SHANGHONG XIE

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Personal website: https://shanghongxie.github.io/

EDUCATION

Columbia University in the City of New York	2019
Ph.D. in Biostatistics	
Dissertation Title: Statistical Methods for Constructing Heterogeneous Biomarker Ne Advisor: Yuanjia Wang	etworks
University of Illinois at Urbana-Champaign M.S. in Statistics	2014
Sichuan University, Chengdu, China B.S. with highest honors in Statistics	2012
PROFESSIONAL EXPERIENCE	
• Assistant Professor, Department of Statistics, University of South Carolina	2024 – Present
• Member, Carolina Autism and Neurodevelopment (CAN) Research Center, University of South Carolina	2024 – Present
• Member, Institute for Mind and Brain (IMB), University of South Carolina	2024 – Present
• Associate Professor, <i>Department of Data Science</i> , School of Statistics, Southwestern University of Finance and Economics	2023 - 2024
• Assistant Professor, <i>Department of Data Science</i> , School of Statistics, Southwestern University of Finance and Economics	2021 - 2023
• Member, Center of Statistical Research, Southwestern University of Finance and Economics	2021 - 2024
• Postdoctoral Research Scientist, <i>Department of Biostatistics</i> , Mailman School of Publicolumbia University	lic Health, $2019 - 2021$
RESEARCH INTERESTS	

Statistical machine learning; network analysis; graphical model; precision medicine; functional data analysis; causal inference; mediation analysis; variable selection; high dimensional analysis; neuroimaging; biomarker; neurological and psychiatric diseases; mental health; COVID-19

AWARDS AND HONORS

• American Statistical Association (ASA) Mental Health Statistics Section Best Student Paper Award	2020
\bullet International Conference on Health Policy Statistics (ICHPS) Student Travel Award	2018
\bullet NYC Datathon (Data Science Competition) 1st Place Winning Team, among 1000+ participants	2017
• Columbia University Fellowship 2014 -	- 2017

PUBLICATIONS

An asterisk (*) is used to indicate corresponding author; An <u>underline</u> is used to indicate students under my supervision; A dagger (†) is used to indicate joint first author, equal contribution.

Peer-Reviewed Journal and Conference Articles

- McDonnell, E., Xie, S., Marder, K., Cui, F., and Wang, Y. (2024). Dynamic Undirected Graphical Models for Time-Varying Clinical Symptom and Neuroimaging Networks. Statistics in Medicine 43 (1), 4131-4147. (An earlier version won ASA Statistics in Imaging Section First Prize Student Paper Award)
- 2. Shi, B., Liu, Y., **Xie**, **S.**, Zhu, X., and Wang, Y. (2024). Network-Assisted Mediation Analysis with High-Dimensional Neuroimaging Mediators. Machine Learning for Healthcare Conference. *Proceedings of Machine Learning Research* 252, 1-22.
- 3. **Xie, S.***, Zeng, D., and Wang, Y. (2024). Identifying Temporal Pathways Using Biomarkers in the Presence of Latent Non-Gaussian Components. *Biometrics* 80 (2), ujae033.
- 4. Xie, S.* and Ogden, R. T. (2024). Functional Support Vector Machine. Biostatistics 25 (4), 1178-1194.
- 5. **Xie, S.***, Tarpey, T., Petkova, E., and Ogden, R. T. (2022). Multiple Domain and Multiple Kernel Outcome-weighted Learning for Estimating Individualized Treatment Regimes. *Journal of Computational and Graphical Statistics* 31 (4), 1375-1383.
- 6. **Xie, S.***, Wang, W., Wang, Q., Wang, Y., and Zeng, D. (2022). Evaluating Effectiveness of Public Health Intervention Strategies for Mitigating COVID-19 Pandemic. *Statistics in Medicine* 41 (9), 3820-3836.
- 7. COVID-19 Forecast Hub Consortium (2022). Evaluation of Individual and Ensemble Probabilistic Forecasts of COVID-19 Mortality in the US. *Proceedings of the National Academy of Sciences* 119 (15), e2113561119.
- 8. COVID-19 Forecast Hub Consortium (2022). The United States COVID-19 Forecast Hub Dataset. Scientific Data 9, 462.
- 9. Xie, S.*, McDonnell, E., and Wang, Y. (2022). Conditional Gaussian Graphical Model for Estimating Personalized Disease Symptom Networks. *Statistics in Medicine* 41 (3), 543-553. (An earlier version won ASA Mental Health Statistics Section Best Student Paper Award)
- 10. **Xie, S.***, Zeng, D., and Wang, Y. (2021). Integrative Network Learning for Multi-modality Biomarker Data. *Annals of Applied Statistics* 15 (1), 64-87.
- 11. Xie, S.*, Li, X., McColgan, P., Scahill, R. I., Zeng, D., and Wang, Y. (2020). Identifying Disease-associated Biomarker Network Features Through Conditional Graphical Model. *Biometrics* 76 (3), 995-1006. (Cover story of *Biometrics* September 2020 issue; An earlier version won the International Conference on Health Policy Statistics (ICHPS) Student Travel Award)
- 12. Goldman, J., Xie, S., Green, D., Naini, A., Mansukhani, M. M., and Marder, K. (2021). Predictive Testing for Neurodegenerative Diseases in the Age of Next-generation Sequencing. *Journal of Genetic Counseling* 30, 553-562.
- 13. Wang, Q., Xie, S., Wang, Y., and Zeng, D. (2020). Survival-Convolution Models for Predicting COVID-19 Cases and Assessing Effects of Mitigation Strategies. Frontiers in Public Health 8, 325. (Our model was used by the Center of Disease Control and Prevention (CDC) for COVID-19 Ensemble Forecast; Our forecasts website: https://github.com/COVID19BIOSTAT/covid19_prediction; CDC ensemble forecast website: https://www.cdc.gov/coronavirus/2019-ncov/covid-data/forecasting-us.html)

