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CONTACT

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EDUCATION

PhD., Biomedical Informatics, 2024 - Present

The Ohio State University, Columbus, OH, USA. (Qin Ma and Anjun Ma)

M.S., Biomedical, 2024

Zhengzhou University, Henan, Zhengzhou, China. (Ying Peng and Xia Xue)

B.S., Software Engineering, 2021

Zhengzhou University, Henan, Zhengzhou, China.

PROFESSIONAL EXPERIENCE

08/2024-present, Graduate Research Assistant

Department of Biomedical Informatics at The Ohio State University, OH

- Analyze bulk-level, single-cell, and spatial multi-omics datasets to derive meaningful insights for immuno-oncology research.
- Developed deep learning frameworks to enable cross-species analysis of single-cell and spatial omics data, providing new opportunities to investigate conserved and divergent molecular mechanisms across organisms.
- To establish a centralized Al-driven knowledgebase for integrating single-cell and spatial omics data across neurodegenerative diseases (ssKIND).

JOURNAL PUBLICATIONS

Full list: https://scholar.google.com/citations?user=kKMLVpUAAAAJ&hl=en

First and co-first author (#)

1. **Wu, W**.*, Liu, B. *, Zhang, Q. *, Zhang, X., Feng, P., Jia, Y., & Xue, X. (2025). Heterogeneity and efficacy of immunotherapy in multiple cancer: insights from a meta-analysis. <u>Biological Procedures</u> Online, 27(1), 17.

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2. Si, Y*., **Wu, W.***, Xue, X., Sun, X., Qin, Y., Li, Y., ... & Zheng, P. (2023). The evolution of SARS-CoV-2 and the COVID-19 pandemic. <u>PeerJ</u>, 11, e15990.

3. **Wu, W.***, Wang, Y., Xu, S., & Yan, K. (2020, September). Sfnn: Semantic features fusion neural network for multimodal sentiment analysis. In 2020 5th International Conference on Automation, Control and Robotics Engineering (<u>CACRE</u>) (pp. 661-665). IEEE.

Contributing author

- 1. Wang, Y., Ma, A., Song, N.J., Shannon, A.E., Amankwah, Y.S., Chen, X., **Wu, W**., Wang, Z., Saadey, A.A., Yousif, A. and Ghosh, G., 2025. Proteotoxic stress response drives T cell exhaustion and immune evasion. <u>Nature</u>, pp.1-11.
- 2. Dong, H., Liang, C., Zhang, J., **Wu, W.**, Kumar, N., Liu, Z., Sun, Y., Liao, Z., Cheng, X., Yu, Y. et al., 2025. O-GlcNAc transferase plays dual antiviral roles by integrating innate immunity and lipid metabolism. <u>Nature Communications</u>, 16, pp.1-14.
- 3. Miller, S.J., Zhong, H., **Wu, W.,** Cordova, A.M., Smith, M.E., Yashchenko, A., Li, Z., Jafree, D.J., Zimmerman, C.N., DeVette, C.I. and Do, V., 2025. A cross model spatial and single-cell atlas reveals the conserved involvement of osteopontin in polycystic kidney disease. <u>bioRxiv</u>, pp.2025-09.
- 4. Wang, X., Duan, M., Su, P.L., Li, J., Krull, J., Jin, J., Chen, H., Sun, Y., **Wu, W**., He, K. and Carpenter, R., 2025. Deep-learning-enabled multi-omics analyses for prediction of future metastasis in cancer. <u>bioRxiv</u>, pp.2025-05.
- 5. Jiang, Y., Wang, S., Feng, S., Wang, C., **Wu, W**., Huang, X., Ma, Q., Wang, J. and Ma, A., 2024. scGNN+: Adapting ChatGPT for Seamless Tutorial and Code Optimization. <u>bioRxiv</u>, pp.2024-09.
- 6. Ye, S., Li, C., Zhao, R. and **Wu, W**., 2019. NOAA-LSTM: A new method of dialect identification. In Artificial Intelligence and Security: 5th International Conference, <u>ICAIS 2019</u>, New York, NY, USA, July 26-28, 2019, Proceedings, Part I 5 (pp. 16-26). Springer International Publishing.

Journal reviewer:

2025, Heliyon, Guest Reviewer

PRESENTATIONS

2025,

MENTORING

2025/08, **Juile Zhu** (Undergraduate, OSU)

B.S. in Mathematics and Neuroscience

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AWARDS

• Second place in the multimodal emotion recognition track of the 'KDDI' Al algorithm competition (Rank:2/456).

- China University Computer Challenge "Network Technology Challenge", Second Prize in Central China Region.
- ACM Programming Competition, Zhengzhou University, First Prize.
- "Coretronic Cup" Future Car Human-Machine Interaction Design Competition, 7 th (Top 0.1%)
- "Challenge Cup" National Competition of Extracurricular Academic Science and Technology Works for University Students, First Prize in Henan Province.

SKILLS

- **Bioinformatics & Data Analysis**: Next-generation sequencing, proteomics, single-cell multi-omics data analysis, high-performance computing (R, Python, HPC systems)
- Software Development: Java Boot Spring, Vue, Flask, Sql.
- Machine Learning & AI: Pytorch.
- Data Visualization: Creation of impactful visualizations (Adobe Illustrator, ggplot2, BioRender)