

Wei Zhang

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EDUCATION

University of Miami <i>Ph.D. in Biostatistics Advisor: Chen, X. Steven, Ph.D.</i> <i>Dissertation: Integrative Multi-Omics Analysis Using Multivariate Random Forest</i>	Miami, FL 08/2024
The George Washington University <i>M.S. in Statistics</i>	Washington, DC 05/2019
State University of New York at Binghamton <i>B.S. in Economics Analysis & Double Majors: Actuarial Math</i>	Binghamton, NY 05/2017

RESEARCH INTERESTS

Multi-omics Integration, Random Forests, Variable Selection, Meta-analysis, Biomarker Detection, Subtype Clustering, Statistical Genomics, Epigenetics, Neurodegenerative Disease and Cancers

RESEARCH EXPERIENCE

Postdoctoral Associate <i>University of Miami Translational Statistical Bioinformatics Lab</i>	Miami, FL 09/2024-Present
<ul style="list-style-type: none">Develop and implement advanced computational and machine learning methods for the analysis of large-scale omics data, including genomics, transcriptomics, proteomics, and epigenomicsDesign and develop innovative bioinformatics software and statistical tools for analyzing single-cell sequencing and spatial transcriptomics dataConduct computational analyses to identify biomarkers and therapeutic targets using multi-omics data integrationDevelop and apply machine learning models for predictive analytics in biomedical researchMaintain and optimize computational clusters and cloud computing environments to support large-scale data analysisPublish in refereed journals in collaboration with the principal investigatorContribute to basic and applied research activities, including authorship of scientific publications, technical and agency reports, or patent preparation	
Graduate Research Assistant <i>University of Miami Translational Statistical Bioinformatics Lab</i>	Miami, FL 05/2022-08/2024
<ul style="list-style-type: none">Collaborated with a diverse team to research and analyze genomic data for association studies, biomarker discoveries, and disease predictions in late-onset Alzheimer's Disease, triple-negative breast cancer, and colorectal cancerPublished multiple research papers contributing to the field of biomarker detection and disease predictionDemonstrated proficiency in R programming for comprehensive statistical analysis, handling diverse genomic data types, including RNA-seq, DNA methylation, and clinical dataSupported in drafting and editing grant proposals, ensuring clarity and alignment with project objectivesDeveloping an advanced R package for comprehensive DNA methylation data analysis	

PUBLICATIONS AND PREPRINTS

Key: * Indicates corresponding authors.

