Andrew C. Patt

☐ (518) 366 4293 • ☑ patt.14@osu.edu • ② andyptt21.github.io

Education

The Ohio State University **Expected Graduation: March 2021**

PhD in Biomedical Science, concentration in Biomedical Informatics, GPA: 3.813/4.0

Columbus, OH

State University of New York at Geneseo

May 2015

B.Sc. in Biochemistry, minor in Biomathematics, GPA: 3.660/4.0

Geneseo, NY

Awards

Finalist, Ruth L. Kirschstein National Research Service Award (NRSA) Individual Predoctoral Fellowship 2020

Received a potentially fundable score of 30 applying for NLM F31 fellowship, ultimately not funded

Systems and Integrative Biology Training Program Fellow

2017-2019

Competitive departmental T32 fellowship offered through Biomedical Sciences Graduate Program at Ohio State

American Society of Mass Spectrometry Travel Award

2018 and 2019

Travel stipend for student attendees of American Society of Mass Spectrometry annual conference

May Institute Travel Award

2018

Travel stipend for attendees of Northeastern University May Institute of mass spectrometry

Multi-Modeling and Integrative Data Analytics Training Program fellow

2016-2017

Departmental T15 fellowship offered through Biomedical informatics department at Ohio State

Best Poster Award

Awarded during NIAMS Scientific Planning Retreat

Summer Undergraduate Research Fellowship at University at Buffalo Funded by American Society for Pharmacology and Experimental Therapeutics 2014

2016

Publications

- 1. Patt, A.; Demoret, B.; Stets, C.; Bill, K.-L.; Smith, P.; Vijay, A.; Patterson, A.; Hays, J.; Hoang, M.; Chen, J.L.; Mathé, E.A. MDM2-Dependent Rewiring of Metabolomic and Lipidomic Profiles in Dedifferentiated Liposarcoma Models. Cancers 2020, 12, 2157.
- 2. An OpenData portal to share COVID-19 drug repurposing data in real time. Kyle R. Brimacombe, Tongan Zhao, Richard T. Eastman, Xin Hu, Ke Wang, Mark Backus, Bolormaa Baljinnyam, Catherine Z. Chen, Lu Chen, Tara Eicher, Marc Ferrer, Ying Fu, Kirill Gorshkov, Hui Guo, Quinlin M. Hanson, Zina Itkin, Stephen C. Kales, Carleen Klumpp-Thomas, Emily M. Lee, Sam Michael, Tim Mierzwa, Andrew Patt, Manisha Pradhan, Alex Renn, Paul Shinn, Jonathan H. Shrimp, Amit Viraktamath, Kelli M. Wilson, Miao Xu, Alexey V. Zakharov, Wei Zhu, Wei Zheng, Anton Simeonov, Ewy A. Mathé, Donald C. Lo, Matthew D. Hall, Min Shen. bioRxiv 2020.06.04.135046; doi: https://doi.org/10.1101/2020.06.04.135046
- 3. Eicher, T.; Kinnebrew, G.; Patt, A.; Spencer, K.; Ying, K.; Ma, Q.; Machiraju, R.; Mathe, E.A. Metabolomics and Multi-Omics Integration: A Survey of Computational Methods and Resources. Metabolites 2020, 5, 15.
- 4. Patt, A.; Eicher, T.; Kautto, E.; Machiraju, R.; Mathe, E.A.; Zhang, Y. Challenges in proteogenomics: a comparison of analysis methods with the case study of the DREAM Proteogenomics Sub-Challenge. BMC Bioinformatics 2019;20(Suppl 24):669.
- 5. Zhang, B.; Hu, S.; Baskin, E.; Patt, A.; Siddiqui, J.K.; Mathe, E.A. RaMP: A Comprehensive Relational Database of Metabolomics Pathways for Pathway Enrichment Analysis of Genes and Metabolites. Metabolites 2018, 8, 16.
- 6. Patt A.; Siddiqui J.; Zhang B.; Mathe E.A. (2019) Integration of Metabolomics and Transcriptomics to Identify Gene-Metabolite Relationships Specific to Phenotype. In: Haznadar M. (eds) Cancer Metabolism. Methods in Molecular Biology, vol 1928. Humana Press, New York, NY

Presentations

Metabolomics Society Annual Meeting

November 2020

Online Improving pathway analysis of lipidomic and metabolomic data through comprehensive functional annotation and network approaches