- 14. Li, X.†, Xie, S.†, McColgan, P., Tabrizi, S. J., Scahill, R. I., Zeng, D., and Wang, Y. (2018). Learning Subject-Specific Directed Acyclic Graphs with Mixed Effects Structural Equation Models from Observational Data. Frontiers in Genetics 9, 430.
- 15. Li, X., Xie, S., Zeng, D., and Wang, Y. (2018). Efficient l_0 -norm Feature Selection Based on Augmented and Penalized Minimization. Statistics in Medicine 37 (3), 473-486.
- 16. Avissar, M.†, **Xie**, **S.**†, Vail, B., Lopez-Calderon, J., Wang, Y., and Javitt, D. C. (2018). Meta-analysis of Mismatch Negativity to Simple versus Complex Deviants in Schizophrenia. *Schizophrenia Research* 19, 25-34.

COLLABORATIVE RESEARCH EXPERIENCE

40 students enrolled

New York State Psychiatric Institute (NYSPI) Role: Statistician	2016 - 2021
Columbia University Vagelos College of Physicians and Surgeons, Department of Neurology Role: Statistician	2019 - 2021
University College London, Institute of Neurology Role: Collaborator	2016 - 2021
Columbia University, School of Social Work Role: Statistician	2016
TEACHING EXPERIENCE	
University of South Carolina, Department of Statistics Instructor	2024 – Present
Classroom Teaching	
 STAT 530, Applied Multivariate Statistics and Data Mining STAT 509, Statistics for Engineers 	Spring 2025 Fall 2024
Computing Session Seminar	
• Computing Session 4: Using GitHub and Git	Fall 2024
Functional and Complex Data Analysis (FUNCODA) Working Group	2024 – Present
• Co-founder with Yuexuan Wu and William Consagra	
Southwestern University of Finance and Economics, School of Statistics $Instructor$	2022 - 2024
• Reading & Writing Scientific Articles: Advanced Topics in Machine Learning (<i>Graduat</i> 11 students enrolled	e) Spring 2024
• Machine Learning and Data Mining (<i>Undergraduate</i>) 72 students enrolled	Fall 2023
• An Introduction to Machine Learning (<i>Graduate</i>) 63 students enrolled	Fall 2023
\bullet Reading & Writing Scientific Articles: Advanced Topics in Machine Learning ($Graduat$ 12 students enrolled	e) Spring 2023
\bullet Machine Learning and Data Mining ($Undergraduate)$	Spring 2023

