# **BRIAN HIE**

brianhie@mit.edu | github.com/brianhie | brianhie.com

### **EDUCATION**

# Massachusetts Institute of Technology, Cambridge, MA

# Electrical Engineering and Computer Science, Ph.D. candidate

2017-Present

• GPA: 5.0/5.0; Areas of concentration: Machine learning, algorithms, computational biology

Stanford University, Palo Alto, CA

# Computer Science, B.S. with Honors and Distinction

2012-2016

Minor in English Literature

• GPA: 3.9/4.0; Areas of concentration: Machine learning, computational biology, distributed systems

### RESEARCH AND WORK EXPERIENCE

# Massachusetts Institute of Technology, Cambridge, MA

### Computer Science and Artificial Intelligence Laboratory (CSAIL)

2017-Present

- Achieved privacy-preserving and scalable neural network training on large, cryptographically hidden data sets, applied to drug-target interaction prediction (<a href="https://github.com/brianhie/secure-dti">https://github.com/brianhie/secure-dti</a>).
- Applying novel algorithmic techniques for analyses of high-dimensional data to derive better biological insight from single cell RNA-sequencing data sets (https://github.com/brianhie/scanorama).

# Stanford University, Palo Alto, CA

### **Hunter Fraser Laboratory, Biology**

2013-2016

- Applied machine learning methods to predict associations between transcription factor binding and gene expression.
- Used statistical methods to explore the relationship between genotype and the formation and function of circular RNAs.

### Stanford University, Palo Alto, CA

### **Shakeosphere Project, Digital Humanities**

2016

• Applied graph-theoretic algorithms to understand the evolution of social networks of early modern English authors, printers, publishers, and booksellers.

#### **WORK EXPERIENCE**

Illumina, Inc., San Diego, CA

## **Machine Learning Intern**

2018

 Analyzed large genomic data sets by developing novel unsupervised and semi-supervised statistical machine learning methods.

#### Salesforce.com, Inc., San Francisco, CA

### Software Engineer, Cloud Infrastructure

2016-2017

• Developed algorithms and a distributed pipeline that scaled to petabytes of system log and metric data to generate accurate key performance indicators visible to top company leadership.

<ul> <li>Microsoft Corporation, Redmond, WA</li> <li>Software Engineering Intern, Azure Compute and Microsoft Research</li> <li>Investigated different distributed scheduling algorithms and their impact on availability on a simulated Azure data center.</li> </ul>	PAGE 2  2015  utilization and
<ul> <li>Synaptics, Inc., San Jose, CA</li> <li>Systems Architecture/Algorithms Intern</li> <li>Developed and implemented an algorithm for optimal touchscreen finger tracking minimum weight bipartite matching problem.</li> </ul>	2014 g by solving the
PUBLICATIONS  B. Hie, B. Bryson, and B. Berger.  "Panoramic stitching of heterogeneous single cell transcriptomic data."  bioRxiv (Preprint).	2018
<b>B. Hie</b> *, H. Cho*, and B. Berger. (*Equal contribution.)  "Realizing private and practical pharmacological collaboration."  Science, 362:6417 (2018).	2018
A.K. Tehranchi, <b>B. Hie</b> , M. Dacre, I.M. Kaplow, K.P. Pettie, P.A. Combs, and H.B. Fraser. "Fine-mapping cis-regulatory variants in diverse human populations." eLife (In Press).	2018
A.K. Tehranchi, M. Myrthil, T. Martin, <b>B. Hie</b> , D. Golan, and H.B. Fraser. "Pooled ChIP-seq links variation in transcription factor binding to complex disease risk." Cell, 165:3 (2016).	2016
AWARDS	

2016

2015

2014

2013

2013

2012

Hoefer Prize for Writing in the Major, Nominated, Stanford University

Lunsford Award for Oral Presentation, Nominated, Stanford University

President's Award for Academic Achievement, Stanford University

Boothe Prize for Excellence in Writing, Honorable Mention, Stanford University

Tau Beta Pi Engineering Honors Society

National Merit Scholarship Finalist