

# YANGYANG LI

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## SELF-INTRODUCTION

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I am a third-year Ph.D. candidate with a focus on deep learning and algorithm development for complex biological problems. My area of expertise is machine learning-based, data-driven domains. I am skilled at applying these technologies to solve scientific issues, and I am eager to contribute and further my knowledge in a fast-paced professional environment.

## RESEARCH EXPERIENCE AND PROJECT

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### Northwestern University

*Ph.D. in Bioinformatics*

Chicago, US

Sep. 2022 – Present

- Developed a deep generative model for sequencing data simulation
- Introduced a graph algorithm to identify non-linear transcripts in long-read data, achieving a 20x speedup
- Crafted a web application for graph algorithm visualization
- Designed a Python interface for a C-based command-line tool, gaining 20% performance boosts

### University of Minnesota

*Ph.D. in Bioinformatics and Computational Biology*

Minnesota, US

Sep. 2020 – Present

- Pioneered an algorithm to discern non-linear structure variations in transcriptomes
- Evaluated the efficacy of prevalent tools for detecting alternative splicing variants

### China Agricultural University

*Master in Crop Bioinformatics*

Beijing, CN

Sep. 2018 – June 2020

- Identified key features in 1,400 maize genomics datasets to enhance agronomic traits
- Examined the correlation between genetic variations and maize ear characteristics in 450 natural populations

## EDUCATION

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### Northwestern University

*Ph.D in Bioinformatics*

Chicago, US

June 2022 – present

### University of Minnesota

*Ph.D. in Bioinformatics and Computational Biology*

Minnesota, US

Sep. 2020 – June 2022

### China Agricultural University

*Master in Crop Bioinformatics*

Beijing, CN

Sep. 2018 – June 2020

### Northeast Agricultural University

*Bachelor of Arts in Agricultural Engineering*

Harbin, CN

Sep. 2014 – June 2018

## TECHNICAL SKILLS

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**Languages and Frameworks:** C/C++, Python, Rust, Pytorch, Jax

**Developer Tools:** Neovim, Git, Docker, GitHub Action, Gcc, Clang

**Specializations:** Algorithm Development, Concurrency Programming, Data Analysis and Visualization, Natural Language Processing

## PUBLICATIONS

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Fry, J., Li, Yangyang, & Yang, R. (2022, 09). ScanExitronLR: characterization and quantification of exon splicing events in long-read RNA-seq data. *Bioinformatics*. doi: 10.1093/bioinformatics/btac626

Li, Yangyang, & Yang, R. (2023). Pxblat: An ergonomic and efficient python binding library for blat. *bioRxiv*. doi: 10.1101/2023.08.02.551686