• Machine Learning and Data Mining (<i>Undergraduate</i>)	Fall 2022
 2 classes, 121 students enrolled in total Reading & Writing Scientific Articles: Advanced Topics in Machine Learning (Graduate 8 students enrolled 	e) Spring 2022
• Machine Learning and Data Mining (<i>Undergraduate</i>) 2 classes, 132 students enrolled in total	Spring 2022
Massive Open Online Course (MOOC: XuetangX) Co-Instructor (Graduate Course)	2023
Machine Learning	
Columbia University, Department of Biostatistics Co-Instructor (Graduate Course)	2015 - 2018
• Statistical Collaboration for Interdisciplinary Research	Spring 2018
Teaching Assistant (Graduate Course)	
• Randomized Clinical Trial II	Fall 2016
• Generalized Linear Models	Fall 2016
• Design of Medical Experiments	Spring 2016
• Analysis of Longitudinal Data	Fall 2015
University of Illinois at Urbana-Champaign, Department of Statistics Teaching Assistant (Graduate Course)	2013 - 2014
• Sampling and Categorical Data	Spring 2014
• Applied Multivariate Analysis	Fall 2013
MENTORING ACTIVITIES	
University of South Carolina, Department of Statistics PhD Dissertation Advisor	
• Sehun An: PhD student (co–advisor with Ting Fung Ma)	2025 - Present
PhD Dissertation Committee Member	
• Jiasong Duan: PhD student	2025 – Present
• Sijian Fan: PhD student	2024 – Present
Southwestern University of Finance and Economics, School of Statistics $Thesis\ Advisor$	
• Hexuan Song (Master thesis)	2022 - 2024
• Guishan Xiang (Master thesis)	2022 - 2024
• Angi Hua (Bachelor thesis)	2023 - 2024
• Yi Jiang (Bachelor thesis)	2023 - 2024
• Lu Li (Bachelor thesis)	2023 - 2024
• Lijuan Guo (Bachelor thesis)	2022 - 2023
• Silu Liu (Bachelor thesis)	2022 - 2023
• Qi Yang (Bachelor thesis)	2022 - 2023
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• Yangjie Yin (Bachelor thesis)	2022 - 2023
• Ruiying Li (Bachelor thesis)	2021 - 2022
• Mengjie Li (Bachelor thesis)	2021 - 2022
• Xing Wang (Bachelor thesis)	2021 - 2022
• Qiyu Wang (Bachelor thesis)	2021 - 2022
$A cademic\ Advisor$ • Supervised over 40 undergraduate students, received excellent evaluations	2022 - 2024
Columbia University, Department of Biostatistics $Mentor$	
• Zexi Cai (PhD student) PhD dissertation projects	2022 – Present

ACADEMIC SERVICES

Practicum project

Dissertation projects

• Bin Yang (Master student)

Research project

Editorial Board

• Frontiers in Public Health

2023 - Present

2021 - 2021

2019 - 2021

2019 - 2020

Journal and Conference Review

- Annals of Applied Statistics
- Biometrics
- Briefings in Bioinformatics
- BMC Bioinformatics
- Computational Statistics and Data Analysis
- Journal of the American Statistical Association

• Erin Mcdonnell (PhD student, Advisor: Professor Yuanjia Wang)

• Bihui Sun (Master student, Advisor: Professor Yuanjia Wang)

- Journal of Nonparametric Statistics
- Scientific Reports
- Statistics in Medicine
- Statistics in Biosciences
- Statistical Methods in Medical Research
- Stat
- IEEE International Conference on Bioinformatics and Biomedicine

Grant Review

• Full-time Member, 2024 – Present NIH/NINDS Huntington's Disease Biospecimen Resource Access Committee (HD-BRAC)

