## Jiangiao Wang

Information

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EMPLOYMENT Harvard University

Department of Biostatistics

Postdoctoral Research Fellow, 2022 - present

Advisor: Xihong Lin

EDUCATION University of Pennsylvania

Ph.D., Biostatistics, 2017 - 2022

Advisor: Hongzhe Li

University of Pennsylvania

M.S., Biostatistics, 2017 - 2019

Renmin University of China

B.S., Statistics, 2013 - 2017

**PUBLICATIONS** 

Wang, J., Li, S., and Li, H. (2023). A Regression-based Approach to Robust Estimation and Inference for Genetic Covariance, Journal of the American Statistical Association (Theory and Methods), Accepted

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Wang, J., Wang, W. and Li, H. (2022). Sparse block signal detection and identification for shared cross-trait association analysis. The Annals of Applied Statistics, 16(2), 866-886.

Wang, J. and Li, H. (2022). Estimation of genetic correlation with summary association statistics. Biometrika, 109(2), 421-438.

Xu, K., Wang, J., Pan, R. and Wang, H., 2019. Photographic diary: a new estimation approach to PM 2.5 monitoring. Statistics and Its Interface, 12(3), 387-395.

Chen, Y., Wang, J., Chubak, J., and Hubbard, R. A. (2019). Inflation of type I error rates due to differential misclassification in EHR-derived outcomes: Empirical illustration using breast cancer recurrence. *Pharmacoepidemiology and drug safety*, 28(2), 264-268.

Deo, R., Dubin, R.F., Ren, Y., Murthy, A.C., **Wang, J.**, Zheng, H., Zheng, Z., Feldman, H., Shou, H., Coresh, J. and Grams, M. (2023). Proteomic cardiovascular risk assessment in chronic kidney disease. European Heart Journal, 44(23), 2095-2110.

Dubin, R., Deo, R., Ren, Y., Wang, J., Zheng, Z., Shou, H., Go, A., Parsa, A., Lash, J., Rahman, M., Hsu, C., Weir, M., Chen, J., Anderson, A., Grams, M., Surapaneni, A., Coresh, J., Lee, H., Kimmel, P., Vasan, R., Feldman, H., Segal, M., Ganz, P. (2023). Proteomics of CKD progression in the chronic renal insufficiency cohort. Nature Communication, Accepted.

Hsu, C. Y., Yang, W., Parikh, R. V., Anderson, A. H., Chen, T. K., Cohen, D. L., He, J., Mohanty, M. J., Lash, J. P., Mills, K. T., Muiru, A. N., Parsa, A., Saunders, M. R., Shafi, T., Townsend, R. R., Waikar, S. S., **Wang, J.**, Wolf, M., Tan, T. C., Feldman, H. I., ... CRIC Study Investigators (2021). Race, Genetic Ancestry, and Estimating Kidney Function in CKD. *The New England journal of medicine*, 385(19), 1750–1760. (2022 Top 10 Clinical Research Achievement Awards).

PREPRINTS & Miao, Z., Wang, J., Park, K., Kuang, D., Kim, J. (2023). Model-based compound Working paper hypothesis testing for snATAC-seq data with PACS, bioRxiv.

Wang, J., Cui, Y. and Su, W. (2023). Two-sample comparison via unsupervised random forest. *Submitted*.

Wang, J. and Lin, X. Variance estimation with similarity encoding. Working paper.

Wang, J. and Li, H. Large-scale shared-signal detection and identification. Working paper.

Talks & Presentations

Contributed talk at JSM 2018: 'Integrating Data from GWAS and EQTL by Estimating Genetic Relatedness'.

Poster at JSM 2020: 'Robust Estimation of Genetic Covariance in High-Dimensional Generalized Linear Model'.

Contributed talk at JSM 2021: 'A Unified Approach to Robust Inference for Genetic Covariance'.

TEACHING EXPERIENCE Teaching assistant, 2018 Fall, BSTA 620 Probability

Teaching assistant, 2019 Fall, BSTA 620 Probability

PROFESSIONAL

Organizer of the 9th China-R Conference

ACTIVITIES

Session chair of the Biostatistics session in 15th China-R Conference

Referee Service Biometrics, Journal of the American Statistical Association

References

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