

Pat Wall
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Education

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|-----------------|---|-----------------------------|
| <i>May 2023</i> | M.S. Informatics <ul style="list-style-type: none">• Concentration: Complex Networks and Systems• Relevant Courses: Network Science, Signal Processing and Information Theory, Nonparametric Statistics, SNP Discovery and Population Genetics• Exited PhD program when advisor left institution | Indiana University |
| <i>May 2019</i> | B.S. Biochemistry <ul style="list-style-type: none">• Minor: Mathematics | Eastern Michigan University |

Academic Projects

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| <i>Jan. 2022—
Present</i> | Dormancy in 2D discrete dynamical systems <p>Advisor: Jay Lennon</p> <ul style="list-style-type: none">• Implemented 2D cellular automaton models to run in parallel using Rust and characterized statistical differences in outcomes for different models.• Produced observations consistent with current theoretical understanding of dormancy and illustrated specific patterns we should expect in empirical data. |
| <i>May 2020—
Jan. 2022</i> | Predicting complex traits in metagenomes <p>Advisor: Jay Lennon</p> <ul style="list-style-type: none">• Leveraged database of thousands of publically available metagenomes to develop random forest classifier for bacterial genomes capacity to enter dormancy.• Enabled further discovery and testing of evolutionary theory by all current and future members of the lab. |
| <i>Feb. 2020—
May 2021</i> | Testing multivariate information theory methodology <p>Advisor: Luis Rocha</p> <ul style="list-style-type: none">• Evaluated several multivariate information theoretic methods for their ability to uncover these structures.• Generated synthetic time series data from networks with known hypergraph structures.• Showed that multivariate information theoretic methods can predict some dynamical features while casting doubt on the ability of these tools to reveal underlying structure. |

Skills & Achievements

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| <i>Languages</i> | Python, Rust, C++, R, Bash, SQL, L ^A T _E X |
| <i>Libraries</i> | NumPy, Pandas, scikit-learn, TensorFlow, graph_tool, Tidyverse, vegan |
| <i>Other Software</i> | GitHub (GitHub actions for CI/CD), Heroku, BLAST and other bioinformatics tools |
| <i>Awards</i> | NSF National Research Traineeship (\$72,000) |