Conference Service

•	Member, Program Committee, 2024 – Pr	resent
	2025 International Chinese Statistical Association (ICSA) China Conference	
•	Chair, Invited session "New developments in the frontiers of precision medicine and data science", 7th International Conference on Econometrics and Statistics (EcoSta)	2024
•	Member, Student Paper Competition Committee for ASA Mental Health Statistics Section 2023 –	- 2024
•	Organizer, Invited session "Topics in healthcare and biostatistics", R conference in China	2023
•	Organizer, Invited session "Novel machine learning methods to advance precision medicine using big biomarker data", ICSA China	2023
•	Reviewer, Student Paper Competition, International Conference on Health Policy Statistics (ICHPS)	2022
•	Chair, Invited session "Statistical research in rapid response to COVID-19 pandemic: forecasts factors, therapeutics, and vaccine trials", Joint Statistical Meetings (JSM)	s, risk 2021
•	Chair, Topic-contributed session "Topics in clustering", JSM	2018
De	partmental and University Committees	

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University of South Carolina

• Member, Data Science and Analytics Programs Committee, Department of Statistics	2024 – Present
• Member, Faculty Hiring Committee, Department of Statistics	2024 - Present

Southwestern University of Finance and Economics

• Organizer, Departmental Seminar, School of Statistics	2023
• Member, Postdoctoral Evaluation Committee, School of Statistics	2022-2024
• Member, Master Thesis Committee, School of Statistics	2022-2024
• Member, Bachelor Thesis Committee, School of Statistics	2022 - 2024
• Member, Graduate Student Admissions Committee, School of Statistics	2022-2024
• Member, Curriculum Committee, School of Statistics	2021 - 2024

Memberships

- American Statistical Association (ASA)
- International Biometric Society, Eastern North American Region (ENAR)
- International Chinese Statistical Association (ICSA)
- Institute of Mathematical Statistics (IMS)
- New England Statistical Society (NESS)

SOFTWARE

R package 'APML0'

Co-maintainer

- Augmented and penalized minimization method for regularized linear, logistic, and Cox models with ℓ_0 penalty, flexible for ℓ_1 , ℓ_2 , and network type regularized regression
- Most intensive computation codes written in C++
- Available on CRAN: https://cran.r-project.org/web/packages/APMLO/index.html
- Downloaded 41,593 times as of 9/2023

R package 'Covariate-dependent-network'

Maintainer

- Estimate covariate-dependent networks through conditional Gaussian graphical model, in which both the mean and precision matrix depend on covariates
- Most intensive computation codes written in C++
- Available on GitHub: https://github.com/shanghongxie/Covariate-dependent-network

R package 'INL'

Maintainer

- Integrative network learning for multi-modality data
- Most intensive computation codes written in C++
- Available on GitHub: https://github.com/shanghongxie/INL

Matlab toolbox 'OWMKL'

Maintainer

- Outcome weighted multiple kernel learning (OWMKL) for estimating individualized treatment rules
- Available on GitHub: https://github.com/shanghongxie/OWMKL

R package 'FSVM'

Maintainer

- Functional support vector machine for classification and regression problems
- Available on GitHub: https://github.com/shanghongxie/FSVM

R package 'ICATemporalNetwork'

Maintainer

- Temporal causal network learning, adjusting for latent non-Gaussian components and separating the temporal network from the contemporaneous network
- Available on GitHub: https://github.com/shanghongxie/ICATemporalNetwork

PRESENTATIONS AND POSTERS

"Identifying Temporal Pathways Using Biomarkers in the Presence of Latent Non-Gaussian Components.", ICSA Applied Statistics Symposium, Storrs, CT, Invited session 6/2025

"Identifying Temporal Pathways Using Biomarkers in the Presence of Latent Non-Gaussian Components.", Lifetime Data Science Conference (LiDS), New York, NY, *Invited session* 5/2025

"Identifying Temporal Pathways Using Biomarkers in the Presence of Latent Non-Gaussian Components.", Eastern North American Region (ENAR), New Orleans, LA, Contributed session 3/2025

"Network-Assisted Mediation Analysis with High-Dimensional Neuroimaging Mediators.", ABCD Insights & Innovation Meeting (AIIM) Meeting, Bethesda, MD, Poster session 3/2025

"Identifying Temporal Pathways Using Biomarkers in the Presence of Latent Non-Gaussian Components.", Statistics Seminar, Department of Mathematical Sciences, Indiana University-Indianapolis, $Invited\ talk\ 2/2025$

"Identifying Temporal Pathways Using Biomarkers in the Presence of Latent Non-Gaussian Components." Functional and Complex Data Analysis (FUNCODA) Working Group, University of South Carolina 12/2024

"Identifying Temporal Pathways Using Biomarkers in the Presence of Latent Non-Gaussian Components."

