ANDREW MCNUTT

EDUCATION

Ph.D. Computational Biology

Expected 2024

Carnegie Mellon University-University of Pittsburgh

B.S. Physics and B.S. Mathematics, Minor: Chemistry

Purdue University

RESEARCH EXPERIENCE

Graduate Student Researcher

University of Pittsburgh

• Pittsburgh, PA

Advisor: David Koes

Aug 2019 - Current

- Analyzed convolutional neural network (CNN) scoring functions within molecular docking pipeline
- Developed Siamese CNN network to predict relative binding free energy of congeneric series ligands
- Open-sourced a docking pipeline to use pairwise statistics for improved docking performance

Student Intern at Regenstrief Institute

Center for Biomedical Informatics

Mentor: Shaun Grannis

May 2019 - Aug 2019

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- Engineered several machine learning approaches for record-linkage, identifying records that are from the same patient

- Analyzed and compared the performance of record-linkage techniques

Undergraduate Research Assistant at Purdue University

Department of MCMP

♥ West Lafayette, IN

Advisor: Markus Lill

- Analyzed a novel coarse-grained approach for modeling protein-ligand interaction
- Customized a random forest scoring function for coarse grained modeling of protein-ligand binding
- Enhanced a convolutional neural network with the ability to use probe-protein interaction data to better classify ligand binding poses

Student Intern at Indiana University-Purdue University Indianapolis

Division of Nephrology

◊ Indianapolis, IN

Advisor: Tarek M. Ashkar

- **May 2018 Aug 2018**
- Researched and determined the strengths and weaknesses of various approaches for clustering and dimensionality reduction
- Collaborated on the development of a full release version of a volumetric cell cytometry software for use on tissue images
- Expanded the capability of the cell cytometry software to include clustering and dimensionality reduction on the data created with the

Indiana University O'Brien Student Intern

Division of Nephrology

Indianapolis, IN

Advisor: Bruce A. Molitoris

May 2017 - Aug 2017

- Prototyped the merging of a 3D visualization software with an application developed for cell cytometry in tissue images
- Developed procedures for integrating native code into Java applications

SKILLS AND CLASSES

- Python, PyTorch, Pandas, NumPy, Java, Bash, Git
- Deep Learning, Computer Vision, Scalable Machine Learning, Metric Learning
- Molecular Dynamics, Docking, Pharmacophore Modeling
- Linear Algebra, Vector Calculus, Discrete Math, Probability
- Thermal Physics, Computational Physics, Quantum Mechanics
- Biochemistry, Genomics, Systems Biology

PUBLICATIONS

"Improving $\Delta \Delta G$ predictions with a multi-task convolutional Siamese Network"

McNutt, A. and Koes, D.

Journal of Chemical and Information Modeling (2022)

"GNINA 1.0: molecular docking with deep learning"

DOI:10.1186/s13321-021-00522-2

DOI:10.1021/acs.jcim.1c01497

McNutt, A.T., Francoeur, P., Aggarwal, R., Masuda, T., Meli, R., Ragoza, M., Sunseri, J. and Koes, D.R. Journal of Cheminformatics (2021)

"Integrated cytometry with machine learning applied to high-content imaging of human kidney tissue for in-situ cell classification and neighborhood analysis"

DOI:10.1101/2021.12.27.474025

Winfree, S., McNutt, A.T., Khochare, S., Borgard, T.J., Barwinska, D., Sabo, A.R., Ferkowicz, M.J., Williams, J.C., Lingeman, J.E., Gulbronson, C.J. and Kelly, K.J.,

bioRxiv (2021)

"In situ classification of cell types in human kidney tissue using 3D nuclear staining"

DOI:10.1002/cyto.a.24274

Woloshuk, A., Khochare, S., Almulhim, A.F., **McNutt, A.T.**, Dean, D., Barwinska, D., Ferkowicz, M.J., Eadon, M.T., Kelly, K.J., Dunn, K.W. and Hasan, M.A.

Cytometry Part A (2021)

PRESENTATIONS

"Open-ComBind: An open-source docking pipeline harnessing pairwise pose statistics" ACS Spring 2023

Mar 2023

"GNINA 1.0: molecular docking with deep learning"

Broad Institute: Machine Learning in Drug Discovery (Poster)

Oct 2022

Molecular Machine Learning Conference (Poster) ACS Spring 2021

Oct 2022 April 2021

"Exploring $\Delta\Delta G$ prediction with Siamese Networks"

Machine Learning for Structural Biology(MLSB) Workshop at NeurIPS (Poster)

Dec 2021

"Comparison of Supervised Machine Learning and Probabilistic Approaches for Record Linkage"

AMIA Summit 2020 (Not presented due to COVID-19)

March 2020

MENTORING

Mentoring Master's Student Project

University of Pittsburgh

- Advised a Master's student through a neural network knowledge distillation project
- Guided the development of a poster for the PhD program orientation and submitted an abstract to ACS Spring 2023

Summer Research Mentor

Student: Maddie Bonanno

University of Pittsburgh, TECBio Research Experience for Undergraduates

• Acted as primary research mentor for an undergraduate student completing graduate-level research in drug discovery

Advised project to completion; final work presented at university student research symposium

LEADERSHIP

Treasurer ## 2021-Current

Carnegie Mellon-University of Pittsburgh Computational Biology Graduate Student Association

- Recorded purchases made by the graduate student association
- Budgeted events to fit within our departmental allocations

Senator #2020-2021

Carnegie Mellon-University of Pittsburgh Computational Biology Graduate Student Association

- Organized several in-person and online events for the PhD program
- Assisted in transitioning our PhD program to virtual events during the COVID-19 pandemic

TEACHING

Intro to Computational Structural Biology

Graduate Teaching Assistant, University of Pittsburgh

Scalable Machine Learning for Big Data Biology

Graduate Teaching Assistant, University of Pittsburgh

Modern Mechanics

Undergraduate Teaching Assistant, Purdue University

Electric And Magnetic Interactions

Undergraduate Teaching Assistant, Purdue University

M Aug 2022-Dec 2022

Instructors: David Koes & James Faeder

Instructors: David Koes & Maria Chikina

🛗 Jan 2019-May 2019

Aug 2016-Nov 2018