1/4

Metabolomics Association of North America Annual Meeting September 2020 Ann Arbor, MI (virtual) Improving pathway analysis of lipidomic and metabolomic data through comprehensive functional annotation and network approaches American Society for Mass Spectrometry June 2020 Poster Houston, TX (virtual) MDM2 copy number aberrations alter ceramide glycosylation in liposarcoma tumors, impacting drug response Wexner Medical Center Trainee Research Day **April 2020** Poster Columbus, OH MDM2 copy number aberrations alter ceramide glycosylation in liposarcoma tumors, impacting drug response Ohio Mass Spectrometry and Metabolomics Symposium October 2019 Poster Columbus, OH MDM2 copy number aberrations alter ceramide glycosylation in liposarcoma tumors, impacting drug response International Conference on Intelligent Biology and Medicine June 2019 20 minute talk Columbus, OH Challenges in proteogenomics: a comparison of analysis methods with the case study of the DREAM Proteogenomics Sub-Challenge American Society for Mass Spectrometry June 2019 Atlanta, GA MDM2 Alterations Reprogram the Metabolic Functioning of Liposarcoma Cell Lines Ohio State University Cancer Care Center Annual Meeting May 2019 Columbus, OH MDM2 Alterations Reprogram the Metabolic Functioning of Liposarcoma Cell Lines American Society for Mass Spectrometry June 2018 Poster San Diego, CA MDM2 alterations reprogram the metabolic functioning of liposarcoma cell lines Ohio Mass Spectrometry and Metabolomics Symposium **April 2018** Poster Columbus, OH MDM2 Alterations Reprogram the Metabolic Functioning of Liposarcoma Cell Lines June 2017 **National Library of Medicine Informatics Training Conference** 5-minute talk La Jolla, CA MDM2 alterations reprogram the metabolic functioning of liposarcoma cell lines Wexner Medical Center Trainee Research Day **April 2017** Columbus, OH MDM2 alterations reprogram the metabolic functioning of liposarcoma cell lines **NIAMS Scientific Planning Retreat April 2016** Bethesda, MD Deep Resequencing of Juvenile Arthritis Patients Identifies Rare Variant Associations Postbac Poster Day **April 2016** Bethesda, MD Poster Deep Resequencing of Juvenile Arthritis Patients Identifies Rare Variant Associations SUNY Geneseo GREAT Day **April 2015** Poster Geneseo, NY Sex Differences in MCT1 Tissue Expression University at Buffalo Summer Research Symposium August 2014 20 minute talk and poster Buffalo. NY Sex Differences in MCT1 Tissue Expression

Research Experience

National Center for Advancing Translational Sciences

March 2020-Present

Research Associate

Rockville, MD (remote position)

Working with Dr. Ewy Mathe in the Bioinformatics core at the National Center for Advancing Translational Sciences. Contracted through Axle Informatics.

- Continued projects from OSU including providing guidance for updates of RaMP database/R package.
- Developing a method for improved functional enrichment analysis of metabolomic data (including pathway analysis) using biological pathway and chemical structure knowledge graphs.
- Contributing to identifying new, publicly available 'omics data generated in the study of COVID-19. See data portal: https://opendata.ncats.nih.gov/covid19/omics

The Ohio State University, Biomedical Informatics Department

August 2016-Present

Graduate Research Assistant

Columbus, OH

Working with Dr. Ewy Mathe in the Biomedical Informatics department through the Biomedical Sciences Graduate Program.

- Investigating metabolomic and transcriptomic differences between dedifferentiated liposarcoma tumors based on copy number of the MDM2 oncogene.
- Assisted in developing RaMP metabolic pathways database/R package and Rshiny app.
- Worked primarily with the R programming language as well as Bash, MySQL and Python.
- Ran memory-intensive scripts on the Ohio Supercomputing Cluster.

National Institute of Arthritis, Musculoskeletal and Skin Diseases

July 2015-July 2016

Postbaccalaureate Fellow

- Bethesda, MD
- Worked in the lab of Dr. Michael Ombrello through NIH postbaccalaureate program. • Searched for ultrarare genetic variants associated with Systemic Juvenile Idiopathic Arthritis in a targeted resequencing project.
- Used Illumina sequencing kits as well as GATK and annovar softwares.
- Worked with R, Excel and Bash for data sorting, manipulation and analysis.
- Gained experience working with supercomputing clusters.

University at Buffalo, Pharmaceutical Sciences Department Summer Undergraduate Research Fellow

May 2014-August 2014

Summer undergraduate research fellowship funded by the American Society of Pharmacology and Experimental Therapeutics. Worked with Dr. Marilyn Morris and Dr. Melanie Felmlee in the Pharmaceutical Science department at University at Buffalo.

