Obblivignes KanchanadeviVenkataraman

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SKILLS

Languages & Tools | Python, MATLAB, C, C++, Java, LaTeX, Git, Bash

Libraries & Frameworks | PyTorch, Scikit-Learn, Pandas, Plotly, Numpy, Matplotlib, Keras, Scipy, Electron JS, React.JS

Machine Learning | Classical (Linear Regression, Classification, Clustering, Tree-based methods), Deep (LiteLLM, GPT)

Data Science | Data Preprocessing, Data Visualization, Statistical Analysis, Monte Carlo Simulations, Predictive Modeling

WORK EXPERIENCE

Graduate Teaching Assistant - CSC 326 Software Engineering | NC State University

August 2025 - Present

- Mentored 60 students in two lab sections collaborate to complete assignments, and build a full-stack development application.
- Held office hours to help students with questions or concerns they have regarding course material and project issues
- Graded assignments and proctored exams for all students

Graduate Research Assistant | NC State University - MAP Lab & GIS Lab

September 2024 - Present

- Built end-to-end data pipelines for super-resolution microscopy using **Python (NumPy, Pandas, Scikit-learn)**, applying clustering, classification, and regression to extract actionable biological insights.
- Generated 500k+ physics-informed Monte Carlo simulations, boosting model accuracy by 25% and training efficiency by 40% with feature engineering and custom normalization.
- Developed and fine-tuned **deep learning models** (**PyTorch: U-Net, Attention U-Net, Conditional Diffusion**), doubling image quality in super-resolution denoising, and up to **50% better** denoising performance compared to baseline models.
- Improved model generalization by 50% from baselines through hyperparameter tuning, cross-validation, and statistical evaluation
- Partnered with cross-functional teams to integrate **AI/ML workflows** into biology and materials science research, accelerating experimentation and interdisciplinary innovation.

Graduate Research Assistant | NC State University - Dr. Bowen Xu's Lab [Link]

January 2024 - May 2024

- Presented, trained, and evaluated the performance of an existing deep **just-in-time SE code-comment consistency checker** technique among existing datasets such as Microsoft's CodeXGlue.
- Integrated the tool into SEEDGuard GitHub repository, providing video-guided tutorials and Docker images for easy deployment.

Graduate Research Assistant | NC State University - Dr. Ruozhou Yu's Lab [Link]

January 2023 - August 2023

- Conducted **microservice-based distributed application benchmarking** by implementing a comprehensive Kubernetes cluster, utilizing Raspberry Pis, an Apache Cassandra database, Flask, Jaeger, and Opentelemetry.
- Trained several machine learning models and achieved a **96% R2-score** using Random Forest models and presented all findings using data visualization techniques in a comprehensive report.

RELEVANT PROJECTS

Applying SpecUNet Towards Spatial Single-Molecule Imaging Analysis

August 2025 - Present

(Python, PyTorch, Scipy, Pandas, matplotlib, imago, h5py)

- Applied SpecU-Net simulation process to generate 100k+ Gaussian-based Monte Carlo single-molecule spectral simulations
- Trained Spec-UNet to achieve significant improvements in performance compared to existing denoising tools

SIMBA: Single-molecule Imaging with Multi-agent Bot Assistant

June 2025 - Present

(Electron JS, ChatGPT, ReactJS, LiteLLM, Python, PyTorch, matplotlib, scipy)

- Led a multi-agent desktop app to perform spectroscopic analyses, image denoising, single-molecule localization, and other tasks
- Significantly improved scientific analysis workflow by seamlessly integrating microscopy tasks with built-in OpenAI GPT chatbot

SpecDiffuse for Accurate Spectroscopic Single Molecule Localization Microscopy

(MATLAB, Python, PyTorch, Scipy, Pandas, matplotlib, imago, h5py)

May 2025 - Present

- Adapted existing **conditional diffusion model** with U-Net backbone on super-resolution denoising for spectroscopic analysis
- Outperformed conventional U-Net (SpecU-Net) by ~50% on several metrics in image quality for super-resolution denoising
- Enabled accurate biological analysis by simulating 400k+ 16-bit images (~25GB) for robust GPU-based training and inference

SA-SpecUNet for Accurate Spectroscopic Single Molecule Localization Microscopy

(MATLAB, Python, PyTorch, SciPy, Pandas, Matplotlib, imago, h5py)

November 2024 - Present

- Integrated spatial attention component into SpecU-Net to capture specific latent representations for better contextual understanding
- Outperformed conventional U-Net (SpecU-Net) by ~25% on several metrics in image quality for super-resolution denoising
- Enabled accurate biological analysis by simulating 400k+ 16-bit images (~25GB) for robust GPU-based training and inference

$\textbf{Framework Development for Accurate Single-Molecule Imaging Spectroscopic Analysis } [\underline{Link}]$

(MATLAB, Python, PyTorch, SciPy, Pandas, matplotlib, imago, h5py)

September 2024 - December 2024

- Improved image resolution by ~70% from conventional denoising methods by developing a U-Net-based model in MATLAB (SpecU-Net) to denoise spectroscopic single-molecule localization images
- Increased prediction precision by ~70% and accuracy by ~50% through generating 100k+ physics-informed Monte Carlo simulation datasets and creating comprehensive evaluation metrics
- Migrated existing SpecU-Net pipeline from MATLAB to Python, achieving 5x GPU acceleration on training and inference process

EDUCATION

North Carolina State University

August 2025 - Present

Ph.D. in Computer Science | GPA: N/A

August 2023 - May 2025

North Carolina State University *M.S. in Textiles* | *GPA:* 3.94/4

August 2020 - May 2023

North Carolina State University

B.S. in Textile Engineering | GPA: 4.00/4

August 2020 - May 2025

PURLICATIONS

• Mao, H., Liu, Y., KanchanadeviVenkataraman O., Shahid, M. A., Laplante C., Xu, D., Song, K., Zhang, Y. (2025). Framework for Accurate Single-Molecule Spectroscopic Imaging Analyses Using Monte Carlo Simulation and Deep Learning. Analytical Chemistry, 97(30), 16250-16258.