# Brian Andrews

# Education

Ph.D. Candidate, Drexel University, Philadelphia, PA.

Fall 2018-Present

Physics - Research in Biophysics & Molecular Simulation. Expected Grad: Early 2023.

M.S., Case Western Reserve University, Cleveland, OH.

Fall 2016-Summer 2018

Physics - Entrepreneurship Track

B.A., Kenyon College, Gambier, OH.

Physics - Math Minor - Scientific Computing Concentration

Fall 2012-Spring 2016

## Research

#### **Publications**

- 6. **Brian Andrews**, Thomas Ruggiero, and Brigita Urbanc. How Do Salt and Lipids Affect Conformational Dynamics of A $\beta$ 42 Monomers in Water? *Phys. Chem. Chem. Phys.*, 2022. Submitted.
- 5. **Brian Andrews**, Jose Guerra, Reinhard Schweitzer-Stenner, and Brigita Urbanc. Do molecular dynamics force fields accurately model ramachandran distributions of amino acid residues in water? *Phys. Chem. Chem. Phys.*, **24**, 3259-3279, 2022.
- 4. **Brian Andrews**, Kaho Long, and Brigita Urbanc. Soluble state of villin headpiece protein as a tool in the assessment of MD force fields. *J. Phys. Chem. B*, **125(25)**, 6897–6911, Jul 2021.
- 3. B. Milorey, R. Schweitzer-Stenner, **B. Andrews**, H. Schwalbe, and B. Urbanc. Short peptides as predictors for the structure of polyarginine sequences in disordered proteins. *Biophys. J.*, **120**, 662–676, 2021.
- 2. S. Zhang, **B. Andrews**, R. Schweitzer-Stenner, and B. Urbanc. Intrinsic conformational dynamics of alanine in water/ethanol mixtures: An experiment-driven molecular dynamics study. *J. Phys. Chem. B*, **124(51)**, 11600–11616, 2020.
- 1. **B. Andrews**, S. Zhang, R. Schweitzer-Stenner, and B. Urbanc. Glycine in water favors the polyproline II state. *Biomolecules*, **10**, 1121, 2020.

## Contributed Talks at Conferences and Meetings

- 3. **B. Andrews**, S. Zhang, R. Schweitzer-Stenner, and B. Urbanc. Assessing the Ability of Molecular Dynamics Force Fields to Capture Conformational Dynamics of Amino Acid Residues in Water. APS March Meeting, 2022.
- 2. **B. Andrews**, S. Zhang, R. Schweitzer-Stenner, and B. Urbanc. Glycine Shows Preference for Polyproline II Indicating Greater Role for Amino Acid Backbone. APS March Meeting, 2021.
- B. Andrews, K. Long, and B. Urbanc. Examining the Self Assembly of the Villin Headpiece Protein: A Combined Experimental and Molecular Dynamics Study. APS Mid-Atlantic Section Fall Meeting, 2020.

## Poster Presentations

2. **B. Andrews**, T. Ruggiero, and B. Urbanc. Analyzing the Conformational Differences of the Intrinsically Disordered Amyloid  $\beta$ -Protein in Varying Lipid and Salt Concentrations. Drexel URCF Showcase, 2022.

1. **B. Andrews**, S. Zhang, R. Schweitzer-Stenner, and B. Urbanc. Hydrogel-forming Ultrashort Oligopeptides in Water/Ethanol Mixtures as a Potential Candidate for Oral Drug Delivery. Merck Emerging Talent Symposium, 2021.

#### Peer Review Contributions

Journal of Physical Chemistry Letters, Proteins, Biochemistry, Journal of Chemical Information and Modeling

### **Awards**

The Guoliang Yang Award, Drexel University, 2022.

# Experience

experiments.

**Systems Administrator**, *Drexel University*, Philadelphia, PA.

Responsible for upkeep of Physics department servers as well as lab computers.

Teaching Assistant, Drexel University, Philadelphia, PA.

Moderate recitations and lab exercises for students with various levels of physics backgrounds.

**Adjunct Instructor**, *Salem County Community College*, Salem, NJ. Taught introductory physics where I designed my own lectures and laboratory

Operations Intern, OptoQuest Inc., Cleveland, OH.

Created a machine learning model to predict postoperative risk profiles of patients undergoing eye surgeries.

Data Science Associate, MedaSync Inc., Cleveland, OH.

Constructed a self-updating machine learning cost prediction model in a production environment (AWS) which estimated medical costs based on diagnosis, projected length of stay, and comorbities.

Fall 2020-Present

Fall 2018-Spring 2022

Summer 2019

Summer 2017-Summer 2018

Fall 2017-Summer 2018

## Skills

Primary Programming Languages: Python, C, Bash, Mathematica, MATLAB, SQL

Software: GROMACS, AMBER, VMD, OpenMM

Computing: Highly Parallelized HPC at Drexel URCF, XSEDE at UT. Proficient with SLURM,

TORQUE, Univa Grid Engine. Proficient with distributed Windows and Linux systems. **Development Experience:** Git, AWS (EC2, S3, Lambda, MongoDB), Google Data Studio

# Open Source Software Contributions

OpenMM. Summer 2022

Add polarizable solvent directly to system. Pull request #3760: https://github.com/openmm/openmm/.

pdb-tools. Winter 2021

Pull Request #112: https://github.com/haddocking/pdb-tools.

# Extracurriculars

**Analytics Contributor and Writer**, *Pro Lacrosse Talk*, Remote. Contribute via analytics articles, modeling win probabilities using machine learning, and podcast appearances.

Summer 2020-Present