

# Maxwell A. Xu

US Citizen

Personal Website: [maxxu05.github.io](https://maxxu05.github.io)

## EDUCATION

### University of Illinois Urbana-Champaign

*PhD. in Computer Science*

*Advised by Dr. James M. Rehg*

Expected 2026

### Georgia Institute of Technology

*M.S. in Computer Science*

### Johns Hopkins University

*B.S. in Applied Math / Biomedical Engineering*

## AWARDS

- National Science Foundation Graduate Research Fellowship
- Georgia Tech President's Fellowship
- Apple AIML Scholar Fellowship Finalist
- 2nd Place at the 11th Annual Jump ARCHES Symposium
- Intuitive Surgical Best Deep Learning Project Award

## SKILLS

**Machine Learning:** Time-Series, Self-Supervised Learning, Contrastive Learning, Imputation, Biosensors, Health

**Programming:** Pytorch, Jax, Flax, Tensorflow

## SELECTED EXPERIENCE

### Google Health

Student Researcher

Seattle, WA

Dec 2024 – Current

- Research on foundation models for wearable health biosignals on real-world production data.
- Developing scalable data and training pipelines for large-scale foundation model training using tensorflow-based infrastructure (i.e. Flax/Jax and TFDS).
- Collaborated across various interdisciplinary research efforts.

### Apple

Health AI Research Intern

Seattle, WA

April 2024 – Sep 2024

- Spearheaded research efforts on self-supervised learning methods for a motion foundation model.
- Collaborated with an interdisciplinary team to use health-domain knowledge in model development.
- Heavily assisted in engineering efforts for machine learning pipeline and infrastructure.

### Rehg Lab

PhD Student

Urbana, IL

Aug 2020 – Current

- Research interests include developing self-supervised learning methods for biosignal foundation models.
- Projects include a state-of-the-art motif-based time-series self-supervised learning method, physiological and health signal imputation via transformers, and disease progression modeling.
- First author published at ICLR (x2), NeurIPS, INSAR.

## PUBLICATIONS

**Xu, M.**<sup>†</sup>, Narayanswamy, G.<sup>†</sup>, Ayush, K., Spathis, D., Liao, S., Tailor, S. A., Metwally, A., Heydari, A. A., Zhang, Y., Garrison, J., Abdel-Ghaffar, S., Xu, X., Gu, K., Sunshine, J., Poh, M.-Z., Liu, Y., Althoff, T., Narayanan, S., Kohli, P., Malhotra, M., Patel, S., Yang, Y., Rehg, J. M., Liu, X.<sup>°</sup>, & McDuff, D.<sup>°</sup> (2025). LSM-2: Learning from Incomplete Wearable Sensor Data. *In Review at NeurIPS 2025*. DOI:

<https://doi.org/10.48550/arXiv.2506.05321> (<sup>†</sup>Co-first authors, <sup>°</sup>Co-last authors)

Zhang, Y., Ayush, K., Qiao, S., Heydari, A. A., Narayanswamy, G., **Xu, M.**, Metwally, A. A., Xu, S., Garrison, J., Xu, X., Althoff, T., Liu, Y., Kohli, P., Zhan, J., Malhotra, M., Patel, S., Mascolo, C., Liu, X., McDuff, D. & Yang, Y. (2025). SensorLM: Learning the Language of Wearable Sensors. *In Review at NeurIPS 2025*. DOI: <https://doi.org/10.48550/arXiv.2506.09108>

Gu, K., Zhang, Z., Lin, K., Zhang, Y., Paruchuri, A., Yu, H., Kazemi, M., Ayush, K., Heydari, A. A., **Xu, M.**, Narayanswamy, G., Liu, Y., Poh, M.-Z., Yang, Y., Malhotra, M., Patel, S., Palangi, H., Xu, X., McDuff, D., Althoff, T. & Liu, X. (2025). RADAR: Benchmarking Language Models on Imperfect Tabular Data. *In Review at NeurIPS 2025*. DOI: <https://doi.org/10.48550/arXiv.2506.08249>

Saha, M.<sup>†</sup>, **Xu, M.**<sup>†</sup>, A., Mao, W., Neupane, S., Rehg, J. M., & Kumar, S. (2025). Pulse-PPG: An Open-Source Field-Trained PPG Foundation Model for Wearable Applications Across Lab and Field Settings. *Ubiquitous Computing (UbiComp), Proceedings of the ACM on Interactive, Mobile, Wearable and Ubiquitous Technologies (IMWUT)*. DOI: <https://doi.org/10.48550/arXiv.2502.01108> (<sup>†</sup>Co-first authors)

