Dechao Tian

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Research Interests

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

Current Employment

Mathematical Statistician

2019 - Present

Centers for Disease Control and Prevention, US

• Surveillance, Information Management, and Statistics Office: research on anomaly cluster detection using heterogeneous and high dimensional time-to-event data

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

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

^{*} represents co-first authors

- 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 three-dimensional genome organization and its functional association with transcriptional regulation

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*

Presentation

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

References

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Associate Professor, Computational Biology Carnegie Mellon University

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National University of Singapore, Singapore

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Zhidong Bai, Ph.D.

Professor, Probability
Northeast Normal University, China

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