# ZHIGUANG (CALEB) HUO

2004 Mowry Road, 5th Floor CTRB, P.O. Box 117450, Gainesville, FL 32611-7450

 $(352)\text{-}294\text{-}5929 \diamond \texttt{zhuo@ufl.edu}$ 

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.

#### 2018

- 1. William Paden, Kelly Barko, Rachel Puralewski, Kelly Cahill, **Zhiguang Huo**, Micah Shelton, George Tseng, Ryan Logan, Marianne Seney. (2018) Developmental origin of sex differences in adult mood and in stress-induced transcriptional coherence across mesocorticolimbic circuitry. (Submitted)
- 2. **Zhiguang Huo**, Yun Zhu, Lei Yu, Jingyun Yang, Philip De Jager, David A. Bennett, Jinying Zhao (2018). Altered DNA methylation variability associated with Alzheimer's Disease. (Submitted)
- 3. 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. (Submitted)
- 4. 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 (2018). Parent Training for Feeding Problems in Children with Autism Spectrum Disorder: Initial Randomized Trial. (Submitted)

<sup>&</sup>lt;sup>0</sup>Last modified: September 8, 2018

- 5. Li Zhu, **Zhiguang Huo**, Tianzhou Ma, George Tseng (2018). Bayesian indicator variable selection to incorporate multi-layer overlapping group structure in multi-omics applications. (Submitted)
- 6. **Zhiguang Huo\***, Li Zhu, Tianzhou Ma, Hongcheng Liu, Song Han, Daiqing Liao, Jinying Zhao and George Tseng\* (2018). Two-way Horizontal and Vertical Omics Integration for Disease Subtype Discovery. (Submitted).
- 7. 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 and George C. Tseng. (2018). MetaOmics Comprehensive Analysis Pipeline and Web-based Software Suite for Transcriptomic Meta-Analysis. (Submitted)
- 8. **Zhiguang Huo**, Shaowu Tang, Yongseok Park, George Tseng. (2018) P-value evaluation, variability index and biomarker categorization for adaptively weighted Fisher's meta-analysis method in omics applications. (Submitted)
- 9. Marianne Seney, **Zhiguang Huo**, Kelly Cahill, Leon French, Rachel Puralewski, Joyce Zhang, Ryan W. Logan, George Tseng, David A Lewis, Etienne Sibille. (2018) Opposite molecular signatures of depression in men and women. *Biological Psychiatry*
- 10. **Zhiguang Huo**, Chi Song, George C. Tseng. (2018) Bayesian latent hierarchical model for transcriptomic meta-analysis to detect biomarkers with clustered meta-patterns of differential expression signals. *Annals of Applied Statistics* (Accepted).
- 11. Kelly Cahill<sup>†</sup>, **Zhiguang Huo**<sup>†</sup>, George Tseng, Ryan W. Logan\*, Marianne L. Seney\* (2018), Improved identification of concordant and discordant gene expression signatures using an updated rank-rank hypergeometric overlap approach. *Scientific Reports* 8.1 (2018): 9588.

#### 2017

- 12. SungHwan Kim, Dongwan Kang, **Zhiguang Huo**, Yongseok Park, George C. Tseng. (2017) Meta-analytic principal component analysis in integrative omics application. *Bioinformatics*, 1, 8.
- 13. 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. *Molecular Psychiatry*.
- 14. **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.
- 15. 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*.

#### 2016

- 16. **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.
- 17. 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

- 18. 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.
- 19. 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.
- 20. 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.
- 21. 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.
- 22. 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.

#### $\mathbf{AW}$

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VARD	
Student Awards	
• Delta Omega Membership	April 2017
<ul> <li>American Statistics Association (ASA) Pittsburgh chapter</li> <li>Student of the year</li> </ul>	March 2016
<ul> <li>Department of Physics, Harbin Institute of Technology</li> <li>National Scholarship of P.R. China.</li> </ul>	May 2009
(Awarded to the top 2 students in my Bachelors degree.)  Travel Awards	
• 2018 ASA Biometrics Section JSM Travel Award, Vancouver, BC, Canada.	Aug 2018
• Objective Bayes meeting 2017 travel award, Austin, TX.	Dec 2017
• SAMSI, Research Triangle Park, NC.	
<ul> <li>Interface of Statistics and Optimization</li> </ul>	Feb 2017
- Optimization Summer School	Aug 2016
- Epigenetics Workshop	Mar 2015
- Beyond Bioinformatics Workshop	June 2014
EACHING EXPERIENCE	
Lecturer, University of Florida	
• PHC6068 - Biostatistical computing	Fall 2018

• PHC6068 - Biostatistical computing	Fall 2018
• PHC6937 - Frontiers in Biostatistics	Feb 2018
• PHC6068 - Biostatistical computing	Fall 2017

#### Lecturer, University of Pittsburgh

• BIOST2094 - Advanced R Computing – (with Tianzhou Ma)	Spring 2017
• BIOST2025 - Special Studies in Bayesian Data Analysis	Fall 2016
- (with George Tseng, Tianzhou Ma and Li Zhu)	

#### Guest Lecturer, University of Pittsburgh

• BIOST2055 - Introductory high-throughput genomic data analysis I: data mining and applications

- Differential and isoform analysis of RNA-seq data

Mar 2016

• BIOST2078 - Introductory high-throughput genomic data analysis II: theories and algorithms Dec 2015 - Reproducible research and parallel computing in R • BIOST2078 - Introductory high-throughput genomic data analysis II: theories and algorithms 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 Fall 2011 • PHYS 0212 - Introduction to Laboratory Physics **PRESENTATIONS Invited Oral Presentation** • 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 • Oral Presentation, ENAR, Washington, DC Mar 2017 - 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. • 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	$\mathrm{Aug}\ 2015$
- Meta-analytic framework for sparse $K$ -means to identify disease	
subtypes in multiple transcriptomic studies.	
• Poster, Pittsburgh ASA banquet, Pittsburgh, PA	Apr 2015
- Meta-analytic framework for sparse K-means to identify disease	
subtypes in multiple transcriptomic studies.	3.5
• Oral Presentation, ENAR Conference, Miami, FL	Mar 2015
- Meta-analytic framework for sparse K-means to identify disease	
subtypes in multiple transcriptomic studies.	M 2015
• Poster, Dean's Day's competition, GSPH, University of Pittsburgh	Mar 2015
- Discover and Characterize Invasive Lobular Breast Carcinoma Subtypes.	M 0014
• Oral Presentation, ENAR Conference, Baltimore, MA	Mar 2014
- Meta-analytic framework for sparse K-means to identify disease	
subtypes in multiple transcriptomic studies.	3.5 001.4
• 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
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- Chair for Session New developments in microbiome sequencing data mode	
<ul> <li>Chair for Session Modern Statistical Development for Biomedical Big dat</li> </ul>	a.
REVIEWER SERVICE	
• Journal of the American Statistical Association (1)	
• Journal of the Royal Statistical Society Series C (1)	
• Bioinformatics (1)	
• BMC Bioinformatics (1)	
• PLOS ONE (2)	
• Scientific Reports (2)	
Scientific respons (2)	
DEPARTMENT SERVICE	
• Member of PhD Qualifying Exam committee	$Jan~2018 \sim Now$
<ul> <li>Department of Biostatistics, University of Florida</li> </ul>	
• Director of the Biostatistics Consulting Lab	Sep $2017 \sim Now$
<ul> <li>Department of Biostatistics, University of Florida</li> </ul>	
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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.	

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# Running Record

• 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 (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