# XIN XING

Department of Statistics Hutcheson Hall, 250 Drillfield Drive E-mail: xinxing@vt.edu Website: https://xin-xing.github.io/ Virginia Tech Blacksburg, VA, USA Phone: +1 (706) 207-9841

#### **APPOINTMENTS**

Assistant Professor		Department of Statistics, Virginia Tech	August 2020 - present	
EDUCATION				
Postdoc	Statistics	Harvard University Mentor: Dr. Jun S. Liu	2018 - 2020	
Ph.D.	Statistics	University of Georgia Advisor: Dr. Wenxuan Zhong Thesis topic: Statistical Methods with Applications in Epigen nomics and Neuroimaging	2013 - 2018 omics, Metage-	
M.S.	Statistics	University of Science and Technology of China (USTC) Advisor: Dr. Yaning Yang	2013	
B.S.	Statistics	University of Science and Technology of China (USTC)	2010	

#### **RESEARCH INTEREST**

<b>Bioinformatics</b> :	metagenomics, single cell, epigenomics, neuroimaging
Statistics:	minimax nonparametric testing, smoothing spline, dimension reduction, con-
	trolled variable selection
Machine learning:	computer vision, neural network compression, variable selection via neural network

## PAPERS

- [1] Yiwen Liu, Xin Xing, Wenxuan Zhong (2018). *Sufficient Dimension Reduction for Tensor Data*. Handbook of Big Data Analytics, Springer.
- [2] Xin Xing, Zuofeng Shang, Pang Du, Ping Ma, Wenxuan Zhong, Jun S. Liu (2019). *Minimax Non-parametric Two-sample Test*, Journal of the American Statistical Association: Theory and Methods, revised. arXiv:1911.02171
- [3] Xin Xing, Zhigen Zhao, Jun S. Liu (2019). *Controlling False Discovery Rate Using Gaussian Mirrors*, Journal of the American Statistical Association: Theory and Methods, revised. arXiv:1911.09761
- [4] Xin Xing, Yu Gui, Chengguang Dai, Jun S. Liu (2020). *Deep Gaussian Mirror for Controlled Variable Selection*, IEEE ICMLA, 2020.
- [5] Xin Xing, Meimei Liu, Wenxuan Zhong, Ping Ma. (2020). *Minimax Nonparametric Parallelism Test*, Journal of Machine Learning Research, 21(94):1-47.

- [6] Xin Xing, Meimei Liu, Weiping Zhang (2014). Joint Semiparametric Mean-Covariance Modeling by Moving Average Cholesky Decomposition for Longitudinal Data, Journal of University of Science and Technology of China, 43(8), 607-621.
- [7] Xin Xing, Long Sha, Pengyu Hong, Zuofeng Shang, Jun S. Liu (2020). Probabilistic Connection Importance Inference and Lossless Compression of Deep Neural Networks. International Conference on Learning Representations (ICLR).
- [8] Xin Xing, Jun S. Liu, Wenxuan Zhong (2018). MetaGen: reference-free learning with multiple metagenomic samples. Genome biology, 18 (1), 187, 2018.
- [9] Xin Xing, Jinjin Hu, and Yaning Yang (2014). *Robust minimum variance portfolio with L-infinity constraints*, Journal of Banking and Finance, 46, 107-117.
- [10] Xin Xing, Peng Zeng, Wenxuan Zhong (2019). Parsimonious Tensor Dimension Reduction, Statistica Sinica, revised.
- [11] Xin Xing, Di Xiao, Rui Xie, Wenxuan Zhong (2019). *Model-based Dictionary Learning: Sparse Coding beyond Gaussian Independent Model*, IEEE Transactions on Signal Processing, under review.
- [12] Wenxuan Zhong, Xin Xing, Kenneth Suslick (2015). Tensor Sufficient Dimension Reduction WIREs Computational Statistics, 7(3), 178-184.
- [13] Terry Ma, Di Xiao, Xin Xing (Corresponding author) (2019). *MetaBMF: A Scalable Binning Algorithm for Large-scale Reference-free Metagenomic Studies*. Bioinformatics.
- [14] Terry Ma, Xin Xing (2018). A Scalable Reference-Free Metagenomic Binning Pipeline. International Symposium on Bioinformatics Research and Applications, 79-83.
- [15] Ping Ma, Xinlian Zhang, Xin Xing, Jinyi Ma Michael Mahoney (2020). Asymptotic Analysis of Sampling Estimators for Randomized Numerical Linear Algebra Algorithms, International Conference on Artificial Intelligence and Statistics (AISTATS).
- [16] Chengguang Dai, Buyu Lin, Xin Xing and Jun S. Liu (2019). A Scale-free Approach for False Discovery Rate Control in Generalized Linear Models, Journal of the American Statistical Association: Theory and Methods, under review arXiv:2007.01237
- [17] Chengguang Dai, Buyu Lin, **Xin Xing** and Jun S. Liu (2019). *False Discovery Rate Control Via Data Splitting*, Journal of the Royal Statistical Society: Series B, under review.

