# Philip Wisniewski

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#### Education

**Purdue University** 

West Lafayette, IN August 2023 – May 2027

B.S. Computer Science

GPA: 4.0/4.0, Honors College

Key Courses: Object Oriented Programming, Data Structures and Algorithms, Computer Architecture, Discrete Math

## Experience

Research Intern

West Lafayette, IN

Rosen Center for Advanced Computing

May 2024 - Present

- Researched and developed a queue time prediction model using historical data from the SLURM workload manager on the ANVIL supercomputing cluster.
- Trained a densely connected feedforward neural network with PyTorch, integrating it into a command line tool for enhanced operational efficiency.
- Utilized Python, R, PyTorch, PostgreSQL, and SLURM to investigate, store, and calculate trends in data.

Research Intern

Rochester, MN

Mayo Clinic

June 2022 – August 2023

- Researched and predicted cancer cell line drug resistance utilizing artificial intelligence from various genetic data.
- Investigated effects of specific omic data and pathway expressions on modeling.
- Presented findings at two poster conferences (Explore the Exposome: Individualizing Medicine Conference and Mayo Clinic SURF Symposium).

#### Technical Skills

Languages: Python, C, C#, Java, SQL (Postgres), R

Developer Tools: Git, Docker, Rancher, VS Code, Visual Studio, PyCharm, IntelliJ, Spyder, Lex, Yacc

Libraries: Pandas, NumPy, Matplotlib, PyTorch, TensorFlow, Scikit-Learn

#### **Publications**

- Lovell A\*, **Wisniewski P\***, Rodenbeck S, Ashish. "A Hierarchical Deep Learning Approach for Predicting Job Queue Times in HPC Systems." *Proceedings of HUST-24: 11th International Workshop on HPC User Support Tools, Supercomputing '24 Conference, 2024.*
- Meng-Lin K, Ung CY, Zhang C, Weiskittel TM, **Wisniewski P**, Zhang Z, Tan SH, Yeo KS, Zhu S, Correia C, Li H. SPIN-AI: A Deep Learning Model That Identifies Spatially Predictive Genes. Biomolecules. 2023 May 27;13(6):895.
- Weiskittel TM, Cao A, Meng-lin K, Lehmann ZJ, Feng B, Correia C, Zhang C, Wisniewski P, Zhu S, Ung CY, Li H. Network biology inspired machine learning features predict cancer gene targets and reveals target coordinating mechanisms. Pharmaceuticals (Basel). 2023 May 16;16(5):752.

### Awards

- Dean's List and Semester Honors for all semesters
- National Merit Scholar Finalist
- AP Scholar with Distinction
- Presidential Scholar Candidate

<sup>\*</sup> Designates co-first authorship