# Obblivignes KanchanadeviVenkataraman

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### **EDUCATION**

North Carolina State University

Doctorate of Philosophy in Computer Science

August 2025 - Expected May 2029

GPA: N/A

August 2023 - May 2025

GPA: 3.94/4

August 2020 - May 2023

GPA: 4.00/4

Bachelor of Science in Computer Engineering

Master of Science in Computer Science.

#### **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 | Classification, Clustering, Tree-based methods), Deep (UNet, Diffusion, LiteLLM, GPT)

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

### **WORK EXPERIENCE**

**Graduate Teaching Assistant** - 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

CV/ML Graduate Research Assistant (GRA) | 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 hyper-parameter 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.

**Software Engineering AI GRA** | 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.

 $\textbf{Distributed Systems RA} \mid \text{NC State University - Dr. Ruozhou Yu's Lab} \ [\underline{\text{Link}}]$ 

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.

## **PROJECTS**

Computer Vision (Applying SpecUNet and SA-SpecUNet Towards Spatial SMLM)

August 2025 - Present

- Applied SpecU-Net to generate 100k+ Gaussian- and Perlin-based Monte Carlo single-molecule spatial simulations
- Trained SpecU-Net to achieve significant improvements in super-resolution denoising performance compared to existing tools

Multi-AI Agent System (SIMBA: Single-molecule Imaging with Multi-agent Bot Assistant)

June 2025 - Present

- · Led a multi-agent desktop app to perform spectroscopic analyses, image denoising, single-molecule localization, and other tasks
- Utilized Electron JS and React JS for front-end integration and utilized GPT-40 and LiteLLM for backend integration
- Significantly improved scientific analysis workflow by seamlessly integrating microscopy tasks (denoising, localization, analysis)

Computer Vision (Applying SpecDiffuse for Accurate Spectroscopic SMLM)

May 2025 - Present

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

Computer Vision (Applying SA-SpecU-Net for Accurate Spectroscopic SMLM)

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

**Computer Vision** (Framework for Accurate SMLM Spectroscopic Analysis) [<u>Link</u>]

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 SpecU-Net pipeline from MATLAB to Python using PyTorch, achieving **5x GPU acceleration** on training and inference

### **PUBLICATIONS**

• 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.