# 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**

†: co-first author; \*: corresponding author.

### 2019

- Nader Shaikh, Judith M Martin, Alejandro Hoberman, Megan Skae, Linette Milkovich, Christi McElheny, Robert W Hickey, Diana Kearney, Massoud Majd, Eglal Shalaby-Rana, George Tseng, John F Alcorn, Jay Kolls, Marcia Kurs-Lasky, William Horne, **Zhiguang Huo**, Timothy R Shope. (2019) Biomarkers that differentiate false positive urinalyses from true urinary tract infection. Pediatric Nephrology (Accepted)
- 2. **Zhiguang Huo**, Lei Yu, Jingyun Yang, Yun Zhu, David A. Bennett, Jinying Zhao. (2019) Brain and blood metabolome for Alzheimer's dementia: Findings from a targeted metabolomics analysis. *Neurobiology of aging* (Accepted)
- 3. Ali Zarrinpar, Ting-Yuan D Cheng, **Zhiguang Huo**. (2019) What Can We Learn About Drug Safety and Other Effects in the Era of Electronic Health Records and Big Data That We Would Not Be Able to Learn from Classic Epidemiology? *Journal of Surgical Research* (Accepted)

<sup>&</sup>lt;sup>0</sup>Last modified: November 23, 2019

- 4. Saunjoo L. Yoon, Oliver Grundmann, Joseph J. Williams, Samuel S. Wu, Christiaan Leeuwenburgh, Zhiguang Huo, and Thomas J. George Jr. Differential response to targeted acupuncture by gender in patients with gastrointestinal cancer cachexia: secondary analysis of a randomized controlled trial. Acupuncture in Medicine (2019): 0964528419873670.
- 5. Kristin Curry Greenwood, Jennifer Kirwin, **Zhiguang Huo**. (2019) Design and Implementation of the Health Professions Simulation Assessment, a Tool to Assess Students Perceptions of Simulation Experiences. *Journal of Acute Care Physical Therapy* (Accepted)
- Yenisel Cruz-Almeida, Puja Sinha, Asha Rani, Zhiguang Huo, Roger Benton Fillingim, Thomas C Foster. (2019) Epigenetic Aging is Associated with Clinical and Experimental Pain in Community-Dwelling Older Adults. *Molecular Pain* (Accepted)
- 7. **Zhiguang Huo**, Shaowu Tang, Yongseok Park, George Tseng. (2019) P-value evaluation, variability index and biomarker categorization for adaptively weighted Fisher's meta-analysis method in omics applications. *Bioinformatics* (Accepted)
- 8. Marianne L. Seney, Kelly Cahill, John F. Enwright III, Ryan W. Logan, **Zhiguang Huo**, Wei Zong, George Tseng, and Colleen A. McClung. (2019) Diurnal rhythms in gene expression in the prefrontal cortex in schizophrenia. *Nature Communications*. (Accepted)
- 9. Theodore Drashansky<sup>†</sup>, Eric Helm<sup>†</sup>, Zhiguang Huo, Upasana Parthasarathy, Ashley Zuniga, Jonathan Cho, Zhiwei Xu, Mohammad Uddin, Kyle Lorentsen, Safiehkhatoon Moshkani, Liang Zhou, Nina Curkovic, Preet Kumar, Xiaoping Luo, Dorina Avram. (2019) Bcl11b prevents fatal autoimmunity by promoting Treg cell program and constraining innate lineages in Treg cells. Science Advances (Accepted)
- Li Zhu, Zhiguang Huo, Tianzhou Ma, George Tseng. (2019) Bayesian indicator variable selection to incorporate multi-layer overlapping group structure in multi-omics applications. Annals of Applied Statistics (Accepted) (An earlier version won ENAR distinguished student paper award 2018)
- 11. **Zhiguang Huo\***, Li Zhu, Tianzhou Ma, Hongcheng Liu, Song Han, Daiqing Liao, Jinying Zhao and George Tseng\*. (2019) Two-way Horizontal and Vertical Omics Integration for Disease Subtype Discovery. *Statistics in Bioscience* (Accepted)
- 12. Song Han\*, **Zhiguang Huo**\*, Kathy Nguyen, Fanchao Zhu, Patrick W Underwood, Kari Basso, Thomas George, and Steven Hughes. (2019) The Proteome of Pancreatic Cancer-Derived Exosomes Reveals Signatures Rich in Key Signaling Pathways. *PROTEOMICS* (Accepted)
- 13. **Zhiguang Huo**, Chi Song\*, George C. Tseng\*. Bayesian latent hierarchical model for transcriptomic meta-analysis to detect biomarkers with clustered meta-patterns of differential expression signals. *Annals of Applied Statistics* 13, no. 1 (2019): 340-366. (An earlier version won ASA Biometrics Section JSM Travel Award 2018)
- 14. **Zhiguang Huo**, Yun Zhu, Lei Yu, Jingyun Yang, Philip De Jager, David A. Bennett, Jinying Zhao. DNA methylation variability in Alzheimer's Disease. *Neurobiology of aging* 76 (2019): 35-44.
- 15. Nader Shaikh, Judith M Martin, Alejandro Hoberman, Megan Skae, Linette Milkovich, Andrew Nowalk, Christi McElheny, Robert W Hickey, Diana Kearney, Massoud Majd, Eglal Shalaby-Rana, George Tseng, John F Alcorn, Jay Kolls, Marcia Kurs-Lasky, **Zhiguang Huo**, William Horne, Greg Lockhart, Hans Pohl, Timothy R Shope. (2019) Host and Bacterial Markers that Differ in Children with Cystitis and Pyelonephritis. *The Journal of Pediatrics*

