

# Obblivignes KanchanadeviVenkataraman

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## EDUCATION

### North Carolina State University

Doctorate of Philosophy in Computer Science	August 2025 - Expected May 2029
	GPA: 4.00/4
Master of Science in Computer Science	August 2023 - May 2025
	GPA: 3.97/4
Bachelor of Science in Computer Engineering	August 2020 - May 2023
	GPA: 4.00/4

## SKILLS

**Languages & Tools** | Python, MATLAB, C, C++, Java, LaTeX, Git, Bash, SLURM  
**Libraries & Frameworks** | PyTorch, Scikit-Learn, Pandas, Plotly, Numpy, Matplotlib, Keras, Scipy, Electron JS, React.JS  
**Machine Learning** | **Classical** (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** - Automated Learning & Data Analysis | NC State University January 2026 - Present
- Held office hours for **60 students** to understand material, complete assignments, and build a real-world machine learning project.
  - Managed grading for all student coursework, including assignments, quizzes, and exams.
- Graduate Teaching Assistant** - Software Engineering | NC State University August 2025 - December 2025
- 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.
- 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.
- Distributed Systems RA** | 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
  - 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-4o 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, physics-informed model with U-Net backbone on SR 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) [\[Link\]](#) 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.