

Zhiguang Huo

CONTACT INFORMATION	4600 Bayard ST, APT 308 Pittsburgh, PA 15213	412-979-0592 zh18@pitt.edu
RESEARCH INTERESTS	Omics data integration, Machine learning and statistical learning, Genomics and epigenetics algorithms and applications, Bioinformatics, Optimization, Graphical model	
EDUCATION	University of Pittsburgh , Pittsburgh, PA, US Ph.D., Biostatistics, <i>Expected</i> : Summer 2017 <ul style="list-style-type: none">Thesis Topic: <i>Statistical integrative omics methods for disease subtype discovery and single cell methylation methods</i>GPA: 3.93/4.00Advisors: George C. Tseng, ScD and Yong Seok Park, PhD M.S., Physics, Apr 2012 <ul style="list-style-type: none">GPA: 3.86/4.00 Harbin Institute of Technology , Harbin, Heilongjiang, China B.S., Physics, June 2011 <ul style="list-style-type: none">GPA: 90.43/100	
RESEARCH EXPERIENCE	Research Assistant	Dec 2011 to present
	Department of Biostatistics, University of Pittsburgh Supervisor: George C. Tseng, ScD	
	Research Assistant	Aug 2014 to present
	Department of Biostatistics, University of Pittsburgh Supervisor: Yong Seok Park, PhD	
	Collaboration	Feb 2016 to present
	Department of Psychiatry, University of Pittsburgh Supervisor: George C. Tseng, ScD Collaborator: Colleen A. McClung, Ph.D, Marianne Seney, PhD, Ryan Logan, PhD	
	Collaboration	Oct 2015 to present
	Department of Pediatrics, University of Pittsburgh Supervisor: George C. Tseng, ScD Collaborator: Nader Shaikh , MD, MPH	
	Collaboration	Jul 2014 to Feb 2016
	Department of Psychiatry, University of Pittsburgh Supervisor: George C. Tseng, ScD Collaborator: David A. Lewis, MD, John F. Enwright, Ph.D., Dominique Arion, Ph.D.	
	Collaboration	Jul 2014 to Sep 2015
	Department of Pharmacology and Chemical Biology, Magee-Womens Research Institute Supervisor: George C. Tseng, ScD Collaborator: Steffi Oesterreich, PhD	
	Collaboration	Jan 2013 to Sep 2014

Department of Pathology,
University of Pittsburgh
Supervisor: George C. Tseng, ScD Collaborator: Jianhua Luo, MD, PhD, Yan Ping
Yu, MD, PhD

Collaboration

Dec 2012 to Nov 2013

Department of Anesthesiology and Neurobiology,
University of Pittsburgh
Supervisor: George C. Tseng, ScD Collaborator: William R. Lariviere, PhD

Collaboration

Mar 2012 to Aug 2012

Department of Environmental and Occupational Health,
University of Pittsburgh
Supervisor: George C. Tseng, ScD Collaborator: George D Leikauf, PhD

REFEREED
JOURNAL
PUBLICATIONS
(STATISTICAL)

1. **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.

REFEREED
JOURNAL
PUBLICATIONS
(APPLICATION)

1. 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.
2. 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 igflr which persists long after parturition. *Cancer Prevention Research*, pages canprevres-0178.
3. 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.
4. 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.

SUBMITTED
JOURNAL
PUBLICATIONS

1. **Zhiguang Huo**, Chi Song, George C. Tseng. (2016) 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 second round of review).
2. **Zhiguang Huo**, George C. Tseng. (2016) Integrative Sparse K -means for disease subtype discovery using multi-level omics data. Submitted to *Annals of Applied Statistics* (under second round of review).
3. Dominique Arion, **Zhiguang Huo**, John F. Enwright, John P. Corradi, George Tseng and David A. Lewis. Transcriptome alterations in prefrontal pyramidal neurons distinguish schizophrenia from bipolar and major depressive disorders. Submitted to *Biological Psychiatry*, (under second round of review).

	<ol style="list-style-type: none"> SungHwan Kim, Dongwan Kang, Zhiguang Huo, Yongseok Park, George C. Tseng. (2016) Meta-analytic principal component analysis. Submitted to <i>Annals of Applied Statistics</i> (under revision). Li Zhu, Ying Ding, Cho-Yi Chen, Lin Wang, Zhiguang Huo, SungHwan Kim, Christos Sotiriou, Steffi Oesterreich and George C. Tseng. (2016) MetaDCN: meta-analysis framework for differential coexpression network detection with an application in breast cancer. <i>Bioinformatics</i> (accepted).
PUBLISHED ABSTRACT	<ol style="list-style-type: none"> Oesterreich, S., Katz, T.A., Logan, G., Levine, K., Nagle, A., Huo, Z., Tseng, G.C., Rui, H., Lee, A.V. and Butler, L.M., 2016. Abstract PD2-08: Potential role of prolactin signaling in development and growth of the lobular subtype of breast cancer. <i>Cancer Research</i>, 76(4 Supplement), pp.PD2-08. Enwright, John, Dominique Arion, John Corradi, Aiqing He, Zhiguang Huo, George Tseng, and David Lewis. (2015) Transcriptome Profiling of Layer 3 Parvalbumin Neurons from the Dorsolateral Prefrontal Cortex of Schizophrenia Subjects. <i>NEUROPSYCHOPHARMACOLOGY</i>, vol. 40, pp. S400-S401.
MANUSCRIPT	<ol style="list-style-type: none"> John Enwright, Dominique Arion, Zhiguang Huo, George Tseng and David A. Lewis. Transcriptome alterations in layer 3 parvalbumin neurons in the dorsolateral prefrontal cortex in schizophrenia differ from those in layer 3 pyramidal cells. Zhiguang Huo, Shaowu Tang, YongSeok Park and George Tseng. Biomarker categorization and fast computing of adaptively weighted Fisher's method for meta-analysis in omics applications. MetaOmics software paper
BOOK AND BOOK CHAPTER	<ol style="list-style-type: none"> George C. Tseng, Zhiguang Huo and Tianzhou Ma. Foundations for High-Throughput Omics Data Analysis: Methods, Theories and Applications. Chapman & Hall/CRC. Zhiguang Huo, Shaowu Tang, 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.
AWARDS	<p>Student Awards – American Statistics Association (ASA) Pittsburgh chapter</p> <ul style="list-style-type: none"> Student of the year March 2016 <p>Travel Awards – SAMSI Research Triangle Park, NC</p> <ul style="list-style-type: none"> Optimization Summer School Aug 2016 Epigenetics Workshop Mar 2015 Beyond Bioinformatics Workshop June 2014 <p>Student Awards – Department of Physics, Harbin Institute of Technology</p> <ul style="list-style-type: none"> National Scholarship of P.R. China. May 2009 (Awarded to the top 2 students in my Bachelors degree.)