#### 2018

- 16. Tianzhou Ma<sup>†</sup>, **Zhiguang Huo**<sup>†</sup>, Anche Kuo<sup>†</sup>, Li Zhu, Fang Zhou, Xiangrui Zeng, Chien-Wei Lin, Silvia Liu, Lin Wang, Tanbin Rahman, Lun-Ching Chang, Sunghwan Kim, Jia Li, Yongseok Park, Chi Song, Steffi Oesterreich, Etienne Sibille and George C. Tseng. (2019). MetaOmics Comprehensive Analysis Pipeline and Web-based Software Suite for Transcriptomic Meta-Analysis. *Bioinformatics* 35, no. 9 (2018): 1597-1599.
- 17. Marianne Seney, **Zhiguang Huo**, Kelly Cahill, Leon French, Rachel Puralewski, Joyce Zhang, Ryan W. Logan, George Tseng, David A Lewis, Etienne Sibille. Opposite molecular signatures of

- depression in men and women. Biological Psychiatry 84, no. 1 (2018): 18-27.
- 18. Cynthia R. Johnson, Kimberly Brown, Susan Hyman, Maria M. Brooks, Courtney Aponte, Lynne Levato, Brianna Schmidt, Victoria Evans, Zhiguang Huo, Roxanna Bendixen, Heather Eng, Theresa Sax, Tristram Smith. Parent Training for Feeding Problems in Children with Autism Spectrum Disorder: Initial Randomized Trial. Journal of pediatric psychology 44, no. 2 (2018): 164-175.
- 19. John Enwright, **Zhiguang Huo**, Dominique Arion, John P. Corradi, George Tseng, and David A. Lewis. Transcriptome alterations of prefrontal cortical parvalbumin neurons in schizophrenia. *Molecular psychiatry* 23, no. 7 (2018): 1606.
- 20. Kelly Cahill<sup>†</sup>, **Zhiguang Huo**<sup>†</sup>, George Tseng, Ryan W. Logan\*, Marianne L. Seney\*. Improved identification of concordant and discordant gene expression signatures using an updated rank-rank hypergeometric overlap approach. *Scientific Reports* 8.1 (2018): 9588.
- 21. Wang, Kai Wen, Xiangrui Zeng, Xiaodan Liang, **Zhiguang Huo**, Eric P. Xing, and Min Xu (2018). Image-derived generative modeling of pseudo-macromolecular structures-towards the statistical assessment of Electron CryoTomography template matching. *The British Machine Vision Conference*

### 2017

- 22. **Zhiguang Huo**, George C. Tseng. (2017) Integrative Sparse K-means with overlapping group lasso in genomic applications for disease subtype discovery. The Annals of Applied Statistics, 11(2), 1011-1039.
- 23. SungHwan Kim, Dongwan Kang, **Zhiguang Huo**, Yongseok Park, and George C. Tseng. Metaanalytic principal component analysis in integrative omics application. *Bioinformatics* 34, no. 8 (2017): 1321-1328.
- 24. Dominique Arion, **Zhiguang Huo**, John F. Enwright, John P. Corradi, George Tseng, and David A. Lewis. Transcriptome alterations in prefrontal pyramidal cells distinguish schizophrenia from bipolar and major depressive disorders. *Biological psychiatry* 82, no. 8 (2017): 594-600.