Conference on Statistical Learning and Data Science (SLDS), California, Contributed session

11/2024

"Identifying Temporal Pathways Using Biomarkers in the Presence of Latent Non-Gaussian Components." International Conference on Econometrics and Statistics (EcoSta), Beijing, China, *Invited session* 07/2024

"Identifying Disease-associated Biomarker Network Features by Integrating Multi-modality Data." Department of Statistics, Virginia Tech, $Invited\ talk$ 08/2023

"Identifying Temporal Pathways Using Biomarkers in the Presence of Latent Non-Gaussian Components." Hangzhou International Conference on Frontiers of Data Science, Hangzhou, China, *Invited session* 08/2023

"Identifying Temporal Pathways Using Biomarkers in the Presence of Latent Non-Gaussian Components." ICSA China, Chengdu, China, Invited session 07/2023

"Identifying Disease-associated Biomarker Network Features through Graphical Models." Department of Epidemiology and Biostatistics, University of Georgia, *Invited talk* 04/2023

"Evaluating Effectiveness of Public Health Intervention Strategies for Mitigating COVID-19 Pandemic." New England Statistics Symposium (NESS), Hybrid, *Invited session* 05/2022

"Integrative Network Learning for Multi-modality Biomarker Data." Center for Statistical Science, Tsinghua University, $Invited\ talk$ 11/2021

"Identifying Temporal Pathways Using High-Dimensional Biomarkers." Joint Statistical Meetings (JSM), Virtual, Topic-contributed session 08/2021

"Evaluating Effectiveness of Public Health Intervention Strategies for Mitigating COVID-19 Pandemic." Columbia University Data Science Day, Oral poster session 04/2021

"Integrative Network Learning for Multi-modality Biomarker Data." Department of Biostatistics and Computational Biology & Del Monte Neuroscience Institute, University of Rochester, *Invited talk* 01/2021

"Integrative Network Learning for Multi-modality Biomarker Data." Division of Biostatistics, Department of Public Health Sciences, University of Virginia, *Invited talk* 10/2020

"Survival-Convolution Models for Predicting COVID-19 Cases and Assessing Effects of Mitigation Strategies."

Data Science Conference on COVID-19, Presentation session

08/2020

"Conditional Gaussian Graphical Model for Estimating Personalized Disease Symptom Networks." JSM, Virtual, Topic-contributed session 08/2020

"Integrative Network Learning for Multi-modality Biomarker Data." Eastern North American Region (ENAR), Virtual, Topic-contributed session 03/2020

"Statistical Methods for Constructing Heterogeneous Biomarker Networks." Division of Biostatistics, Department of Population Health, New York University School of Medicine, *Invited talk* 11/2019

"Integrative Network Learning for Multi-modality Biomarker Data." ICSA Applied Statistics Symposium, Raleigh, NC, *Invited session* 06/2019

"Estimating Heterogeneous Biomarker Networks and Their Effects on Disease Outcome." JSM, Vancouver, Canada, Topic-contributed session 07/2018

"Learning Subject-Specific Directed Acyclic Graphs (DAGs) from High-Dimensional Biomarker Data." Conference on Statistical Learning and Data Science (SLDS), New York, NY, Poster session 06/2018

"Learning Subject-Specific Directed Acyclic Graphs (DAGs) from High-Dimensional Biomarker Data." ENAR, Atlanta, GA, Poster session 03/2018

"High-dimensional Subject-Specific Network Analysis for Disentangling Genetic Mutation-Phenotype Pathways." ICHPS, Charleston, SC, Poster session 01/2018

CAREER DEVELOPMENT

• FreeSurfer Course	04/	/20	1'	7
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• Structural and Functional Connectivity via MRI

10/2016