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## Education

#### Ph.D. Candidate in Biostatistics

Expected: Apr 2022

Graduate School of Public Health, University of Pittsburgh

Pittsburgh, PA

- Research Interests: Bayesian data analysis, High-dimensional statistics, Statistical genetics, Meta-analysis, Subgroup analysis
- Cumulative GPA: 4.00 / 4.00

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

May 2017

School of Mathematical Sciences, Xiamen University

Xiamen, China

- Major GPA: 3.94 / 4.00 (top 2/71 in the program)
- Awards: Outstanding Undergraduate Thesis, Scholarship of Academic Excellence (2015-2017)

# Research Experience \_\_\_\_\_

### **Graduate Student Research**

July 2017 - Present

Department of Biostatistics, University of Pittsburgh; Advisor: George C. Tseng

Pittsburgh,PA

- Performing theoretical analysis of existing meta-analysis methods and Developing methodology for detecting sparse signal among multiple studies by combining multiple p-values
- · Developing methodology for outcome guided disease subtyping based on high throughput molecular data
- Performing single cell RNA-sequencing data analysis and developing methodology for subject classification model utilizing single cell RNA-sequencing data
- Preprocess bulk RNA-sequencing data using pipelines, such as TopHat2, HiSeq2, SAMtools and HTSeq. Perform downstream analysis of Bulk RNA-sequencing data, such as DE-analysis based on edgeR2, Limma and DESeq2 and pathway analysis
- · Perform variant identification and analysis for sequencing data based on GATK

Research Assistant July 2016 - May 2017

School of Mathematical Sciences, Xiamen University; Advisor: Wei Liang

Xiamen, China

Developed algorithm for kernel-based semi-supervised Bayesian quantile regression and applied it to cell lineage data for detection of abnormal asynchrony of division between sister cells

## **Publications**

- Fang Y, Tang S, Huo Z, Tseng GC, Park Y. Properties of adaptively weighted Fisher's method. arXiv preprint arXiv:1908.00583. 2019 Aug 1.ready to submit
- Grabosch S, Bulatovic M, Zeng F, Ma T, Zhang L, Ross M, Brozick J, **Fang Y**, Tseng G, Kim E, Gambotto A. Cisplatin-induced immune modulation in ovarian cancer mouse models with distinct inflammation profiles. *Oncogene*. 2019 Mar;38(13):2380.
- Liang W, Yang Y, Fang Y, Zhao Z, Hu J. Bayesian Detection of Abnormal Asynchrony of Division Between Sister Cells in Mutant Caenorhabditis elegans Embryos. *Journal of Computational Biology*. 2019 May 1;26(5):495-505.

## **Relevent Coursework**

• Bayesian Data Science, Longitudinal and Clustered Data Analysis, Statistical Estimation Theory, Linear Models, Applied Regression Analysis, Applied Mixed Model Analysis, Likelihood Theory and Applications, Asymptotic Methods in Statistics, High-Dimensional Statistics, Introduction to Genomics Analysis, High-Dimensional Data with Omics Application, Advanced R Computing, Nonparametic Theory, Introduction to Bioinformatics Programming in Python

## Skills \_

**Languages** English and Chinese

Statistical Packages R, STATA, SPSS and MATLAB

Programming Languages Python, C, shell scripting(Linux) and MFX