### 2016

- 25. **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.
- 26. 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* 33, no. 8 (2016): 1121-1129.

### 2015 and before

- 27. 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.
- 28. 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. 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.
- 29. 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.

- 30. 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.
- 31. 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.

# A١

Academic Awards	
<ul> <li>Department of Biostatistics, University of Florida</li> <li>Outstanding Teacher</li> </ul>	March 2019
Student Awards	
<ul> <li>Delta Omega Membership</li> <li>American Statistics Association (ASA) Pittsburgh chapter <ul> <li>Student of the year</li> </ul> </li> <li>Department of Physics, Harbin Institute of Technology</li> <li>National Scholarship of P.R. China. <ul> <li>(Awarded to the top 2 students in my Bachelors degree.)</li> </ul> </li> </ul>	April 2017 March 2016 May 2009
Travel Awards	
• 2018 ASA Biometrics Section JSM Travel Award, Vancouver, BC, Canada.	Aug 2018
• Objective Bayes meeting 2017 travel award, Austin, TX.	Dec 2017
<ul> <li>SAMSI, Research Triangle Park, NC.</li> <li>Interface of Statistics and Optimization</li> <li>Optimization Summer School</li> <li>Epigenetics Workshop</li> <li>Beyond Bioinformatics Workshop</li> </ul>	Feb 2017 Aug 2016 Mar 2015 June 2014
EACHING EXPERIENCE	
Lecturer, University of Florida	
<ul> <li>PHC6068 - Biostatistical computing</li> <li>PHC6937 - Frontiers in Biostatistics</li> <li>PHC6068 - Biostatistical computing</li> </ul>	Fall 2018 Feb 2018 Fall 2017
Lecturer, University of Pittsburgh	
<ul> <li>BIOST2094 - Advanced R Computing – (with Tianzhou Ma)</li> <li>BIOST2025 - Special Studies in Bayesian Data Analysis</li> <li>– (with George Tseng, Tianzhou Ma and Li Zhu)</li> </ul>	Spring 2017 Fall 2016
Guest Lecturer, University of Pittsburgh	
<ul> <li>BIOST2055 - Introductory high-throughput genomic data analysis I: data mining and applications         <ul> <li>Differential and isoform analysis of RNA-seq data</li> </ul> </li> <li>BIOST2078 - Introductory high-throughput genomic data analysis II:</li> </ul>	Mar 2016

theories and algorithms  $\mathrm{Dec}\ 2014$ - Reproducible research Teaching Assistant, University of Pittsburgh • BIOST 2078 - Introductory high-throughput genomic data analysis II: theories and algorithms Fall 2014 • PHYS 0212 - Introduction to Laboratory Physics Spring 2012 • PHYS 0212 - Introduction to Laboratory Physics Fall 2011 **PRESENTATIONS Invited Oral Presentation**  Division of Biostatistics, Albert Einstein College of Medicine, Bronx, NY Oct 2019 Statistical methods on omics data integration for disease subtype discovery. • Rush Alzheimer's Disease Center ROSMAP Investigator's Meeting, Chicago, IL April 2019 - Brain and blood metabolome for Alzheimer's dementia. • Department of Physiology and Functional Genomics at UF, Gainesville, FL October 2018 - Statistical methods on omics data meta-analysis and integration, for disease subtype discovery and differential expression analysis. • Joint Statistical Meetings, Vancouver, BC, Canada August 2018 Bayesian meta-analysis for biomarkers of meta-patterns. • International Chinese Statistical Association conference, New Brunswick, NJ June 2018 - Two-way Horizontal and Vertical Omics Integration for Disease Subtype Discovery. • Rush Alzheimer's Disease Center ROSMAP Investigator's Meeting, Chicago, IL May 2018 - Identifying variably methylated regions (VMRs) associated with AD. • International Indian Statistical Association conference, Gainesville, FL May 2018 - Two-way Horizontal and Vertical Omics Integration for Disease Subtype Discovery. • Department of Biostatistics, University of Florida, Gainesville, FL Feb 2017 - Meta-analytic and integrative framework for sparse K-means to identify disease subtypes. Contributed Poster and Oral Presentation • Poster, ENAR converence, Atlanta, GA Mar 2018 - p-value evaluation, variability index and biomarker categorization for adaptively weighted Fisher's meta-analysis method in omics applications • Poster, 2018 UF Stats Winter Workshop, University of Florida, Gainesville, FL Jan 2018 Bayesian latent hierarchical model for transcriptomic meta-analysis to detect biomarkers with clustered meta-patterns of differential expression signals • Poster, Objective Bayes meeting 2017, University of Texas, Austin, TX Dec 2017 - Bayesian latent hierarchical model for transcriptomic meta-analysis to detect biomarkers with clustered meta-patterns of differential expression signals • Poster, Dean's Day's competition, GSPH, University of Pittsburgh April 2017 - Circadian rhythms of gene expression in the human prefrontal cortex reveal distinct pattern between schizophrenia and control subjects Mar 2017 • Oral Presentation, ENAR, Washington, DC - Bayesian latent hierarchical model for transcriptomic meta-analysis to detect biomarkers with clustered meta-patterns of differential expression signals. • Poster, SAMSI optimization summer school, Research Triangle Park, NC Aug 2016 - Integrative Sparse K-means for disease subtype discovery using multi-level omics data.

