# ZHIGUANG (CALEB) HUO

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(352)-294-5929  $\Leftrightarrow$  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

• Director of the Biostatistics Consulting Lab

Sep  $2017 \sim Now$ 

- Department of Biostatistics, University of Florida
- Assistant Professor

July  $2017 \sim Now$ 

- Department of Biostatistics, University of Florida

# **PUBLICATIONS**

#### 2017

- 1. 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*.
- 2. 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 and George C. Tseng. (2017). MetaOmics Comprehensive Analysis Pipeline and Web-based Software Suite for Transcriptomic Meta-Analysis. (Submitted) (\*: co-first author).
- 3. **Zhiguang Huo**, Shaowu Tang, Yongseok Park, George Tseng. (2017) P-value evaluation, variability index and biomarker categorization for adaptively weighted Fisher's meta-analysis method in omics applications. (Submitted)

<sup>&</sup>lt;sup>0</sup>Last modified: November 18, 2017

- 4. 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)
- 5. **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).
- 6. **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.
- 7. 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*,
- 8. SungHwan Kim, Dongwan Kang, **Zhiguang Huo**, Yongseok Park, George C. Tseng. (2017) Meta-analytic principal component analysis in integrative omics application. (accepted).

## 2016

- 9. **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.
- 10. 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

- 11. 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.
- 12. 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.
- 13. 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.
- 14. 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.
- 15. 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

• 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.
- George C. Tseng, **Zhiguang Huo** and Tianzhou Ma. (2017) Foundations for High-Throughput Omics Data Analysis: Methods, Theories and Applications. Chapman & Hall/CRC.

# A

AWARD	
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> </ul>	April 2017 March 2016 May 2009
<ul> <li>Department of Fhysics, narbin institute of Technology</li> <li>National Scholarship of P.R. China.</li> <li>(Awarded to the top 2 students in my Bachelors degree.)</li> </ul>	May 2009
Travel Awards	
<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
TEACHING EXPERIENCE (UNIVERSITY OF PITTSBURGH)	
Main Lecturer (teaching fellow)	
<ul> <li>BIOST2094 - Advanced R Computing – (with Tianzhou Ma)</li> <li>BIOST2025 - Special Studies in Bayesian Data Analysis – (with George Tseng, Tianzhou Ma and Li Zhu)</li> </ul>	Spring 2017 Fall 2016
Guest Lecturer	
<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  - 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	Fall 2014

## **PRESENTATIONS**

# Poster and Oral Presentation

• Poster, Dean's Day's competition, GSPH, University of Pittsburgh

• PHYS 0212 - Introduction to Laboratory Physics

• PHYS 0212 - Introduction to Laboratory Physics

April 2017

Spring 2012

Fall 2011

- Circadian rhythms of gene expression in the human prefrontal cortex reveal distinct pattern between schizophrenia and control subjects

• Invited talk, University of Florida, Gainesville, FL	Feb 2017
– Meta-analytic and integrative framework for sparse ¡i¿K¡/i¿-means to id	lentify disease sub-
types.  • Poster, SAMSI optimization summer school, Research Triangle Park, NC	Aug 2016
- Integrative Sparse K-means for disease subtype discovery using	1148 2010
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.	A . 001F
<ul> <li>Oral Presentation, JSM, Seattle, WA</li> <li>Meta-analytic framework for sparse K-means to identify disease</li> </ul>	Aug 2015
subtypes in multiple transcriptomic studies.	
• Poster, Pittsburgh ASA banquet, Pittsburgh, PA	Apr 2015
<ul> <li>Meta-analytic framework for sparse K-means to identify disease</li> </ul>	1
subtypes in multiple transcriptomic studies.	
• Oral Presentation, ENAR Conference, Miami, FL	Mar 2015
- Meta-analytic framework for sparse K-means to identify disease	
subtypes in multiple transcriptomic studies.	Mar 2015
<ul> <li>Poster, Dean's Day's competition, GSPH, University of Pittsburgh</li> <li>Discover and Characterize Invasive Lobular Breast Carcinoma Subtypes.</li> </ul>	Mar 2015
• Oral Presentation, ENAR Conference, Baltimore, MA	Mar 2014
<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 2014
- Meta-analytic framework for sparse K-means to identify disease	
subtypes in multiple transcriptomic studies.	
Seminar Talk	
• Department of Biostatistics, University of Pittsburgh	Nov 2015
<ul> <li>How to use Latex to make slides</li> </ul>	
REVIEWER SERVICE	
• PLOS ONE	
• Scientific Reports	
• Journal of the Royal Statistical Society Series C	
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
- Dielys Charting Coods Dittelyungh Monethon Dittelyungh DA	05/01/2016

• Dick's Sporting Goods Pittsburgh Marathon, Pittsburgh, PA.

• Dick's Sporting Goods Pittsburgh Marathon, Pittsburgh, PA

• Dick's Sporting Goods Pittsburgh Marathon, Pittsburgh, PA

• First National Bank Pittsburgh Triathlon (Olympic standard), Pittsburgh, PA

05/01/2016

08/09/2015

05/03/2015

05/04/2014