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.

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)

⁰Last modified: September 11, 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[†], Zhiguang Huo[†], Anche Kuo[†], 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. (2018). MetaOmics Comprehensive Analysis Pipeline and Web-based Software Suite for Transcriptomic Meta-Analysis. Bioinformatics (Accepted).
- 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[†], **Zhiguang Huo**[†], 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.

AWARD

Student	Awards
Student	Awarus

• Delta Omega Membership	April 2017
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• American Statistics Association (ASA) Pittsburgh chapter

March 2016

Student of the vear

• Department of Physics, Harbin Institute of Technology

May 2009

- National Scholarship of P.R. China.

(Awarded to the top 2 students in my Bachelors degree.)

Travel Awards

• 2018 A	SA Biometrics Sec	tion JSM Travel A	ward, Vancouver,	, BC, Canada.	Aug 2018
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• Objective Bayes meeting 2017 travel award, Austin, TX.

Dec 2017

• 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

- Beyond Bioinformatics Workshop

TEACHING EXPERIENCE

Lecturer, University of Florida

• 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

Mar 2016

theories and algorithms - Reproducible research and parallel computing in R • BIOST2078 - Introductory high-throughput genomic data analysis II:	Dec	
theories and algorithms — Reproducible research	Dec :	2014
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 2	
• PHYS 0212 - Introduction to Laboratory Physics	Fall	2011
PRESENTATIONS		
Invited Oral Presentation		
• Joint Statistical Meetings, Vancouver, BC, Canada	August	2018
- Bayesian meta-analysis for biomarkers of meta-patterns.	T	2010
 International Chinese Statistical Association conference, New Brunswick, NJ Two-way Horizontal and Vertical Omics Integration for Disease Subtype Disc 	June :	2018
 Rush Alzheimer's Disease Center ROSMAP Investigator's Meeting, Chicago, IL Identifying variably methylated regions (VMRs) associated with AD. 	May 2	2018
• International Indian Statistical Association conference, Gainesville, FL	May	2018
- Two-way Horizontal and Vertical Omics Integration for Disease Subtype Disc		
 Department of Biostatistics, University of Florida, Gainesville, FL Meta-analytic and integrative framework for sparse K-means to identify disease subtypes. 	Feb 2	2017
Contributed Poster and Oral Presentation		
 Poster, ENAR converence, Atlanta, GA p-value evaluation, variability index and biomarker categorization for 	Mar	2018
adaptively weighted Fisher's meta-analysis method in omics applications		
 Poster, 2018 UF Stats Winter Workshop, University of Florida, Gainesville, FL Bayesian latent hierarchical model for transcriptomic meta-analysis to detect biomarkers with clustered meta-patterns of differential expression signals 	Jan :	2018
 Poster, Objective Bayes meeting 2017, University of Texas, Austin, TX Bayesian latent hierarchical model for transcriptomic meta-analysis to detect biomarkers with clustered meta-patterns of differential expression signals 	Dec 2	2017
 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 2	2017
 Oral Presentation, ENAR, Washington, DC Bayesian latent hierarchical model for transcriptomic meta-analysis to detect 	Mar	2017
 biomarkers with clustered meta-patterns of differential expression signals. Poster, SAMSI optimization summer school, Research Triangle Park, NC Integrative Sparse K-means for disease subtype discovery using 	Aug 2	2016
multi-level omics data. • Poster, Pittsburgh ASA banquet, Pittsburgh, PA — Integrative Sparse K-means for disease subtype discovery using	Mar	2016
multi-level omics data. • Department of Biostatistics University of Pittsburgh	Nov '	2015

 $-\,$ Differential and isoform analysis of RNA-seq data

 $\bullet\,$ BIOST2078 - Introductory high-throughput genomic data analysis II:

 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 – Meta-analytic framework for sparse K-means to identify disease	Apr 2015
subtypes in multiple transcriptomic studies. • Oral Presentation, ENAR Conference, Miami, FL — Meta-analytic framework for sparse K-means to identify disease	Mar 2015
 subtypes in multiple transcriptomic studies. 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 Meta-analytic framework for sparse K-means to identify disease subtypes in multiple transcriptomic studies. 	Mar 2014
CONFERENCE SERVICE	
• ICSA, New Brunswick, NJ — Chair for Session New developments in microbiome sequencing data mod — Chair for Session Modern Statistical Development for Biomedical Big da	
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) 	
DEPARTMENT SERVICE	
Member of PhD Qualifying Exam committee	$Jan \ 2018 \sim Now$
 Department of Biostatistics, University of Florida Director of the Biostatistics Consulting Lab Department of Biostatistics, University of Florida 	Sep 2017 \sim Now
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

• 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