

# Dechao Tian

1600 Clifton Rd  
Atlanta, GA 30329

(412) 583-6800  
dechao.tian@gmail.com  
<https://tian-dechao.github.io>

## Research Interests

---

- Machine learning application in biomedicine and precision health
- Statistical modeling for data science
- Computational biology, systems biology, and genomics

## Current Employment

---

**Mathematical Statistician** 2019 - Present  
*Centers for Disease Control and Prevention, US*

- Develop solid statistical models to detect anomaly clusters using integrated whole-genome sequencing data and epidemiologic data in precision health

## Education and Training

---

**Postdoctoral Fellow in Computational Biology** 2015 - 2019  
*Carnegie Mellon University, US*

Advisor: Dr. Jian Ma

**Ph.D. in Statistics and Applied Probability** 2010 - 2015  
*National University of Singapore, Singapore*

Advisor: Dr. Kwok Pui Choi

**M.S. in Probability and Mathematical Statistics** 2009 - 2011  
*Northeast Normal University, China*

Advisor: Dr. Zhidong Bai

**B.S. in Mathematics and Applied Mathematics** 2005 - 2009  
*Northeast Normal University, China*

Advisor: Dr. Jianhua Guo

## Publications

---

\* represents co-first authors

10. **Tian D\***, Zhang R\*, Zhang Y, Zhu X, and Ma J. MOCHI enables discovery of heterogeneous interactome modules in 3D nucleome, **Genome Research** 30, 2 (2020).

- Featured cover article
  - Reported by Carnegie Mellon University. Exploring a genome’s 3D organization through a social network lens, **Phys.org** February 20, 2020
9. **Tian D**, Gu Q and Ma J. Identifying gene regulatory network rewiring using latent differential graphical models, **Nucleic Acids Research** 44, 17 (2016).
  8. Koh V, Cheung C, Li X, **Tian D**, Wang J.J, Mitchell P, Cheng C.Y, and Wong T.T. Retinal vein occlusion in a multi-ethnic asian population: the singapore epidemiology of eye disease study, **Ophthalmic Epidemiology** 23, 1 (2016).
  7. Chen L, Cheng C.Y, Choi H, Ikram M.K, Sabanayagam C, Tan G.S, **Tian D**, Zhang L, Venkatesan G, Tai E.S, Wang J.J, Mitchell P, Cheung C.M.G, Beuerman R.W, Zhou L, Chan E.C.Y, Wong T.T. Plasma metabonomic profiling of diabetic retinopathy, **Diabetes** 65, 4 (2016).
  6. Yam G.H.F, Yusoff N.Z.B.M, Kadaba A, **Tian D**, Myint H.H, Beuerman R.W, Zhou L, Mehta J.S. Ex vivo propagation of human corneal stromal “activated keratocytes” for tissue engineering, **Cell Transplantation** 24, 9 (2015).
  5. Chen L, Li J, Guo T, Ghosh S, Koh S.K, **Tian D**, Zhang L, Jia D, Beuerman R.W, Aebersold R, Chan E.C.Y, Zhou L. Global metabonomic and proteomic analysis of human conjunctival epithelial cells (IOBA-NHC) in response to hyperosmotic stress, **Journal of Proteome Research** 14, 9 (2015).
  4. Tong L, Zhou X, Jylha A, Aapola U, Liu D.N, Koh S.W, **Tian D**, Quah J, Uusitalo H, Beuerman R.W, Zhou L. Quantitation of 47 human tear proteins using high resolution multiple reaction monitoring (HR-MRM) based-mass spectrometry, **Journal of Proteomics** 115, (2015).
  3. Zhang S\*, **Tian D\***, Tran N.H, Choi K.P, and Zhang L.X. Profiling human cell-type specific transcription factor regulatory networks, **Nucleic Acids Research** 42, 20 (2014).
  2. Barathi V.A, Chaurasia S.S, Poidinger M, Koh S.K, **Tian D**, Ho C, Iuvone P.M, Beuerman R.W, Zhou L. Involvement of GABA transporters in atropine-treated myopic retina as revealed by iTRAQ quantitative proteomics, **Journal of Proteome Research** 13, 11 (2014).
  1. **Tian D** and Choi K.P. Sharp bounds and normalization of wiener-type indices, **PLOS ONE** 8, 11 (2013).

## Manuscripts in Preparation

---

2. **Tian D**, Zhu X, and Ma J. Diffdomain enables detection of reshaped chromatin domains using random matrix theory.

1. **Tian D** and Ma J. Exploiting the interplay between chromatin interactome and transcriptional regulatory network.

## Research Experience

---

4. **Carnegie Mellon University, US** 2015 - 2019  
*Postdoctoral Fellow. Advisor: Jian Ma, Ph.D.*
  - Develop advanced statistical and machine learning algorithms to study the interplay between three-dimensional genome architecture and gene regulation in disease
3. **National University of Singapore, Singapore** 2014 - 2015  
*Research Assistant. Advisor: Kwok Pui Choi, Ph.D.*
  - Develop machine learning algorithms to identify essential genes using regulatory networks
2. **Singapore Eye Research Institute (SERI), Singapore** 2011 - 2015  
*Research Collaborator with Lei Zhou, Ph.D.*
  - Provide statistical analysis and consultation for Proteomics & Microanalysis Laboratory
  - Collaborate with other members from SERI
1. **Center for Quantitative Medicine, Duke-NUS, Singapore** 2012 - 2015  
*Associate Member*

## Teaching Experience

---

2. **Centers for Disease Control and Prevention, US** 2019  
Lecture on *Introduction to R*
1. **National University of Singapore, Singapore** 2010 - 2014  
Teaching assistant for *Introduction to Statistics, Statistics for Life Science, and Probability*

## Presentations

---

2. **Centers for Disease Control and Prevention, DRIVE: an open source web application using Data Reduction for Interactive Visualization and Analysis of Epidemiologic and Genomic Data**, DataViz day, Atlanta, GA, 2019
1. **The Jackson Laboratory, Regulatory network inference and comparison**, Farmington, CT, 2018

## References

---

### **Anna Blackstock, Ph.D.**

Mathematical Statistician

Centers for Disease Control and Prevention

#### Contact Information

Division of Foodborne, Waterborne, and Environmental Diseases

Centers for Disease Control and Prevention

1600 Clifton Rd NE

Atlanta, GA 30329, USA

Tel:(404) 718-4720

hyp9@cdc.gov

### **Jian Ma, Ph.D.**

Associate Professor, Computational Biology

Carnegie Mellon University

#### Contact Information

School of Computer Science

Carnegie Mellon University

5000 Forbes Avenue

Pittsburgh, PA 15213, USA

Tel:(412) 268-2776

jianma@cs.cmu.edu

### **Kwok Pui Choi, Ph.D.**

Professor, Statistics

National University of Singapore, Singapore

#### Contact Information

National University of Singapore, 6 Science Drive 2, Singapore 117546

Tel:(+65) 6516-4387

stackp@nus.edu.sg