# **HAOWEN ZHANG**

Georgia Institute of Technology  $\diamond$  Atlanta, GA 30332, United States (+1) 404 200 3022  $\diamond$  hwzhang@gatech.edu

#### **EDUCATION**

## Georgia Institute of Technology

08/2017 - Present

Ph.D. student in Computational Science and Engineering

## Shandong University

09/2013 - 06/2017

B.E., Hons, in Computer Science and Technology

#### RESEARCH EXPERIENCE

## Georgia Institute of Technology

Atlanta, GA

Graduate Research Assistant; Advisor: Dr. Srinivas Aluru

08/2017 - Present

- Working on long/short read algorithm design and implementation.
- Using high-performance architectures to accelerate sequencing data analysis.

## Tsinghua University & National Institutes for Food and Drug Control

Beijing, China

Visiting Student; Advisor: Dr. Tao Jiang and Dr. Tai Guo

01/2017 - 07/2017

- Worked on viral deep sequencing data analysis.
- Set up a pipeline to detect low-frequency variants using deep sequencing data of viruses.

#### **Shandong University**

Jinan, China

Research Assistant; Advisor: Dr. Weiquo Liu

08/2015 - 01/2017

- Proposed a new index data structure and two novel seeding algorithms for short read mapping.
- Accelerated edit distance computing on Intel Many Integrated Core Architecture (MIC).

#### The University of Hong Kong

Hong Kong

Research Intern; Advisor: Dr. Siu-Ming Yiu

08/2016

- 1/19 students accepted by HKU CS Research Internship Programme.
- Conducted a comprehensive evaluation of structural variation (SV) detection tools using simulated reads.
- Gave suggestions on developing a new SV calling algorithm.

#### **SKILLS**

- Programming: C/C++, Shell, Java, Python, LATEX
- Parallel computing: OpenMP, MPI, code vectorization

#### **PREPRINTS**

1. **Haowen Zhang**, Chirag Jain, and Srinivas Aluru. A comprehensive evaluation of long read error correction methods. *bioRxiv*, 2019. (submitted)

#### **PUBLICATIONS**

- 1. Chirag Jain<sup>†</sup>, **Haowen Zhang**<sup>†</sup>, Yu Gao, and Srinivas Aluru. On the complexity of sequence to graph alignment. In *International Conference on Research in Computational Molecular Biology (RECOMB)*. Springer, 2019. (†contributed equally; accepted)
- 2. Chirag Jain, Sanchit Misra, **Haowen Zhang**, Alexander Dilthey, and Srinivas Aluru. Accelerating sequence alignment to graphs. In *Parallel and Distributed Processing Symposium (IPDPS)*, 2019 *IEEE International*. IEEE, 2019. (accepted)
- 3. Yueyue Liu, Tai Guo, Qingchuan Yu, **Haowen Zhang**, Jialiang Du, Yunqi Zhang, Shengli Xia, Huan Yang, and Qihan Li. Association of human leukocyte antigen alleles and supertypes with immunogenicity of oral rotavirus vaccine given to infants in china. *Medicine*, 97(40):e12706, 2018
- 4. **Haowen Zhang**<sup>†</sup>, Yuandong Chan<sup>†</sup>, Kaichao Fan, Bertil Schmidt, and Weiguo Liu. Fast and efficient short read mapping based on a succinct hash index. *BMC bioinformatics*, 19(1):92, 2018. (†contributed equally)

#### SELECTED HONORS AND AWARDS

• Workshop on String Algorithms in Bioinformatics (StringBio) Travel Award	2018
• Workshop on the Future of Algorithms in Biology (FAB) Travel Fellowship	2018

### **TALKS**

• On the complexity of sequence to graph alignment, HotCSE, Gatech 03/2019

#### **SERVICE**

• External reviewer: IPDPS 2018

• Journal reviewer: Nucleic Acids Research