- Xu, M.**, Narain, J., Darnell, G., Hallgrimsson, H., Jeong, H., Forde, D., Fineman, R., Raghuram, K., Rehg, J. M. & Ren, S. (2025). RelCon: Relative Contrastive Learning for a Motion Foundation Model for Wearable Data. The Thirteenth International Conference on Learning Representations (ICLR). DOI: <https://doi.org/10.48550/arXiv.2411.18822>
- Chow, W., Gardiner, L., Hallgrimsson, H., **Xu, M.**, & Ren, S. (2024). Towards Time-Series Reasoning with LLMs. (2024). Oral Presentation at 2024 NeurIPS Time Series in the Age of Large Models Workshop. DOI: <https://doi.org/10.48550/arXiv.2409.11376>
- Xu, M.**, Moreno, A., Wei, H., Marlin, B., & Rehg, J. M. (2024). REBAR: Retrieval-Based Reconstruction For Time-series Contrastive Learning. The Twelfth International Conference on Learning Representations (ICLR). DOI: <https://doi.org/10.48550/arXiv.2311.00519>
- Wei, H., **Xu, M.**, Samplawski, C., Rehg, J. M., Kumar, S., & Marlin, B. (2024). Temporally Multi-Scale Sparse Self-Attention for Physical Activity Data Imputation. Conference on Health, Inference, and Learning (CHIL). DOI: <https://doi.org/10.48550/arXiv.2406.18848>
- Xu, M.**, Moreno, A., Nagesh, S., Aydemir, V., Wetter, D., Kumar, S., & Rehg, J. M. (2022). PulseImpute: A Novel Benchmark Task for Pulsative Physiological Signal Imputation. Advances in Neural Information Processing Systems, 35, 26874-26888. DOI: <https://doi.org/10.48550/arXiv.2212.07514>
- Xu, M.**, Rehg, J., Rozga, A., McDaniel, J., Yoder, P., Watson, L., & Brady, N. (2022). Discovering Novel Predictors of Minimally Verbal Outcomes in Autism through Computational Modeling. International Society for Autism Research (Oral + Press Conference < 1% acceptance rate). Press release: <https://twitter.com/AutismINSAR/status/1524427451069345825>
- Liu, Y., Moreno, A.<sup>†</sup>, **Xu, M.**<sup>†</sup>, Li, S., McDaniel, J., Brady, N., Rozga, A., Li, F., Song, L., Rehg, J. Efficient Learning and Decoding of the Continuous-Time Hidden Markov Model for Disease Progression Modeling. arXiv [cs.LG]. 2021 Retrieved From <http://arxiv.org/abs/2110.13998> (†Co-second authors)
- Pomeranz Krummel, D., Nasti T., Kaluzova, M., Kallay, L., Melms, J., Izar, B., **Xu, M.**, Bhattacharya, D., Burnham, A., Ahmed, T., Li, G., Lawson, D., Kowalski, J., Cook, J., Medvedovic, M., Jenkins, A., Khan, M., Sengupta, S. Melanoma cell intrinsic GABAA receptor enhancement potentiates radiation and immune checkpoint inhibitor response by promoting direct and T cell-mediated anti-tumor activity. International Journal of Radiation Oncology. 2020. DOI: <https://doi.org/10.1016/j.ijrobp.2020.10.025>
- Pomeranz Krummel, D.<sup>‡</sup>, Nasti, T.<sup>‡</sup>, Izar, B.<sup>†</sup>, Press, R.<sup>†</sup>, **Xu, M.**<sup>†</sup>, Lowder, L., Kaluzova, M., Kallay, L., Rupji, M., Rosen, H., Su, J., Curran, W., Olson, J., Weinberg, B., Schniederjan, M., Neill, S., Lawson, D., Kowalski, J., Khan, M., Sengupta, S. Impact of sequencing radiation therapy and immune checkpoint inhibitors in the treatment of melanoma brain metastases. Radiation Oncology. 2020. DOI: <https://doi.org/10.1016/j.ijrobp.2020.01.043> (‡Co-first authors; †Co-second authors)
- Kallay, L., Keskin, H., Ross, A., Rupji, M., Moody, O., Wang, X., Li, G., Ahmed, T., Rashid, F., Rajesh Stephen, M., Cottrill, K., Nuckols, T., **Xu, M.**, Martinson, D., Tranchese, F., Pei, Y., Cook, J., Kowalski, J., Taylor, M., Jenkins, A., Pomeranz Krummel, D., Sengupta, S. Modulating native GABAA receptors in medulloblastoma with positive allosteric benzodiazepine-derivatives induces cell death. Journal of Neuro-Oncology. 2019; 142(3):411-422. doi: 10.1007/s11060-019-03115-0
- Pomeranz Krummel, D.<sup>‡</sup>, Tahseen, N.<sup>‡</sup>, Izar, B.<sup>†</sup>, **Xu, M.**<sup>†</sup>, Lowder, L., Press, R., Rupji, M., Kaluzova, M., Kallay, L., Burnham, A., Li, G., Ahmed, T., Chen, H., Curran, W., Kudchadkar, R., Olson, J., Schniederjan, M., Neill, S., Lawson, D., Cook, J., Weinberg, B., Jenkins, A., Kowalski, J., Khan, M., Sengupta, S. EXTH-12. Radiation enhances melanoma response to immunotherapy and synergizes with benzodiazepines to promote anti-tumor activity. Neuro-Oncology, 21(Supplement 6), November 2019, Page vi84, <https://doi.org/10.1093/neuonc/noz175.346> (‡Co-first authors; †Co-second authors)
- Kaluzova, M., Nasti, T., Chen, H.-R., Lowder, L., Press, R., Rosen, H., Rupji, M., Kallay, L., Patel, R., Burnham, A., **Xu, M.**, Ross, A., Keskin, H., Connelly, E., Izar, B., Adamson, C., Olson, J., Su, J., Curran, W., Kudchadkar, R., Schniederjan, M., Neill, S., Lawson, D., Chan, M., Kowalski, J., Khan, M., Pomeranz