- Performed RNA extraction/qPCR using rat tissue
- Recieved weekly coaching on oral/poster presentations of research
- Discovered a significantly lower level of monocarboxylate transporter expression in the livers of female rats per gram of tissue, potentially leading to differences in metabolism of monocarboxylate compounds.

State University of New York at Geneseo, Biology Department

September 2013-May 2015

Undergraduate Research Assistant

Geneseo, NY

Buffalo, NY

- Worked in the Biomathematics research interest group of Dr. Gregg Hartvigsen, learning basic biological network modeling in the R programming language.
- Worked with Dr. Isidro Bosch identifying species of starfish larvae samples using targeted sequencing.

Work Experience

Pace Analytical June 2013-January 2014

Lab Assistant

Schenectady, NY

- Organized/analyzed chemical and supply purchasing data
- Organized stock room
- Performed sample delivery
- General lab support

Pace Analytical January 2013-May 2013

Assistant Lab Instructor Geneseo, NY

Instructed single-semester human biology lab, aided students with experiments and lab reports.

Aqueduct Rowing Club

May 2012-August 2012

Adult Learn-to-Row Coach

Schenectady, NY

Coached two 5-week sessions of an introductory rowing course. Required to devise and execute practice routines to instruct novice adult sweep rowers.

Volunteering and Leadership

Friday Afternoon Bioinformatics Workshops

January 2019-June 2019

Organizer

Columbus, Ohio

Organized and presented at a series of seminars that provided overviews of useful software tools for bioinformaticians, including Git, Bash, Docker, Tidyverse, Emacs and mzMine

Locust Grove Nature Center

August 2015-November 2015

Nature Center Greeter Bethesda, MD

Helped orient visitors, aided in park events, tracked park usage and maintained trail and visitor center on weekends.

Wadeable Assessment by Volunteer Evaluation Program

May 2012-August 2012

WAVE Volunteer

Collected samples of Benthic Macroinvertebrates and performed an amateur assessment of stream health from various wadeable stream habitats to aid in a large scale evaluation of freshwater ecosystems health in upstate New York.

Relevant Coursework

Graduate.....

Northeastern May Institute

May 2018

Computation and Statistics for Mass Spectrometry and Proteomics (week-long workshop)

The Ohio State University

August 2016-October 2018

- Introduction to Biomedical Informatics
- Analysis and Applications of Genome-Scale Data
- Seminars in Bioinformatics
- Data Structures and Algorithms
- Probability for statistical inference
- Metabolomics, Principles and Practice (audited)

Foundation for Advanced Education in the Sciences

September 2015-May 2016

- Practical Bioinformatics
- Introduction to Python

Undergraduate.....

State University of New York at Geneseo

September 2011-May 2015

- Lab: Biochemistry, Molecular Techniques, Organic Chemistry, Biology, Plant Physiology, Analytical Physics
- Biology: Microbiology, Cell Biology, Genetics, Ecology
- Chemistry: Organic Chemistry, Biochemistry, Physical Chemistry
- Math/Physics: Calculus 1,2,3, Linear Algebra, Probability, Differential Equations, Biostatistics, Modeling Biological Systems, Analytical Physics

Relevant Skills

- Programming Languages: R, Python, Bash, MySQL
- o Mass Spec Data Analysis: mzMine, XCMS, LipidMatch, Metaboanalyst, RaMP, MS-DIAL, Excel
- o Genomic Data Analysis: Picard, GATK, Annovar
- Wet Lab: RNA extraction and purification (Qiagen), Illumina MiSeq library preparation (Nextera Rapid Capture Custom Enrichment kits)
- o Development Tools: Emacs, Linux, RStudio, Git/Github/Gitlab, RShiny, RMarkdown, Tidyverse, Devtools
- Public Speaking: Throughout my undergraduate and graduate careers I have given a large number of oral presentations
 describing my work, from small group settings to major international conferences.
- **Technical Writing:** I have been the primary author/coauthor on three peer-reviewed manuscripts, as well as secondary author on three more. I have also been the primary submitter on several grant proposals, including a proposal for a departmental T32 award which funded two years of my tuition/stipend at OSU, and an F31 proposal which recieved a potentially fundable score of 30.

Society Memberships

- Metabolomics Society: 2020-Present
- Metabolomics Association of North America: 2020-Present
- o American Society for Mass Spectrometry: 2018-Present