ZHIGUANG (CALEB) HUO

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https://caleb-huo.github.io

EDUCATION

University of Pittsburgh,

Pittsburgh, PA, US

• Ph.D. in Biostatistics,

April 2017

- Dissertation: Statistical integrative omics methods for disease subtype discovery
- GPA: 3.93/4.00
- Advisors: George C. Tseng, ScD and Yong Seok Park, PhD

• M.S. in Physics,

Apr 2012

- GPA: 3.86/4.00

Harbin Institute of Technology,

Harbin, Heilongjiang, China

• B.S. in Physics,

June 2011

- GPA: 90.43/100

RESEARCH INTEREST

My research interest lies in the intersection between statistical methodology and its applications to genomics and bioinformatics. I am particularly interested in genomic data integration, models and variable selection in high-dimensional data, graphical models, Bayesian methods, optimization and statistical computing. I have collaborated with biologists in the fields of cancer and psychiatry, analyzing a broad range of genomic data. These experiences motivate me to develop methodology and software that are practical, user-friendly and easy to use.

PROFESSIONAL EXPERIENCE

• Assistant Professor

July $2017 \sim Now$

- Department of Biostatistics, University of Florida

PUBLICATIONS

2017

- 1. Marianne Seney, **Zhiguang Huo**, Leon French, Rachel Puralewski, Joyce Zhang, David A Lewis, George Tseng, Etienne Sibille. (2017) Distinct molecular signatures of depression in men and women. (Submitted)
- 2. **Zhiguang Huo**, Chi Song, George C. Tseng. (2017) Bayesian latent hierarchical model for transcriptomic meta-analysis to detect biomarkers with clustered meta-patterns of differential expression signals. Submitted to *Annals of Applied Statistics* (under revision).
- 3. **Zhiguang Huo**, George C. Tseng. (2017) Integrative Sparse K-means with overlapping group lasso in genomic applications for disease subtype discovery. Submitted to Annals of Applied Statistics (In press).
- 4. Dominique Arion, **Zhiguang Huo**, John F. Enwright, John P. Corradi, George Tseng and David A. Lewis. (2017) Transcriptome alterations in prefrontal pyramidal neurons distinguish schizophrenia from bipolar and major depressive disorders. *Biological Psychiatry*,
- 5. SungHwan Kim, Dongwan Kang, **Zhiguang Huo**, Yongseok Park, George C. Tseng. (2017) Meta-analytic principal component analysis in integrative omics application. (under revision).

⁰Last modified: July 12, 2017

- 6. Abraham Apfel, **Zhiguang Huo**, George Tseng and Stewart J. Anderson (2017) Achieving Accurate and Stable Feature Selection via Sparse K-means. (submitted)
- 7. Enwright, John, **Zhiguang Huo**, Dominique Arion, John Corradi, Aiqing He, George Tseng, and David Lewis. (2017) Transcriptome alterations of prefrontal cortical parvalbumin neurons in schizophrenia. (Under revision).

2016

- 8. **Zhiguang Huo**, Ying Ding, Silvia Liu, Steffi Oesterreich, and George Tseng. Meta-Analytic Framework for Sparse K-Means to Identify Disease Subtypes in Multiple Transcriptomic Studies. *Journal of the American Statistical Association*, 111, no. 513 (2016): 27-42.
- 9. Zhu, Li, Ying Ding, Cho-Yi Chen, Lin Wang, **Zhiguang Huo**, SungHwan Kim, Christos Sotiriou, Steffi Oesterreich, and George C. Tseng. "MetaDCN: meta-analysis framework for differential co-expression network detection with an application in breast cancer." *Bioinformatics* (2016): btw788.

