# SHANGHONG XIE

Department of Statistics, University of South Carolina LeConte College, Columbia, SC 29208 SX2@mailbox.sc.edu

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

## **EDUCATION**

Columbia University in the City of New York Ph.D. in Biostatistics Dissertation Title: Statistical Methods for Constructing Heterogeneous Biomarker Net	2019
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, Department of Data Science, 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 Public Columbia University	e 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
• International Conference on Health Policy Statistics (ICHPS) Student Travel Award	2018
• 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.<sup>†</sup>, Xie, S.<sup>†</sup>, 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.

#### GRANT SUPPORT

## • Principal Investigator

01/2023 - 07/2024

"Learning Temporal Causal Network from Biomarker Time Series Data" National Natural Science Foundation of China (NSFC), Department of Mathematical and Physical Sciences, Grant No 12201511. Funding rate 17%

## • Principal Investigator

01/2022 - 12/2022

"Dynamic Network Learning Using Neuroimaging Data" Southwestern University of Finance and Economics Startup Grant

## COLLABORATIVE RESEARCH EXPERIENCE

## New York State Psychiatric Institute (NYSPI)

11/2016 - 08/2021

Role: Statistician

- Investigated the mental health impact of the COVID-19 pandemic on healthcare workers
- Developed design and statistical analysis plan to investigate the effects of cannabis use with varying concentrations on an NIH R01 grant proposal (funded)
- Designed a clinical trial for schizophrenia patients on an NIH grant proposal (funded)
- Provided statistical support (e.g., power analysis, statistical analysis) for clinicians and fellows
- Designed a study to compare a new short-form survey with a standard long-form for Alzheimer's disease, conducted power analysis and computed sensitivity, specificity, etc
- Conducted meta-analysis for a Schizophrenia study with clinicians

# Columbia University Vagelos College of Physicians and Surgeons, Department of Neurology

09/2019 - 08/2021

Role: Statistician

- Conducted statistical analysis to investigate the psychological impact of predictive testing for neurodegenerative diseases using next-generation sequencing panels
- Provided statistical support (e.g., power analysis)

## University College London, Institute of Neurology

11/2016 - 09/2021

Role: Collaborator

• Investigated brain connectivities

#### Columbia University, School of Social Work

01/2016 - 03/2016

Role: Statistician

• Conducted statistical analysis for a 25-year long period longitudinal cardiovascular disease study

