Control Technologies*, Pages: 1-5, April, 2017.
- 12. **Qimin Zhang**, Zihe Liu, Jieru Zhao, Shuguang Zhang. Modeling and attitude control of Bi-copter. 2016 IEEE International Conference on Aircraft Utility Systems (AUS), Pages:172 176, Oct, 2016.

### Academic Services

30+ reviews for conferences and journals:

- Conference Reviewer : RECOMB 2021/2022/2023/2024, ISMB/ECCB 2020/2021/2022, WABI 2021, ACM-BCB 2020/2022, APBC 2020.
- Journal Reviewer: Heliyon, Biochemical Genetics.