2015 and before

- 10. Silvia Liu, Wei-Hsiang Tsai, Ying Ding, Rui Chen, Zhou Fang, Zhiguang Huo, SungHwan Kim, Tianzhou Ma, Ting-Yu Chang, Nolan Michael Priedigkeit, Adrian V. Lee, Jianhua Luo, Hsei-Wei Wang, I-Fang Chung, George C. Tseng. (2015). Comprehensive evaluation of fusion transcript detection algorithms and a meta-caller to combine top performing methods in paired-end RNA-seq data. Nucleic Acids Research, 10.1093/nar/gkv1234.
- 11. Tiffany A. Katz, Serena G. Liao, Vincent J. Palmieri, Robert K. Dearth, Thushangi Pathiraja, **Zhiguang Huo**, Patricia Shaw, Sarah Small, Nancy E. Davidson, David G. Peters, George C. Tseng, Steffi Oesterreich, Adrian V. Lee. (2015) Targeted DNA Methylation Screen in the Mouse Mammary Genome Reveals a Parity-Induced Hypermethylation of IGF1R That Persists Long after Parturition. *Cancer Prevention Research* 8, no. 10 (2015): 1000-1009.
- 12. Yan P. Yu, Silvia Liu, **Zhiguang Huo**, Amantha Martin, Joel B. Nelson, George C. Tseng and Jian-Hua Luo. (2015) Genomic copy number variations in the genomes of leukocytes predict prostate cancer clinical outcomes. *PloS one*, 10(8):e0135982.
- 13. SungHwan Kim, Zhiguang Huo, YongSeok Park and George Tseng. (2015) MetaOmics: transcriptomic meta-analysis methods for biomarker detection, pathway analysis and other exploratory purposes. Book chapter in Integrating omics data: statistical and computational methods. Edited by George C. Tseng, Debashis Ghosh, Xianghong Jasmine Zhou. Cambridge University Press. Page 39-67.
- 14. Xingbin Wang, Dongwan Kang, Kui Shen, Chi Song, Shuya Lu, Lunching Chang, Serena G. Liao, Zhiguang Huo, Naftali Kaminski, Etienne Sibille, Yan Lin, Jia Li and George C. Tseng. (2012) A Suite of R Packages for Quality Control, Differentially Expressed Gene and Enriched Pathway Detection in Microarray Meta-analysis. *Bioinformatics*, 28:2534-2536.

Under preparation

- Zhiguang Huo, Shaowu Tang, YongSeok Park and George Tseng. (2017) Biomarker categorization and fast computing of adaptively weighted Fisher's method for meta-analysis in omics applications.
- YongSeok Park, **Zhiguang Huo**, Shaowu Tang and George Tseng. (2017) Asymptotic properties of adaptive weighted Fisher's method.
- Li Zhu, **Zhiguang Huo**, Tianzhou Ma, George Tseng. (2017) Bayesian indicator variable selection model with multi-layer overlapping groups.
- Tianzhou Ma*, **Zhiguang Huo***, Anche Kuo*, Xiangrui Zeng, Li Zhu, Ark Fang, Lin Wang, Chien-Wei Lin, Tanbin Rahman, Shuchang Liu, YongSeok Park, Sunghwan Kim, Jia Li, Lun-Ching Chang, Chi Song, George Tseng. (2017) MetaOmics a Comprehensive Software Suite

with Interactive Visualization for Transcriptomic Meta-Analysis. (*: co-first author).

- Circadian rhythms of gene expression in the human prefrontal cortex reveal distinct pattern between schizophrenia and control subjects. In preparation.
- Rachel Puralewski, **Zhiguang Huo**, George Tseng, Ryan Logan, Marianne Seney. (2017) Sex differences in basolateral amygdala transcriptome associated with effects of chronic stress. In preparation.
- Zhiguang Huo, Rachel Puralewski, George Tseng, Ryan Logan, Marianne Seney. (2017) Circuitwide transcriptional profiling reveals sex differences in coherence patterns across brain regions. In preparation.
- Nader Shaikh, MD, MPH1; Alejandro Hoberman, MD1; Judith M. Martin, MD; Megan Skae, RN; Linette Milkovtch; Andrew Nowalk, MD; Christi Mcelheny; Robert Hickey, MD; Timothy Shope, MD; Diana Kearney; Massoud Majd; Eglal Shalaby-Rana; George Tseng; John Alcorn; Jay Kolls; William Horne; Zhiguang Huo; Greg Lockhart; Hans Pohl. (2017) Host and bacterial markers that differ in children with cystitis and pyelonephritis. In preparation.
- Nader Shaikh, MD, MPH1; Alejandro Hoberman, MD1; Judith M. Martin, MD; Megan Skae, RN; Linette Milkovtch; Andrew Nowalk, MD; Christi Mcelheny; Robert Hickey, MD; Timothy Shope, MD; Diana Kearney; Massoud Majd; Eglal Shalaby-Rana; George Tseng; John Alcorn; Jay Kolls; William Horne; **Zhiguang Huo**; Greg Lockhart; Hans Pohl. (2017) Characteristics that differentiate children with false positive dips from children with true urinary tract infection. In preparation.
- Nader Shaikh, MD, MPH1; Alejandro Hoberman, MD1; Judith M. Martin, MD; Megan Skae, RN; Linette Milkovtch; Andrew Nowalk, MD; Christi Mcelheny; Robert Hickey, MD; Timothy Shope, MD; Diana Kearney; Massoud Majd; Eglal Shalaby-Rana; George Tseng; John Alcorn; Jay Kolls; William Horne; **Zhiguang Huo**; Greg Lockhart; Hans Pohl. (2017) Host RNA expression in children with UTI An exploratory analysis of potential markers for pyelonephritis and UTI. In preparation.
- George C. Tseng, **Zhiguang Huo** and Tianzhou Ma. (2017) Foundations for High-Throughput Omics Data Analysis: Methods, Theories and Applications. *Chapman & Hall/CRC*.

