YUSI FANG

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EDUCATION

Graduate School in Public Health, University of Pittsburgh

Ph.D Candidate in Biostatistics Cumulative GPA: 4.0/4.0

Research Interest: High-dimensional Statistics, Meta-analysis, Statistical Genetics and Machine Learning

Methods

Relevant coursework: Bayesian Data Science, Applied Mixed Model Analysis, SAS, High-Dimensional Statistics, Introduction to Genomics Analysis, High-Dimensional Data with Omics Application, Advanced R Computing, Nonparametric Theory, Asymptotic Methods, Survival analysis

School of Mathematical Sciences, Xiamen University

Xiamen, China

Pittsburgh, PA

Expected: Apr.2022

B.S. in Mathematics and Applied Mathematics

Aug. 2013 - May 2017

Major GPA: 3.94 / 4.00 (top 2/71 in the program)

EXPERIENCE

Research Assistant

Jul. 2017 – Present

Department of Biostatistics, School of Public health, University of Pittsburgh Advisors: George Tseng and Zhao Ren

Pittsburgh, PA

- Developed methodology for combining p-values for dependent tests with heavy-tailed distributions
- Developed methodology for optimal detection of weak and sparse signals via adaptive Fisher's method
- Developed methodology for outcome-guided disease subtyping for high-dimensional omics data
- Developing methodology for uniformly optimal detection of signals under multiple scenarios via p-values combination
- Collaborated with biologists for data pre-processing and analysis of clinical and genetics data
- Laboratory server daily maintenance and management

Undergraduate Research Assistant

Jul. 2016 – May 2017

School of Mathematical Sciences, Xiamen University

Advisor: Wei Liang

Xiamen, China

• Developed algorithm for kernel-based semi-supervised Bayesian quantile regression with application to cell lineage data for the detection of abnormal asynchrony of division between sister cells

SELECTED RESEACH PROJECTS

Robust testing for combining dependent p-values

Department of Biostatistics, University of Pittsburgh

Oct. 2019 – Sep. 2020

- Developed a family of robust tests for combining dependent p-values via transformation of heavy-tailed distribution
- Theoretically proved the proposed family of tests enjoys asymptotic robustness under arbitrary dependency structure and optimality for detection of sparse signals
- Extensive simulations to demonstrate type-I error control and power of our methods comparing to other existing methods
- Application to a neuroticism GWAS application

Outcome-guided disease subtyping for high-dimensional omics data Department of Biostatistics, University of Pittsburgh

Jun. 2019 – Jul. 2020

- Developed unify latent generative model to perform outcome-guided disease subtyping for continuous clinical outcome implemented by EM algorithm, with simultaneously feature selection for omics data and latent subtype characterization constructed from omics data
- Modified the model for survival outcome by embedding the accelerated failure time model into our model
- Ran simulations to compare the performance of our model modified for survival outcome with other methods

Jan. 2020– Present

Optimal detection of weak and sparse signals via adaptive Fisher's method Department of Biostatistics, University of Pittsburgh

- Developed a novel adaptive Fisher's method for the detection of weak and sparse signals with improvement of computational efficiency
- Theoretically proved the proposed tests enjoys asymptotic optimality for detection of weak and sparse signals
- Implemented comprehensive simulations for finite sample power comparison between our proposed method and other methods

Optimal detection of signals under multiple scenarios via p-values combination

Department of Biostatistics, University of Pittsburgh

Jun. 2020– Present

- Proposed a novel p-values combination method via modification on Fisher's method
- Theoretically proved the proposed method enjoys asymptotic optimality for multiple scenarios (scenarios of weak and sparse signals and scenarios of dense signals)
- Comprehensive simulations to investigate finite sample performance of our proposed method comparing to other methods in both cases of weak and sparse or dense signals
- Developing algorithm based on importance sampling and cross-entropy method for efficient computation of our proposed method

Data analysis of women over 70 years old with clinically node negative breast cancer May. 2020–Sep. 2020

Magee-Womens Research Institute and Foundation, University of Pittsburgh Medical Center

- Implemented Cox-proportional hazards model for overall survival, disease free survival
- Implemented propensity score matching over selected baseline covariates and rerun the survival analysis on the propensity score matched cohort

TEACHING EXPERIENCE

Department of Biostatistics, University of Pittsburgh

• 2020 Spring, BIOST 2094 Advanced R Computing Teaching Fellow.

Publication

- Fang, Yusi & Tseng, George & Chang, Chung. (2020). Robust Testing for Combining Dependent P-values Using Transformation of Heavy-tailed Distribution. submitted to JASA
- Liu, Peng & Fang, Yusi & Ren, Zhao & Tang, Lu & Tseng, George. (2020). Outcome-Guided Disease Subtyping for High-Dimensional Omics Data. *submitted to JASA*
- Liu, Peng & Liu, Silvia & Fang, Yusi & Xue, Xiangning & Zou, Jian & Tseng, George & Konnikova, Liza. (2020). Recent Advances in Computer-Assisted Algorithms for Cell Subtype Identification of Cytometry Data. Frontiers in Cell and Developmental Biology. 8. 234. 10.3389/fcell.2020.00234.
- Lin, Chien-Wei & Chang, Lun-Ching & Ma, Tianzhou & Oh, Hyunjung & French, Beverly & Puralewski, Rachel & Mathews, Fasil & Fang, Yusi & Lewis, David & Kennedy, James & Mueller (Müller), Daniel J. & Marshe, Victoria & Jaffe, Andrew & Chen, Qiang & Ursini, Gianluca & Weinberger, Daniel & Newman, Anne & Lenze, Eric & Nikolova, Yuliya & Sibille, Etienne. (2020). Older molecular brain age in severe mental illness. Molecular Psychiatry. 1-11. 10.1038/s41380-020-0834-1.
- Grabosch, Shannon & Bulatovic, Mirna & Zeng, Feitianzhi & Ma, Tianzhou & Zhang, Lixin & Ross, Malcolm & Brozick, Joan & Fang, Yusi & Tseng, George & Kim, Eun & Gambotto, Andrea & Elishaev, Esther & Edwards, Robert & Vlad, Anda. (2019). Cisplatin-induced immune modulation in ovarian cancer mouse models with distinct inflammation profiles. Oncogene. 38. 10.1038/s41388-018-0581-9.
- Liang, Wei & Yuxiao Yang & Yusi Fang& Zhongying Zhao & Jie Hu. "Bayesian Detection of Abnormal Asynchrony of Division Between Sister Cells in Mutant Caenorhabditis elegans Embryos." Journal of Computational Biology 26, no. 5 (2019): 495-505.

TECHNICAL SKILLS

Languages: English and Chinese

Statistical Packages: R, STATA, SPSS and SAS Programming Languages: Python, C and LATEX