Krummel, D., Sengupta, S. Abstract 247: Identification of the GABAA receptor in melanoma brain metastases patient tumors and demonstration that it is a viable drug target using benzodiazepine-derivatives. In: Proceedings of the American Association for Cancer Research Annual Meeting 2019; 2019 Mar 29-Apr 3; Atlanta, GA. Philadelphia (PA): AACR; Cancer Res 2019;79(13 Suppl). <https://doi.org/10.1158/1538-7445.AM2019-247>

Kallay, L., Keskin, H., Ross, A., Rupji, M., Moody, O., Wang, X., Li, G., Ahmed, T., Rashid, F., Rajesh Stephen, M., Cottrill, K., Nuckols, A., **Xu, M.**, Martinson, D., Tranchese, F., Pei, Y., Cook, J., Kowalski, J., Taylor, M., Jenkins, A., Pomeranz Krummel, D., Sengupta, S. Abstract 2623: Modulating native GABAA receptors in medulloblastoma with positive allosteric benzodiazepine-derivatives induces cell death. Proceedings of the American Association for Cancer Research Annual Meeting 2019; 2019 Mar 29-Apr 3; Atlanta, GA. Philadelphia (PA): AACR; Cancer Res 2019;79(13 Suppl). <https://doi.org/10.1158/1538-7445.AM2019-2623>

Kowalski, J., Pomeranz Krummel, D., Rupji, M., Dwivedi, B., Keskin, H., Kallay, K., **Xu, M.**, Ross, A., Press, R., Rosen, H., Connelly, E., Patel, R., Izar, B., Adamson, C., Olson, J., Su, J., Kudchadkar, R., Schniederjan, M., Lowder, L., Neill, S., Curran, W., Lawson, D., Chan, M., Khan, M., Sengupta, S. COMP-22: Large scale transcriptomic analysis of melanoma brain metastases, Neuro-Oncology, Volume 20, Issue suppl\_6, 1 November 2018, Page vi68, <https://doi.org/10.1093/neuonc/noy148.277>

Kowalski, J., Pomeranz Krummel, D., Rupji, M., Dwivedi, B., Keskin, H., Kallay, K., **Xu, M.**, Ross, A., Press, R., Rosen, H., Connelly, E., Patel, R., Izar, B., Adamson, C., Olson, J., Su, J., Kudchadkar, R., Schniederjan, M., Lowder, L., Neill, S., Curran, W., Lawson, D., Chan, M., Khan, M., Sengupta, S. CD131: Large scale transcriptomic analysis of melanoma brain metastases. Annals of Neurology 84(suppl 22), 2018.