1. **Zhang W***, Huang H, Wang L, Lehmann BD, Chen XS* (2025) An Integrative Multi-Omics Random Forest Framework for Robust Biomarker Discovery. *GigaScience*, In review [[preprint](#)] [[code](#)]
2. Lukacsovich D, Zambare W, Wu C, Huang H, **Zhang W**, Kim MJ, Alvarez J, Bercz A, Paty, PB, Romesser PB, Wang L, Smith JJ*, Chen XS* (2025) Integrating Tumor and Organoid DNA Methylation Profiles Reveals Robust Predictors of Chemotherapy Response in Rectal Cancer (2025) [[preprint](#)]
3. **Zhang W**, Wu C, Huang H, Bleu P, Zambare W, Alvarez J, Wang L, Paty, PB, Romesser PB, Smith JJ*, Chen XS* (2025) Enhancing chemotherapy response prediction via matched colorectal tumor-organoid gene expression analysis and network-based biomarker selection. *Translational Oncology*, 52, 102238. ISSN: 1936-5233 [[paper](#)] [[code](#)]
4. **Zhang W**, Young JI, Gomez L, Schmidt MA, Lukacsovich D, Kunkle B, Chen XS, Martin ER, Wang L* (2024) Blood DNA Methylation Signature for Incident Dementia: Evidence from Longitudinal Cohorts. *Alzheimer's & Dementia*, In Press [[preprint](#)] [[code](#)]
5. **Zhang W**, Lukacsovich D, Young JI, Gomez L, Schmidt MA, Martin ER, Kunkle BW, Chen X, O'Shea DM*, Galvin JE*, Wang L* (2024). DNA Methylation Signature of a Lifestyle-based Resilience Index for Cognitive Health. *Alzheimer's Research & Therapy*, In review [[preprint](#)] [[code](#)]
6. **Zhang W**, Young JI, Gomez L, Schmidt MA, Lukacsovich D, Varma A, Chen XS, Kunkle B, Martin ER, Wang L* (2024) Critical evaluation of the reliability of DNA methylation probes on the Illumina MethylationEPIC BeadChip microarrays *Epigenetics*, 19(1) [[paper](#)] [[code](#)]
7. Lukacsovich D, Deirdre O'Shea, Huang H, **Zhang W**, Young JI, Chen XS, Dietrich ST, Kunkle B, Martin ER, Wang L* (2024) MIAMI-AD (Methylation in Aging and Methylation in AD): an integrative knowledgebase that facilitates explorations of DNA methylation across sex, aging, and Alzheimer's disease. *Database*, 2024, baae061 [[paper](#)] [[miami-ad.org](#)]
8. **Zhang W**, Young JI, Gomez L, Schmidt MA, Lukacsovich D, Varma A, Chen XS, Martin ER, Wang L* (2023) Distinct CSF biomarker-associated DNA methylation in Alzheimer's disease and cognitively normal subjects. *Alzheimer's Research & Therapy*, 15: 78 [[paper](#)] [[code](#)]
9. **Zhang W**, Li E, Wang L, Lehmann BD*, Chen XS* (2023) Transcriptome meta-analysis of triple-negative breast cancer response to neoadjuvant chemotherapy. *Cancers*, 2023; 15(8):2194 [[paper](#)] [[code](#)]
10. Silva TC, **Zhang W**, Young JI, Gomez L, Schmidt MA, Varma A, Chen XS, Martin ER, Wang L* (2022) Distinct sex-specific DNA methylation differences in Alzheimer's disease. *Alzheimer's Research & Therapy*, 14: 133 [[paper](#)] [[code](#)]

PRESENTATIONS

Oral

1. Upcoming: An Integrative Multi-Omics Random Forest Framework for Robust Biomarker Discovery, *STATGEN: Conference on Statistics in Genomics and Genetics*. May 2025. Minneapolis, MN, USA
2. Unlocking the potential of multi-omics data integration using multivariate random forest approach, *International Biometric Society Eastern North American Region (ENAR) Annual Meeting*. Mar 2024. Baltimore, MD, USA

Poster

1. An X chromosome-wide DNA methylation study of Alzheimer's disease, *Alzheimer's Association International Conference (AAIC)*, July 2024, Virtual Poster
2. Distinct CSF biomarker-associated DNA methylation in Alzheimer's disease and cognitively normal subjects, *Alzheimer's Association International Conference (AAIC)*, July 2023, Virtual Poster
3. Iterative Multivariate Random Forest for Feature Selection in Integrating Multi-Omics Datasets, *Annual American Statistical Association (ASA) Florida Chapter Meeting*, Mar 2023, Gainesville, FL, USA

TEACHING EXPERIENCE

Teaching Assistant

EPH705 Advanced Statistical Methods, Professor: Wang, Lily | University of Miami

Spring 2022-2024

STAT6201 Applied Linear Models, Professor: Barut, Emre | The George Washington University

Fall 2018

HONERS & AWARDS

Student Competition Award, ASA Florida Chapter Meeting

03/2023

Travel Award, University of Miami

03/2023

PROFESSIONAL SERVICES & ACTIVITIES

Manuscript Peer Review

Trusted Reviewer, Springer Nature

Manuscript Reviewer, Nature Communication; Discover Applied Sciences; Biology Direct; Medicine in Omics

Membership

International Biometric Society (ENAR)

American Statistical Association (ASA)

International Society to Advance Alzheimer's Research and Treatment (ISTAART)

TECHNICAL SKILLS

Proficient in R/Rstudio and Python for package building, data analysis, and visualization

Comprehensive skills in SAS for various statistical applications

Familiar with Linux system and command