

# Damir Temir

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## EDUCATION

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### University of Illinois Springfield

August 2019 - May 2023

*B.Sc. in Computer Science, GPA: 4.0/4.0*

*Springfield, IL*

- Relevant Coursework: Programming Concepts, Programming Languages, Data Structures & Algorithms, Intro to Neural Networks, Intro to Operating Systems, Applied Statistics, Calculus 2, Linear Algebra
- Activities: Capital Honors Program, Peer Mentor for First-Year Students, Peer Tutor in Computer Science
- Achievements: Full-Ride Scholarship Recipient, Dean's List Honoree, Outstanding Sophomore Award

## TECHNICAL SKILLS

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**Languages:** Python, SQL, JavaScript, Java, Go

**Frameworks:** PyTorch, Hugging Face, Flask, Django, React

**Developer Tools:** Linux, Git, GitHub Actions, Docker, BentoML, Heroku

**Libraries:** NumPy, pandas, Matplotlib, scikit-learn

## EXPERIENCE

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### Machine Learning Intern

May 2022 - August 2022

*Northwestern Medicine*

*Chicago, IL*

- Integrated Transformer-based NLP models to patient ECG-data, generating useful, clinically correct analyses
- Assembled React Web Interfaces for clinicians interacting with NLP models
- Built out internal data workflows for Power BI dashboard generation across business infrastructure metrics

### Software Engineering Fellow

January 2022 – April 2022

*Major League Hacking*

*Remote*

- Contributed to Facebook Research [LabGraph](#), building real-time messaging monitor for graph benchmarking
- Supported CI/CD using GitHub Actions and maintained high quality of code with automated testing

### Open Source Fellow

September 2021 – December 2021

*Major League Hacking*

*Remote*

- Supported PyCaret & LightGBM in an open-source project [BentoML](#) to deploy ML models in production
- Used data-validation tools for deployment schema verification, supporting the [Bento Deployment Tool](#)

### Research Assistant

June 2021 – August 2021

*Stanford University*

*Stanford, CA*

- Developed a statistical interface to control amino acids distributions produced by a Convolutional Neural Network
- Updated the *Protein Design Algorithm* using a Graph Neural Network speeding sequence initialization by 300%
- Documented and presented [research findings](#) at the Protein Design Conference, *Summer RosettaCON 2021*

## PROJECTS

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### Facebook LabGraph Monitor | *TypeScript, Node.js, Python, GitHub Actions, Git*

January 2022 - April 2022

- Equipped LabGraph with an extension to visualize data streaming between graphs and nodes
- Serialized graph topology along with Messages, Streams, and Groups
- Setup [Sphinx Documentation](#) that auto-deploys on GitHub Pages with each commit using GitHub Actions

### Fellowship Prediction | *Python, BentoML, Docker, Flask, React, Git*

September 2021

- Constructed a web tool to analyze GitHub Profiles of aspiring MLH Fellows to gain insights into their statistics
- Mined data on 600 GitHub users to deliver a BentoML Prediction Service in production on Heroku with Docker
- Collaborated with a team, winning the [Fellowship Orientation Hackathon](#) and getting 49 stars on GitHub

### Resfile Interface | *Python, PyTorch, NumPy, PyRosetta, Git*

June 2021 - September 2021

- Built a statistical interface to modify the logits in PyTorch tensors under the Pseudo-Log-Likelihood (PLL) mathematical model to produce *internal hydrogen bonding networks*
- Substituted the Baseline Convolutional Neural Network Model with a Graph Neural Network, increasing the time efficiency of initializing the starting sequence for protein design by 300%