Kallay, L., Keskin, H., Ross, A., Moody, O., Cottrill, K., Nuckols, A., Li, G., Ahmed, T., Rashid, F., Rajesh Stephen, M., **Xu, M.**, Martinson, D., Macdonald, T., Kowalski, J., Wang, X., Taylor, M., Cook, J., Jenkins, A., Pomeranz Krummel, D., Sengupta, S. PDTM-45: Positive modulation of native gabaa receptors in medulloblastoma cancer cells with benzodiazepines induces rapid mitochondrial fragmentation and tp53-dependent, cell cycle-independent apoptosis. Neuro-Oncology, Volume 20, Issue suppl\_6, 1 November 2018, Page vi213, <https://doi.org/10.1093/neuonc/noy148.884>

## FURTHER EXPERIENCE

### Systems & Technology Research

Boston, MA

Machine Learning Researcher

Jan 2020 – Aug 2020

- Conducted machine learning research for identifying vulnerabilities in speech-processing programs with reinforcement learning and representation learning techniques.
- Worked on frontend UI and backend data retrieval, visualization, and unit testing

### Medtronic plc

New Haven, CT

Data Science Engineer Contractor

May 2019 – Dec 2019

- Predicted lung cancer recurrence from tabular clinical data.
- Utilized survival analysis with Kaplan-Meier curve visualizations to identify cancer recurrence risk factors.

### BioSwift Biomedical Engineering Design Team

Baltimore, MD

Former Chief Executive Officer and Co-Founder

Aug 2018 – Dec 2019

- Patent pending design of inhaler attachment to assist pediatric asthmatic patients
- Secured over \$10,000 in funding, including from the Johns Hopkins Student Initiatives Fund
- Won 1<sup>st</sup> place in the 2019 ASAIOfyi Student Design Competition, 1<sup>st</sup> place at Fall 2019 Johns Hopkins FastForward U Spark Accelerator Competition, 3<sup>rd</sup> place at 2019 Johns Hopkins BPC
- Press release:
  - <https://ventures.jhu.edu/news/bioswift-aquatas-fast-forward-u-accelerator-demo-days/>
  - <https://www.jhunewsletter.com/article/2019/11/fastforward-u-teams-innovate-with-new-and-old-technologies>
  - <https://www.facebook.com/JohnsHopkinsBME/photos/a.131397713949264/692413781180985/?type=3&theater>

### Modeling A Transferable Histopathological Image Analysis System

Baltimore, MD

Personal Project

Dec 2019

- Developed a deep clustering representation learning method for histopathology images

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- Won Intuitive Surgical Best Deep Learning Project Award (\$800 cash prize)
- Press Release: <https://www.cs.jhu.edu/2020/01/28/deep-learning-course-prepares-students-for-success-in-ai-careers/>

## **Johns Hopkins Department of Computer Science**

Baltimore, MD

Data Structures Teaching Assistant

Jan 2019 – Dec 2019

- Collaborated with team of faculty during weekly meetings and actively contributed to course content
- Worked with students to enhance student understanding of content such as heaps, AVL trees, hashmaps

## **Sengupta-Krummel Lab at the Emory Winship Cancer Institute**

Atlanta, GA

Cancer Genomics Bioinformatics Researcher

May 2017 – Dec 2019

- Employed hierarchical clustering techniques for analysis of RNA-seq gene expression data
- Used DESeq2 method to identify differentially expressed genes followed by pathway analysis
- Conducted survival analysis with the Kaplan-Meier estimator and Cox PH Regression Model

## **Johns Hopkins Department of Applied Mathematics and Statistics**

Baltimore, MD

Probability and Statistics Teaching Assistant

Jan 2019 – May 2019

- Nominated for Spring 2019 Teaching Award for demonstrating an extraordinary commitment to pedagogy

## **Johns Hopkins University PILOT Learning Program**

Baltimore, MD

Head PILOT Leader of Organic Chemistry

Aug 2017 – May 2019

- Gave lectures to enhance student understanding of content, directed other PILOT leaders in their roles to teach students, and developed problem sets to be distributed among the entire program

## **Centers for Disease Control and Prevention**

Atlanta, GA

Outbreak Surveillance Biostatistics Intern

May 2018 – Aug 2018

- Utilized whole genome multilocus sequence typing data to detect disease clusters
- Employed machine learning techniques to develop this disease cluster detection method for *Listeria monocytogenes*. Validated identified clusters with Simpson's Index of Diversity with PulseNet cluster codes and Chi-Squared ranked overlap between multi-dimensional independent hierarchical clustering