Hong Zhang

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EDUCATION

Worcester Polytechnic Institute (WPI)

Worcester, MA

PhD, Statistics

08/2013 - 04/2018

Dissertation: Goodness-of-Fit Tests for Signal Detection and Genetic Association Studies.

Johns Hopkins University

Baltimore, MD

MS, Financial Mathematics

08/2011 - 12/2012

Wuhan University *BA and BS, Mathematics and Finance*

Wuhan, China 08/2007 – 06/2011

Professional Experience

Merck & Co., Inc.

Rahway, NJ

Senior Scientist

05/2018 – present

- Developed powerful statistical tests for genetic association analyses in Pharmacogenetics studies;
- Performed large-scale simulation studies to compare powers of different association tests;
- Identified genetic biomarkers that potentially could influence drug responses.

Publications and Preprints

- 1. <u>Hong Zhang</u>, Jiashun Jin and Zheyang Wu. "Distributions and Statistical Power of Optimal Signal Detection Methods in Finite Samples". *IEEE Transactions on Signal Processing*, vol. 68 (2020), pp. 1021-1033, doi:10.1109/TSP.2020.2967179.
- 2. <u>Hong Zhang</u>, Tiejun Tong, John Landers and Zheyang Wu. "TFisher Tests: Optimal and Adaptive Thresholding for Combining *p*-Values." *Annals of Applied Statistics*, vol. 14, no. 1 (2020), 178–201. doi:10.1214/19-AOAS1302.
- 3. <u>Hong Zhang</u>, Ni Zhao, Devan Mehrotra and Judong Shen. "Composite Kernel Association Test (CKAT) for SNP-set Joint Analysis of Genotype and Genotype-by-treatment Interaction in Pharmacogenetics Studies". *Bioinformatics*, pii: btaa125 (2020). doi:10.1093/bioinformatics/btaa125.
- 4. <u>Hong Zhang</u>, and Roger Lui. "Releasing Wolbachia-infected Aedes aegypti to prevent the spread of dengue virus: a mathematical study." *Infectious Disease Modelling*, vol. 5 (2020), pp. 142-160. doi:10.1016/j.idm.2019.12.004.
- Lan Luo, Judong Shen, <u>Hong Zhang</u>, Aparna Chhibber, Devan Mehrotra and Zheng-zheng Tang. "Multi-trait analysis of rare-variant association summary statistics using MTAR". *Nature Communications, Accepted*.
- 6. <u>Hong Zhang</u>, Judong Shen and Zheyang Wu. "An efficient and accurate approximation to the distribution of quadratic forms of Gaussian variables". *Submitted*. arXiv:2005.00905.
- 7. <u>Hong Zhang</u> and Zheyang Wu. "A Generalized Family of Fisher's Combination Tests and Their Null Distributions Under Correlation". *Submitted*. arXiv:2003.01286.

8. <u>Hong Zhang</u> and Zheyang Wu. "Generalized Goodness-Of-Fit Tests for Correlated Data". *Submitted*. arXiv:1806.03668.

REFEREE SERVICES

- 1. IEEE Internet of Things Journal
- 2. Computational Statistics and Data Analysis
- 3. PLOS Computational Biology
- 4. PLOS ONE
- 5. BMJ Open

Presentations

- 1. Advancing Pharmacogenomics Analysis of Drug Response in Early-Phase Clinical Trials. *JSM* 2019, *Denver, CO, July* 2019.
- 2. Novel Statistical Methods for Polygenic Prediction of Drug Responses in PGx Studies. *Princeton/Branchburg Seminar, Princeton, NJ,* November 2018.
- 3. A Novel Subgroup Identification Method: Exhaustive Search with Repeated Cross-validation. *ICSA Applied Statistics Symposium, New Brunswick, NJ,* June, 2018.
- 4. Novel P-value Combination Methods with Application to Genetic Association Studies. WPI Bioinformatics & Computational Biology Seminar, Worcester, MA, March, 2018.
- 5. Power of Optimal Detection Methods. *The 10th ICSA International Conference, Shanghai, China,* December, 2016.

Posters

- 1. CC-CKAT: A fast and powerful method for SNP set joint test of prognostic and predictive effects in pharmacogenomics studies. *American Society of Human Genetics* 2019, *Houston*, TX, October 2019.
- 2. Exhaustive Search for Patient Subgroup with Repeated Cross-validation., *Merck Data Science Symposium, Upper Gwynedd, PA,* May 2019.
- 3. Thresholding Fisher's P-Value Combination Method (TFisher) for Set-based Genetic Association Studies. *American Society of Human Genetics* 2018 *Conference, San Diego, CA,* October 2018.
- 4. CK-SKAT: Composite Kernel Machine Association Test for Biomarker Discovery in PGx Studies. *Institute for Mathematics and its Applications Workshop: Innovative Statistics and Machine Learning for Precision Medicine, Twin Cities, MN,* September 2017.

Software

Authored R packages SetTest, TFisher and CKAT.

• SetTest: Group Testing Procedures for Signal Detection and Goodness-of-Fit.

https://cran.r-project.org/package=SetTest

• *TFisher*: Thresholding Fisher's P-value Combination Methods.

https://cran.r-project.org/package=TFisher

• CKAT: Composite Kernel Association Test for Pharmacogenetics Studies.

https://cran.r-project.org/package=CKAT

Awards and Honors

- o PhD Research Achievement Prize, Math Department WPI, 2018.
- o Institute for Mathematics and its Applications PI Travel Fund, WPI, 2017.
- o Graduate Student Travel Fund, WPI, 2016.
- o Teaching Assistant of the Year, Math Department WPI, 2015.