# **Dechao Tian**

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# **Current Employment**

ORISE Fellow 2019 - Present

Center for Disease Control and Prevention

• Surveillance, Information Management, and Statistics Office - Machine Learning: research on anomaly cluster detection

# **Past Employment**

#### Postdoctoral Research Associate

2015 - 2019

Carnegie Mellon University

• Computational Biology Department, Ma Laboratory - Computational Genomics: research on regulatory network, systems biology, and 3D chromatin organization

## **Education**

### Ph.D. in Statistics and Applied Probability

2010 - 2015

National University of Singapore, Singapore

Thesis title: Biological Network Analysis and Comparison

Thesis advisor: Dr. Kwok Pui Choi

#### M.S. in Probability and Mathematical Statistics

2009 - 2011

Northeast Normal University, China

Thesis title: Random Network Models' Discrimination

Thesis advisor: Dr. Zhidong Bai

### **B.S.** in Mathematics and Applied Mathematics

2005 - 2009

Northeast Normal University, China

# Manuscripts in Preparation

- 3. **Tian D**\*, Zhang R\*, Zhang Y, and Ma J, MOCHI enables discovery of heterogeneous interactome modules in 3D nucleome. bioRxiv, doi: https://doi.org/10.1101/542092. Preparing for resubmission to *Genome Research*.
- 2. **Tian D** $^*$ , Zhu X $^*$ , and Ma J, Diffdomains: model-based identification of significantly reshaped chromatin domains from Hi-C contact matrices between normal and disease conditions.

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

### **Publications**

- 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).
- Tian D and Choi K.P, Sharp bounds and normalization of wiener-type indices, PLOS ONE 8, 11 (2013).

Note:  $\star$  represents co-first authors.

# Research Experience

3. National University of Singapore, Singapore

2014 - 2015

Research Assistant. Advisor: Kwok Pui Choi, Ph.D.

• Develop model to identify essential genes by network motifs in 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 Assistant**

Introduction to R ST1131 Introduction to Statistics ST1232 Statistics for Life Science ST2131 Probability

## **Skills**

### Programming (Most used first)

R, Python, Unix/Linux bash script command, SAS programming language (Certified Base Programmer for SAS 9)

### Bioinformatics tools & databases

MEME Suite, bedtools, jucier, UCSC Genome Browser, ENCODE, TCGA

### **Presentation**

1. DRIVE: an open source web application using <u>Data Reduction for Interactive Visualization</u> and Analysis of <u>Epidemiologic</u> and Genomic Data, DataViz day, CDC 2019

### **Posters**

- 3. Advanced Molecular Detection day, CDC, 2019
- 2. Systems Biology, Global Regulation & Gene Expression, CSHL, 2018
- 1. 4DN Annual Meeting, NIH, 2017

## References

### Jian Ma, Ph.D.

Associate Professor, Computational Biology Carnegie Mellon University

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### Kwok Pui Choi, Ph.D.

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National University of Singapore, Singapore

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### Zhidong Bai, Ph.D.

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