

# Damir Temir

312-478-7890 | [damir@temir.dev](mailto:damir@temir.dev) | [linkedin.com/in/damirtemir](https://linkedin.com/in/damirtemir) | [github.com/dtemir](https://github.com/dtemir) | [temir.dev](https://temir.dev)

## EDUCATION

### University of Illinois Springfield

Springfield, IL

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

*Aug. 2019 – May 2023*

- Relevant Coursework: Programming Concepts, Discrete Structures, Data Structures & Algorithms, Applied Statistics, Calculus 2, Linear Algebra, Computer Organization, Introduction to Neural Networks
- 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

**Languages:** Python, SQL (Postgres), JavaScript, HTML/CSS

**Frameworks:** PyTorch, Django, BentoML

**Developer Tools:** Git, Docker, Heroku

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

## EXPERIENCE

### Open-Source Fellow

September 2021 – Present

*Major League Hacking*

*Remote*

- Built support for an NLP framework in the core of an open-source project BentoML used by Data Scientists to deliver their Machine Learning projects in production
- Learned the best practices of Git and GitHub Project Management, exploring ways to work at a startup

### Pre-Fellow

July 2021 – August 2021

*Major League Hacking*

*Remote*

- Assembled a Discord interface to automate daily GitHub discussion posts using REST APIs
- Contributed 20 Pull Request to team projects, actively developing group ideas
- Collaborated with a team of international developers to build group projects in a professional setting with daily standup meetings

### Deep Learning Research Assistant

June 2021 – August 2021

*Stanford University*

*Stanford, CA*

- Developed a statistical interface to control distributions of amino acids produced by a Convolutional Neural Network
- Remodeled the *Protein Sequence Design Algorithm* using a Graph Neural Network to speed up sequence initialization by 300%
- Documented and presented research findings at the Protein Design Conference, *Summer RosettaCON 2021*

## PROJECTS

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

September 2021

- Constructed a web tool to analyze GitHub Profiles of aspiring MLH Fellows for them to gain insights into their statistics
- Mined data on over 600 GitHub users to deliver a BentoML Prediction Service in production on Heroku with Docker
- Utilized the best collaborative practices with GitHub Projects, Pull Request Templates, and Extensive Documentation, winning an Orientation Hackathon

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

June 2021 – August 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%

### U.S. Consumer Complaint Analysis | Python, Pandas, NumPy, Scikit-Learn, Folium, Git

April 2021

- Analyzed an extensive federal dataset about the type of consumer complaints on financial institutions, building a heuristic model to rank their actions based on the volume, response type, disputes, and location
- Reported results with the most highly and poorly rated financial institutions, presenting results in graphs, plots, and maps