

# Jihyeon Je

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## Education

### Duke University

Durham, NC

BSE in Biomedical Engineering and Electrical & Computer Engineering,

May 2023

BS in Computer Science with a concentration in Artificial Intelligence and Machine Learning

Coursework in BME (Medical Instrumentation, Materials and Synthetic Biology), ECE (Microelectronics, Quantum

Engineering/Information Theory), Computer Science (Computer Security, Algorithms, Neurosymbolic Machine Learning)

2020 Woo Fellow, 2022/21 DTech Scholar, 2021-22 Duke Bass Connections Fellow | GPA: 3.83/4.0

## Research Experience

### Duke University, Lafata Lab

Durham, NC

Undergraduate Researcher

May 2020 – present

- Developing an automatized feature extraction tool for the analysis of next generation pathomic signatures of inflammation from renal biopsy data
- Generation of synthetic time series image data using Fokker-Planck dynamics to create prediction models for head-and-neck cancer

### Schrodinger, Machine Learning Team

New York, NY

Machine Learning Intern

May 2022 – Aug 2022

- Implemented diffusion-based generative model for ligand conformation generation to improve conformer quality
- Introduced new features and Gaussian processes to supervised regression and classification models and improved small molecule property prediction
- Devised and implemented automated routine for benchmarking and backtesting

### Broad Institute of MIT and Harvard, Imaging Platform, Cimini Lab

Cambridge, MA

Research Intern

May 2021 – Aug 2021

- Built ML-based image analysis tools and workflows for 2D and 3D image segmentation and reconstruction
- Devised optimized strategies and network architectures to efficiently utilize sparse and limited bioimage data

### Duke University, Caron Lab

Durham, NC

Undergraduate Intern

Sep 2019 – Mar 2020

- Worked on 3D reconstruction and statistical analysis of dendritic spines from EM images

### NCMIR (National Center for Microscopy and Imaging Research), Mark Ellisman Lab

San Diego, CA

Intern

May 2017 – Mar 2020

- Worked on 3D computational reconstruction and segmentation of electron microscope images
- Contributed to CDeep3M-Preview by building the augmentation pipeline and wrote additional image analysis scripts for feature extraction from large-scale biological data

### UNIST (Ulsan National Institute of Science and Technology), Protein Engineering Lab

Ulsan, South Korea

Research Intern

Mar 2015 – Jul 2016

- Developed a fusion nano probe using affibody molecules for targeted cancer therapy

## Additional Experience and Teaching

### Wolfram Alpha

Remote

2018 Wolfram Summer School Alumni, Student Ambassador

May 2018 – present

- Research focus on mathematical and computational analysis of the structure of viral capsids
- Developed the Protein Data Bank importer function for the Mathematica functionality repository

### Duke University Department of Electrical and Computer Engineering

Durham, NC

Teaching Assistant

Jan 2021 – Jan 2022

- Held lab sessions, office hours, and provide tutoring for students in Electrical & Computer Engineering 280: Signals and Systems

## Leadership & Activities

### Duke Undergraduate Machine Learning

Co-President

Durham, NC

Sep 2019 - present

- Organize yearly Datathon and Machine Learning Day events
- Host guest speaker sessions, manage funding, and plan monthly activities.

### Duke iGEM (international Genetically Engineered Machine)

Subteam Lead

Durham, NC

Jan 2020 - present

- 2022 iGEM Silver Medal
- Developing a microfluidics-based organoid-tumor coculture platform for high throughput drug screening
- Creating a automatized computational tool to quantify organoid growth and drug efficiency

### Duke ARAC (America Reads America Counts)

Volunteer

Durham, NC

Sep 2020 - present

- Volunteered at Durham Public Schools tutoring children to help them enhance primary-level reading and math skills

## Publications

Haberl M.G., Wong W., Penticoff S., **Je J.**, Madany M., Borchhardt A., Boassa D., Peltier S.T., Ellisman M.H. *CDeep3M-preview: Online segmentation using the deep neural network model zoo*.

**Preprint** at: <https://doi.org/10.1101/2020.03.26.010660>

## Abstracts and Conference Proceedings

Stevens J, **Je J**, Gao Y, Wang C, Mowery Y, Brizel D, Yin F, Liu J, Lafata K. Radiomics on spatial-temporal manifolds via Fokker-Planck dynamics. American Association of Physicists in Medicine. 2022. **Poster presentation** delivered at the AAPM meeting, September, 2022.

Sotolongo G, **Je J**, Li X, Wang Y, Zee J, Wang B, Chen Y, Talawalla T, Hodgins J, Madabhushi A, Ozeky T, Mariani L, Holzman L, Janowczyk A, Barisoni L, Lafata K. Segmentation and Classification of Lymphocytes in the NEPTUNE Digital Kidney Biopsies via PatchSorter. United States and Canadian Academy of Pathology abstract. 34:847. **Poster presentation** delivered at the USCAP meeting, March, 2022.

**Je J**, Lucas A, Sterling D, Cimini B. Network Optimization with Limited Bioimage Data for Robust Semantic Segmentation. Society of Biomolecular Imaging and Informatics. **2<sup>nd</sup> Place Best Poster Award**, presentation delivered at the 2021 High Content meeting, remote, October, 2021.

Sotolongo G, **Je J**, Zee J, Chen Y, Li X, Wang Y, Hodgins J, Madabhushi A, Janowczyk A, Lafata K, Barisoni L. Cortical Tubulointerstitial Mononuclear Inflammation in Renal Biopsies is a Quantitative Biomarker of Clinical Outcomes in NEPTUNE Glomerular. United States and Canadian Academy of Pathology abstract 34:847. **Poster presentation** delivered at the USCAP meeting, remote, October, 2020.

## Technical Skills

### Programming

**Advanced:** Python, Java, Pytorch, TensorFlow, MATLAB, Scikit-learn, RDKit

**Intermediate:** CSS, JavaScript, HTML, SQL

### DevOps

Jenkins CI/CD

### Chemical

QSAR, ADMET prediction, conformer generation

### Biological

CellProfiler, Fiji/ImageJ, IMod, QuPath