AWARD

Student Awards

Delta Omega Membership
 American Statistics Association (ASA) Pittsburgh chapter
 Student of the year
 Department of Physics, Harbin Institute of Technology
 National Scholarship of P.R. China.
 (Awarded to the top 2 students in my Bachelors degree.)

Travel Awards

• SAMSI Research Triangle Park, NC.

 Interface of Statistics and Optimization 	Feb 2017
- Optimization Summer School	Aug 2016
- Epigenetics Workshop	Mar 2015
- Beyond Bioinformatics Workshop	June 2014

TEACHING EXPERIENCE (UNIVERSITY OF PITTSBURGH)

Main Lecturer (teaching fellow)

 BIOST2094 - Advanced R Computing – (with Tianzhou Ma) BIOST2025 - Special Studies in Bayesian Data Analysis – (with George Tseng, Tianzhou Ma and Li Zhu) 	Spring 2017 Fall 2016
Guest Lecturer	
 BIOST2055 - Introductory high-throughput genomic data analysis I: data mining and applications Differential and isoform analysis of RNA-seq data BIOST2078 - Introductory high-throughput genomic data analysis II: 	Mar 2016
theories and algorithms - Reproducible research and parallel computing in R • BIOST2078 - Introductory high-throughput genomic data analysis II:	Dec 2015
theories and algorithms - Reproducible research	Dec 2014
Teaching Assistant	
 BIOST 2078 - Introductory high-throughput genomic data analysis II: theories and algorithms PHYS 0212 - Introduction to Laboratory Physics PHYS 0212 - Introduction to Laboratory Physics 	Fall 2014 Spring 2012 Fall 2011
PRESENTATIONS	
Poster and Oral Presentation	
 Poster, Dean's Day's competition, GSPH, University of Pittsburgh Circadian rhythms of gene expression in the human prefrontal cortex reveal distinct pattern between schizophrenia and control subjects 	April 2017
 Invited talk, University of Florida, Gainesville, FL Meta-analytic and integrative framework for sparse ¡i¿K¡/i¿-means to identify types. 	Feb 2017 disease sub-
 Poster, SAMSI optimization summer school, Research Triangle Park, NC Integrative Sparse K-means for disease subtype discovery using multi-level omics data. 	Aug 2016
 Poster, Pittsburgh ASA banquet, Pittsburgh, PA Integrative Sparse K-means for disease subtype discovery using multi-level omics data. 	Mar 2016
 Oral Presentation, JSM, Seattle, WA Meta-analytic framework for sparse K-means to identify disease subtypes in multiple transcriptomic studies. 	Aug 2015
 Poster, Pittsburgh ASA banquet, Pittsburgh, PA Meta-analytic framework for sparse K-means to identify disease subtypes in multiple transcriptomic studies. 	Apr 2015
 Oral Presentation, ENAR Conference, Miami, FL Meta-analytic framework for sparse K-means to identify disease subtypes in multiple transcriptomic studies. 	Mar 2015
 Poster, Dean's Day's competition, GSPH, University of Pittsburgh Discover and Characterize Invasive Lobular Breast Carcinoma Subtypes. 	Mar 2015
 Oral Presentation, ENAR Conference, Baltimore, MA Meta-analytic framework for sparse K-means to identify disease subtypes in multiple transcriptomic studies. 	Mar 2014
• Poster, Dean's Day's competition, GSPH, University of Pittsburgh	Mar 2014

- Meta-analytic framework for sparse $K\mbox{-means}$ to identify disease subtypes in multiple transcriptomic studies.

Seminar Talk

• Department of Biostatistics, University of Pittsburgh

Nov 2015

- How to use Latex to make slides

MEMBERSHIP

• Member of International Chinese Statistical Association	$Mar~2015 \sim Now$
• Member of Eastern North American Region International Biometric Society	$Oct\ 2013 \sim Now$
• Member of American Statistical Association	$Nov \ 2013 \sim Now$

HOBBIES

Running, skiing, other endurance activities.

Running Record

• Bank of America Chicago Marathon, Chicago, IL.	10/09/2016
• First National Bank Pittsburgh Triathlon (Sprint), Pittsburgh, PA	08/14/2016
• Dick's Sporting Goods Pittsburgh Marathon, Pittsburgh, PA.	05/01/2016
• First National Bank Pittsburgh Triathlon (Olympic standard), Pittsburgh, PA	08/09/2015
• Dick's Sporting Goods Pittsburgh Marathon, Pittsburgh, PA	05/03/2015
• Dick's Sporting Goods Pittsburgh Marathon, Pittsburgh, PA	05/04/2014
• Dick's Sporting Goods Pittsburgh Marathon, Pittsburgh, PA	05/05/2013