- Reproducible research and parallel computing in R

• BIOST2078 - Introductory high-throughput genomic data analysis II:

• Poster, Pittsburgh ASA banquet, Pittsburgh, PA	Mar 2016
- Integrative Sparse K-means for disease subtype discovery using	
multi-level omics data.	
• Department of Biostatistics, University of Pittsburgh	Nov 2015
- How to use Latex to make slides	
• Oral Presentation, JSM, Seattle, WA	Aug 2015
- Meta-analytic framework for sparse $K$ -means to identify disease	
subtypes in multiple transcriptomic studies.	
• Poster, Pittsburgh ASA banquet, Pittsburgh, PA	$\mathrm{Apr}\ 2015$
- Meta-analytic framework for sparse $K$ -means to identify disease	
subtypes in multiple transcriptomic studies.	
• Oral Presentation, ENAR Conference, Miami, FL	Mar 2015
<ul> <li>Meta-analytic framework for sparse K-means to identify disease</li> </ul>	
subtypes in multiple transcriptomic studies.	
• Poster, Dean's Day's competition, GSPH, University of Pittsburgh	Mar 2015
<ul> <li>Discover and Characterize Invasive Lobular Breast Carcinoma Subtypes.</li> </ul>	
• Oral Presentation, ENAR Conference, Baltimore, MA	Mar 2014
- Meta-analytic framework for sparse K-means to identify disease	
subtypes in multiple transcriptomic studies.	
• Poster, Dean's Day's competition, GSPH, University of Pittsburgh	Mar 2014
- Meta-analytic framework for sparse K-means to identify disease	
subtypes in multiple transcriptomic studies.	

### CONFERENCE SERVICE

• ICSA, New Brunswick, NJ

June 2018

- Chair for Session New developments in microbiome sequencing data modeling and analysis.
- Chair for Session Modern Statistical Development for Biomedical Big data.

# REVIEWER SERVICE

# Statistical

Journal	Count
Journal of the American Statistical Association	1
Journal of the Royal Statistical Society Series C	1
Biometrics	1
Bioinformatics	1
BMC Bioinformatics	1

# **Biological**

Journal	Count
PLOS ONE	3
Scientific Reports	2
Journal of Alzheimer's Disease	2

### DEPARTMENT SERVICE

 $\bullet\,$  Member of PhD Qualifying Exam committee

 $Jan \ 2018 \sim Now$ 

- Department of Biostatistics, University of Florida

• Director of the Biostatistics Consulting Lab

Sep  $2017 \sim June \ 2019$ 

# PH.D. ADVISING

### Committee member

Name	Department	Graduation Time	First position
Shaojun Zhang	Statistics, UF	July 2018	84.51°
Satyajit Ghosh	Statistics, UF	July 2018	postdoc at Rutgers University
Jalalli Peyman	Statistics, UF	April 2019	Wells Fargo
Eric Helm	Anatomy and Cell Biology, UF	Active	
Zhongkai Wang	Biostatistics, UF	Active	
Tamal Ghosh	Statistics, UF	Active	
Minji Lee	Statistics, UF	Active	
Ruoyang Zhang	Statistics, UF	Active	

# **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

Name	Location	Distance	Date
HITS half ironman	Sarasota, FL		01/05/2020
Santa Fe Century	Gainesville, FL		10/26/2019
Las Olas International Triathlon	Fort Lauderdale, FL		03/10/2019
Five Points of Life Half Marathon	Gainesville, FL		02/16/2019
The Southernmost Marathon	Key West, FL		10/06/2018
Dick's Sporting Goods Pittsburgh Half Marathon	Pittsburgh, PA		05/06/2018
Five Points of Life Race Marathon	Gainesville, FL		02/18/2018
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	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