# **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

2019-Present

Electrical Engineering and Computer Science, M.S.

2017-2019

• GPA: 4.9/5.0; Areas of concentration: Algorithms, computational biology, machine learning, statistics

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: Computational biology, distributed systems, machine learning

#### RESEARCH AND WORK EXPERIENCE

Massachusetts Institute of Technology, Cambridge, MA

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

2017-Present

- Insightful and efficient geometric algorithms for single-cell biology (https://github.com/brianhie/scanorama and https://github.com/brianhie/geosketch).
- Cryptographically secure neural network training (https://github.com/brianhie/secure-dti).

#### Massachusetts Institute of Technology, Cambridge, MA

#### Teaching Assistant, Algorithms for Inference (6.438)

2019

• Graduate-level course on statistical inference with probabilistic graphical models. Responsible for preparing exams/assignments, leading discussion sections, and holding office hours.

#### Google LLC, Mountain View, CA

#### Artificial Intelligence/Machine Learning Resident, X – The Moonshot Factory

2019

Machine learning for early-pipeline moonshots.

Illumina, Inc., San Diego, CA

#### Machine Learning Intern, Bioinformatics

2018

• Statistical signal processing for genomics-based health monitoring.

Salesforce.com, Inc., San Francisco, CA

### Software Engineer, Cloud Infrastructure

2016-2017

Robust performance monitoring of globally distributed core application infrastructure.

#### Stanford University, Palo Alto, CA

## Hunter Fraser Laboratory, Biology

2013-2016

• Statistics and machine learning for computational genomics, including fundamental problems involving transcription factor binding, chromatin accessibility, and circular RNA structure.

Stanford	University	Dala Al	to C A
Stantord	University	'. Paio Ai	IO. CA

#### Shakeosphere Project, Digital Humanities

2016

• Graph-theoretic analysis of the social network of early modern authors and publishers.

#### Microsoft Corporation, Redmond, WA

## Software Engineering Intern, Azure Compute and Microsoft Research

2015

• Distributed scheduling algorithms and their impact on data center utilization and availability.

#### Synaptics, Inc., San Jose, CA

#### Systems Architecture/Algorithms Intern

2014

Algorithm design and implementation for embedded touchscreen firmware.

#### **PUBLICATIONS**

B. Hie\*, J. Peters\*, S. Nyquist\*, A. Shalek, B. Berger, and B. Bryson. (\*Equal contribution.)

"Coexpression enables multi-study cellular trajectories of development and disease." Annual Review of Biomedical Data Science, 3:1.

2020

B. Hie, H. Cho, B. Bryson, and B. Berger.

"Coexpression enables multi-study cellular trajectories of development and disease." bioRxiv, 10.1101/719088.

2020

R. Singh, A. Narayan, B. Hie, and B. Berger.

"Schema: A general framework for integrating heterogeneous single-cell modalities." bioRxiv, 10.1101/834549.

2019

**B. Hie\***, H. Cho\*, B. DeMeo, B. Bryson, and B. Berger. (\*Equal contribution.)

 $\hbox{\it ``Geometric sketching of single-cell data preserves transcriptional structure.''}$ 

Cell Systems, 8:6.

2019

B. Hie, B. Bryson, and B. Berger.

Nature Biotechnology, 37:6.

"Efficient integration of heterogeneous single-cell transcriptomes using Scanorama."

2019

A.K. Tehranchi, **B. Hie**, 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, 8:e39595.

2019

**B. Hie\***, H. Cho\*, and B. Berger. (\*Equal contribution.)

"Realizing private and practical pharmacological collaboration."

Science, 362:6417.

2018

A.K. Tehranchi, M. Myrthil, T. Martin, B. Hie, D. Golan, and H.B. Fraser.

"Pooled ChIP-seq links variation in transcription factor binding to complex disease risk." Cell, 165:3.

2016

## **PATENTS**

B. Hie, B. Berger, and H. Cho.

"Realizing private and practical pharmacological collaboration."

US Patent App. 16/235,606. 2019

# **AWARDS**

National Defense Science and Engineering Graduate (NDSEG) Fellow	2019-Present
Hoefer Prize for Writing in the Major, Nominated, Stanford University	2016
Tau Beta Pi Engineering Honors Society	2015-Present
Lunsford Award for Oral Presentation, Nominated, Stanford University	2014
Boothe Prize for Excellence in Writing, Honorable Mention, Stanford University	2013
President's Award for Academic Achievement, Stanford University	2013
National Merit Scholarship Finalist	2012