TEACHING
EXPERIENCE

Lecture

- Lecture on BOST2094 - Advanced R Computing Jan 2017
(6 lectures on Advanced R Computation)
- Lecture on BOST2025 - Special Studies in Bayesian Data Analysis Oct 2016
(4 lectures on Advanced Bayesian Computation)

Guest lecture

- Guest lecture on BOST2055 - Introductory high-throughput genomic data analysis I: data mining and applications Mar 2016
(Differential and isoform analysis of RNA-seq data)
- Guest lecture on BOST2078 - Introductory high-throughput genomic data analysis II: theories and algorithms Dec 2015
(Reproducible research and parallel computing in R)
- Guest lecture on BOST2078 - Introductory high-throughput genomic data analysis II: theories and algorithms Dec 2014
(Reproducible research)

Teaching assistant

- BOST 2078 - Introductory high-throughput genomic data analysis II: theories and algorithms Sep 2014 - Dec 2014
with George C. Tseng
Department of Biostatistics, University of Pittsburgh
- PHYS 0212 - Introduction to Laboratory Physics Jan 2012 - Apr 2012
with Russell J. Clark
Department of Physics and Astronomy, University of Pittsburgh
- PHYS 0212 - Introduction to Laboratory Physics Aug 2011 - Dec 2011
with Russell J. Clark
Department of Physics and Astronomy, University of Pittsburgh

PRESENTATIONS

Statistical Meetings

- Poster, SAMSI optimization summer school, Research Triangle Park, NC Aug 2016
Topic: Integrative Sparse K -means for disease subtype discovery using multi-level omics data.
- Poster, Pittsburgh ASA banquet, Pittsburgh, PA Mar 2016
Topic: Integrative Sparse K -means for disease subtype discovery using multi-level omics data.
- Oral Presentation, JSM, Seattle, WA Aug 2015
Topic: Meta-analytic framework for sparse K-means to identify disease subtypes in multiple transcriptomic studies.
- Poster, Pittsburgh ASA banquet, Pittsburgh, PA Apr 2015
Topic: Meta-analytic framework for sparse K-means to identify disease subtypes in multiple transcriptomic studies.
- Oral Presentation, ENAR Conference, Miami, FL Mar 2015
Topic: 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 2015
Topic: Discover and Characterize Invasive Lobular Breast Carcinoma Subtypes.
- Oral Presentation, ENAR Conference, Baltimore, MA Mar 2014
Topic: 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
Topic: Meta-analytic framework for sparse K-means to identify disease subtypes in multiple transcriptomic studies.

Department of Biostatistics, University of Pittsburgh

- Seminar talk (How to use Latex to make slides) Nov 2015

REFERENCES

George C. Tseng
 Professor Phone: 412-624-5318
 Department of Biostatistics (primary appointment)
 Department of Human Genetics
 Department of Computational & Systems Biology E-mail: ctseng@pitt.edu
 University of Pittsburgh

Yong Seok Park
 Assistant Professor Phone: 412-624-3028
 Department of Biostatistics E-mail: yongpark@pitt.edu
 University of Pittsburgh

David A. Lewis, MD
 Distinguished Professor of Psychiatry and Neuroscience,
 Thomas Detre Professor of Academic Psychiatry, Chair Phone: 412-246-6010
 Department of Psychiatry E-mail: lewisda@upmc.edu
 University of Pittsburgh

Steffi
 Colleen
 Luo

SKILLS

Computer Programming:

- gitbub (1 years),
- R (5 years),
- Linux (5 years),
- Matlab (2 year),
- Python (1 year),
- Java, C++, C (1 year).

RUNNING

- Bank of America Chicago Marathon, Chicago, IL 10/09/2016
[TIME 04:08:27](#)
- First National Bank Pittsburgh Triathlon (Sprint), Pittsburgh, PA 08/14/2016
[TIME 1:10:30](#) (only biking + running)
- Dick's Sporting Goods Pittsburgh Marathon, Pittsburgh, PA 05/01/2016
[TIME 3:58:46](#)
- First National Bank Pittsburgh Triathlon (Olympic standard), Pittsburgh, PA 08/09/2015
[Time 3:29:24](#)
- Dick's Sporting Goods Pittsburgh Marathon, Pittsburgh, PA 05/03/2015
[TIME 3:49:38](#)
- Dick's Sporting Goods Pittsburgh Marathon, Pittsburgh, PA 05/04/2014
[TIME 4:07:44](#)
- Dick's Sporting Goods Pittsburgh Marathon, Pittsburgh, PA 05/05/2013
[TIME 4:48:49](#)