#### PAPERS IN PREPARATION

- [1] Representation Learning of T-cell Receptor in cancer immunotherapy (with Songpeng Zu, Xiaole Liu and Jun S. Liu)
- [2] Bayesian Slicing Methods for Detecting Variable Dependency (with Jiexing Wu and Jun S. Liu)
- [3] Variable Hunting: New Promise in Binary Predictor Selection (with Wenxuan Zhong and Ping Ma)

#### **RESEARCH HIGHLIGHTS**

- Research is highlighted on the website of graduate school of UGA. Link: https://grad.uga.edu/index.php/xin-2017/
- Research is highlighted on the website of Pittsburgh Supercomputer Center. Link: https://www.psc.edu/161-news/psc-highlights/2414-bridges-reveals-diabetes-gut-microbe-links-2

#### GRANT

• NVIDIA GPU Grant for Accelerated Data Science, 2019.

#### HONORS AND AWARDS

- Best Senior PhD Student, UGA, May. 2018.
- Best Beginning PhD Student, UGA, Jul. 2014.
- Outstanding Dissertation For Bachelor's Degree, USTC, Jul. 2010.
- Third Prize of The Second Statistical Contest in Modeling, USTC, May. 2010.
- Third Prize of The First Statistical Contest in Modeling, USTC, May. 2009.
- Outstanding Student Scholarship (Silver Award), USTC, Oct. 2006.

#### TEACHING

- CMDA 3654 Lecture: Intro to Data Analytics & Visualization, Fall, 2020.
- STAT 221 Guest Lecturer : Monte Carlo Methods for Statistical Learning and Intro to Deep Learning, Harvard, Fall, 2019.
- STAT 8200 TA : Experimental Designs, University of Georgia, Fall, 2017.
- STAT 6210 TA : Introduction to Statistics I, University of Georgia, Fall, 2016.
- MSIT 3000 TA : Stat Analysis for Business I, University of Georgia, Spring, 2015.

### **CONFERENCES & INVITED TALKS**

- Neural Gaussian Mirror for Controlled Feature Selection in Neural Networks The International Conference on Learning Representations (ICLR), Online, April, 2020 (Invited talk)
- To Knockoff or To Disturb? Controlled Variable Selection in Regression Problems Department of Statistics, Virginia Tech, VA, Feb. , 2020 (Invited talk)
- PCII: Probabilistic Connection Importance Inference for Neural Network Lossless Compression International Chinese Statistical Association (ICSA), NJ, 2019 (Invited talk)
- Minimax Nonparametric Test for Density Comparison Department of Mathematical Science, IUPUI, IN, Mar 2018 (Invited talk)
- Reference-free Learning with Multiple Metagenomic Samples Department of Statistics, Harvard University, MA, Feb 2017 (Invited talk)
- Identifying Nonparallel Differentially Methylated DNA Regions University of Science and Technology of China, Hefei, China, Jan. 2017 (Invited talk)
- Identifying Nonparallel Differentially Methylated DNA Regions Fudan University, Shanghai, China, Jan. 2017 (Invited talk)
- Minimax Nonparametric Test for Density Comparison The Georgia Statistics Day Conference, GA, Oct 2015 (Invited talk)
- Nonnegative Matrix Factorization for Metagenomic Deconvolution Algorithm of Threat Detection Conference, CO, Sep 2014 (Invited talk)
- Nonnegative Matrix Factorization for Metagenomic Deconvolution Department of Automation, Tsinghua University, Beijing, China, Jun. 2014 (Invited talk)

#### Software

• Parallelism: R package "Parallelism" is to provide a nonparametric testing approach to test whether the spatial or temporal signals in treatment and control groups are parallel or not. Link: https://github.com/BioAlgs/Parallelism

- MetaMat: a fast algorithm for large-scale reference-free metagenomic studies. The pipeline outputs all binned species in multiple metagenomic samples and their estimated relative abundances. Link: https://github.com/didi10384/MetaBMF
- MetaGen: a statistically based algorithm to simultaneously identify microbial species and estimate their abundances in multiple metagenomic samples without using any reference genome. Link: https://github.com/BioAlgs/MetaGen

## **PROFESSIONAL SERVICE**

Referee Service, Journal of the American Statistical Association	2018-present
Referee Service, Electronic Journal of Statistics	2018-present
<ul> <li>Organizer, ICSA 2018 Applied Statistics Symposium</li> </ul>	2018
• Referee Service, IEEE Transactions on Computational Biology and Bioinformatics	2018-present
Referee Service, Statistical Analysis and Data Mining	2018-present
Referee Service, Frontiers in Genetics	2016-present
Organizer, The Georgia Statistics Day Conference, Athens, GA	Oct. 2016
Member, American Statistical Association	Since 2015
<ul> <li>Member, International Chinese Statistical Association</li> </ul>	Since 2015
<ul> <li>Judge, Georgia Science and Engineering Fair</li> </ul>	2015