#### TEACHING EXPERIENCE

University of South Carolina, Department of Statistics Instructor	2024 -	- Present
Classroom Teaching		
<ul> <li>STAT 530, Applied Multivariate Statistics and Data Mining</li> <li>STAT 509, Statistics for Engineers</li> </ul>	Sp	ring 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 $Instructor$	202	22 - 2024
$\bullet$ Reading & Writing Scientific Articles: Advanced Topics in Machine Learning ( $Graduate)$ 11 students enrolled	Sp	oring 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 ( $Graduate)$ 12 students enrolled	Sp	oring 2023
• Machine Learning and Data Mining ( <i>Undergraduate</i> ) 40 students enrolled	Sp	oring 2023
• Machine Learning and Data Mining ( <i>Undergraduate</i> ) 2 classes, 121 students enrolled in total		Fall 2022
$\bullet$ Reading & Writing Scientific Articles: Advanced Topics in Machine Learning ( $Graduate)$ 8 students enrolled	Sp	oring 2022
• Machine Learning and Data Mining ( <i>Undergraduate</i> ) 2 classes, 132 students enrolled in total	Sp	oring 2022
Massive Open Online Course (MOOC: XuetangX) Co-Instructor (Graduate Course)		2023
Machine Learning		
Columbia University, Department of Biostatistics Co-Instructor (Graduate Course)	201	15 - 2018
• Statistical Collaboration for Interdisciplinary Research	$\operatorname{Sp}$	oring 2018
Teaching Assistant (Graduate Course)		
• Randomized Clinical Trial II		Fall 2016
• Generalized Linear Models		Fall 2016
• Design of Medical Experiments	Sp	oring 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  Mentee	
• Sehun An: PhD student (co–mentor with Ting Fung Ma)	2024 – Present
PhD Dissertation Committee	
• Sijian Fan: PhD student	2024 - Present
• Jiasong Duan: PhD student	2025 - Present
Southwestern University of Finance and Economics, School of Statistics $Mentee$	
• Hexuan Song (Master thesis)	09/2022 - Present
• Guishan Xiang (Master thesis)	09/2022 - 06/2024
• Anqi Hua (Bachelor thesis)	10/2023 - 06/2024
• Yi Jiang (Bachelor thesis)	10/2023 - 06/2024
• Lu Li (Bachelor thesis)	10/2023 - 06/2024
• Lijuan Guo (Bachelor thesis)	01/2022 - 06/2023
• Silu Liu (Bachelor thesis)	01/2022 - 06/2023
• Qi Yang (Bachelor thesis)	01/2022 - 06/2023
• Yangjie Yin (Bachelor thesis)	01/2022 - 06/2023
• Ruiying Li (Bachelor thesis)	09/2021 - 06/2022
• Mengjie Li (Bachelor thesis)	09/2021 - 06/2022
• Xing Wang (Bachelor thesis)	09/2021 - 06/2022
• Qiyu Wang (Bachelor thesis)	09/2021 - 06/2022
$A cademic \ A dvisor$	
• Supervised over 40 undergraduate students, received excellent evaluations	09/2022 - 07/2024
Columbia University, Department of Biostatistics $Mentee$	
• Zexi Cai (PhD student) PhD dissertation projects	09/2022 - Present
• Bin Yang (Master student) Research project	01/2021 - 08/2021
• Erin Mcdonnell (PhD student, Advisor: Professor Yuanjia Wang) Dissertation projects	09/2019 - 08/2021
• Bihui Sun (Master student, Advisor: Professor Yuanjia Wang) Practicum project	11/2019 - 05/2020

#### ACADEMIC SERVICES

#### **Editorial Board**

• Frontiers in Public Health 2023 – Present

#### 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
- 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 Present 2025 International Chinese Statistical Association (ICSA) China Conference
- Chair, Invited session "New developments in the frontiers of precision medicine and data science", 2024 7th International Conference on Econometrics and Statistics (EcoSta)
- 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 2023 big biomarker data", ICSA China
- Reviewer, Student Paper Competition, 2022 International Conference on Health Policy Statistics (ICHPS)
- Chair, Invited session "Statistical research in rapid response to COVID-19 pandemic: forecasts, risk factors, therapeutics, and vaccine trials", Joint Statistical Meetings (JSM) 2021
- Chair, Topic-contributed session "Topics in clustering", JSM 2018

#### Departmental Committees

University of South Carolina

Member, Data Science and Analytics Programs Committee
 Member, Professional-Track Faculty Hiring Committee
 2024 - Present
 2024 - Present

Southwestern University of Finance and Economics

Organizer, Departmental Seminar, School of Statistics
 Member, Postdoctoral Evaluation Committee, School of Statistics
 01/2023 - 07/2023
 09/2022 - 07/2024

• Member, Master Thesis Committee, School of Statistics	03/2022 - 07/2024
• Member, Bachelor Thesis Committee, School of Statistics	03/2022 - 07/2024
• Member, Graduate Student Admissions Committee, School of Statistics	01/2022 - 07/2024
• Member, Curriculum Committee, School of Statistics	09/2021 - 07/2024

#### 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  $\ell_0$  penalty, flexible for  $\ell_1$ ,  $\ell_2$ , 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* 5/2025

"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 Mathematics, Indiana University-Indianapolis, *Invited session* 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

#### **Neuroimaging Short Courses**

Harvard University, Martinos Center for Biomedical Imaging

• FreeSurfer Course 04/2017

• Structural and Functional Connectivity via MRI 10/2016