# Jennifer Williams

Curriculum Vitae

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#### **Publications**

Conference

Same cause; different effects in the brain.

Proceedings

M. Toneva\*, J. Williams\*, A. Bollu, C. Dann, L. Wehbe.

Causal Learning and Reasoning, CLeaR 2022

Preprints

Behavior measures are predicted by how information is encoded in an individual's brain.

J. Williams, L. Wehbe.

arXiv 2021 (arXiv:2112.06048)

Discriminative subtyping of lung cancers from histopathology images via contextual deep Learning.

B.J. Lengerich, M. Al-Shedivat, A. Alavi, **J. Williams**, S. Labbaki, E.P. Xing. *medRxiv* 2020 (DOI: 10.1101/2020.06.25.20140053)

Journal Papers

LSD1 dual function in mediating epigenetic corruption of the vitamin D signaling in prostate cancer.

S. Battaglia, E. Karasik, B. Gillard, **J. Williams**, T. Winchester, M.T. Moser, D.J. Smiraglia, B.A. Foster.

Clinical Epigenetics 2017 (DOI: 10.1186/s13148-017-0382-y)

#### Education

Expected 2022

**PhD in Computational Biology**, Carnegie Mellon University (CMU).

Advisor: Dr. Leila Wehbe, Machine Learning Department

Topic: Machine Learning Method Development to Model Individual Differences in the Brain in the Presence and Absence of Disease

2013-2016

Cancer and Systems Biology EU-USA Atlantis Dual Degree Program.

Master of Science in Natural Science, Roswell Park, University at Buffalo.
 Advisor: Dr. Barbara Foster

Topic: Computational and Experimental Interrogation of Vitamin D's Role in Prostate Cancer

- Master of Science in Integrated Systems Biology, University of Luxembourg.
- Courses Towards MSc, University of Amsterdam and VU University Amsterdam.

2013 Bachelor of Science in Biology, Canisius College. Magna Cum Laude.

## Relevant PhD Coursework and Computing Skills

Languages

Python, R, MATLAB, Java, LATEX

Libraries

Scikit-learn, Pandas, NumPy, Pycortex

Courses

Cognitive Neuroscience, Machine Learning, ABCDE of Statistical Methods in Machine Learning, Intermediate Statistics, Probabilistic Graphical Models, Computational Medicine

## Fellowships

2020-2021

Digital Health Fellowship - Center for Machine Learning and Health (CMLH)

• Full tuition and stipend for 12 months and \$3,000 for research-related expenses

2017-2019

NIH T32 Training Grant - National Institute of Biomedical Imaging and Bioengineering

• Full tuition and stipend for 2 years and \$6,000 for research-related expenses

- 2013-2015 CanSys MS Scholarship Atlantis EU-USA Training Program
  - Stipend for 12 months
- 2012-2013 Canisius Earning Excellence Program (CEEP) Research Grant

### Current Research Projects

#### Graduate Research Assistant - CMU

#### Predicting Behavior From How Information is Encoded in an Individual's Brain.

 Creating a framework to identify individual differences in how information is encoded in the brain, and testing if these differences predict behavior measures. Utilizing functional connectivity methods and encoding/decoding models with visual and language semantic features to analyze neuroimaging data.

#### Creating Interpretable Integrated Analysis of Imaging and Genomic Data.

 Collaborating with a cross-functional team to develop a deep learning framework to incorporate complex covariate data (e.g. histology images) into an interpretable graphical model framework.

#### Discovering How Individual Differences in a Stroke Impact Vision Recovery.

 Extending causal mediation methods to model how fMRI activity and a stroke lesion impact a participant's visual field recovery.

#### Identifying Individual Brain Differences in Healthy and Depressed Participants.

 Developing algorithms to model individual differences in time series fMRI or EEG data elicited by naturalistic stimuli (audio and videos) that are clinically relevant.

#### Conference Posters and Oral Presentations

- 2022 Behavior Measures are Predicted by How Information is Encoded in an Individual's Brain. Computational and Systmes Neuroscience (COSYNE), poster
- 2021 Behavior Measures are Predicted by How Information is Encoded in an Individual's Brain. Neuromatch Conference 4 (NMC4), poster
- 2021 Leveraging Brain Encoding Models to Predict Individual Differences in Behavior. Society for Neuroscience (SFN), poster
- 2019 All Terrain Graph-Learner-CPC: Causal Model Discovery for Missing Not at Random Data. Intelligent Systems for Molecular Biology/European Conference on Computational Biology (ISMB/ECCB), poster
- 2018 Contextual Explanation Networks Enable Integrated Analysis of Imaging and Genomic Data. ISMB, poster
- 2018 Learning Causal Models from Missing Not at Random Data in Clinical Datasets.

  NIH Training Grantees Meeting of the National Institute of Biomedical Imaging and
  Bioengineering, poster
- 2013 Preliminary Results on the Prevalence of Physiology Students' Homeostatic Misconceptions. Experimental Biology (EB) poster
- 2012 Mentoring: A Collaboration Between Higher Education and Urban After School Outreach Programs. Science Teachers Association of New York State Conference (STANYS), oral presentation

#### **Awards**

- 2021 Top Reviewer Machine Learning for Health (ML4H) Conference
  - Top 10 reviewer out of several hundred

- 2019 Attended 2019 Machine Learning Summer School (MLSS) Moscow to collaborate and learn state of the art techniques to advance machine learning research
- 2019 Carnegie Mellon University Graduate Student Assembly/Provost Conference Award
- 2019 International Conference on Machine Learning (ICML) Travel Award
- 2018 CMU-Pitt Computational Biology (CPCB) PhD Program Student Service Award
- 2013 Canisius College Biology Department Excellence in Science Education Award
- 2012 Beta Beta Beta (Biological Honor Society National Inductee)

## Selected Previous Research Experiences

#### Graduate Research Assistant - CMU

#### Extending Causal Model Discovery for Missing Not at Random Data (MNAR).

 Proposed, developed and implemented a causal learning algorithm to learn a causal graph when data is MNAR.

#### Graduate Research Assistant - Roswell Park

#### Interrogated Vitamin D's Role in Prostate Cancer.

- Profiled the vitamin D receptor and lysine-specific demethylase 1 (LSD1) interactomes
  using Prostate Cancer clinical samples and data I collected from cell lines and mice.
- Thesis published by ProQuest, "Inhibiting LSD1 restores vitamin D responsiveness in castration resistant prostate cancer", 2016.

### Relevant Service and Leadership

#### 2021-present **BrAIn Seminar Co-organizer**.

Co-organized a weekly seminar at the intersection of neuroscience and AI research

#### 2021-present CPCB Steering Committee Student Representative.

Collaborated with senior leadership in the CPCB PhD program to decide strategic priorities

#### 2018-present **Peer Reviewer**.

ML4H Conference 2021

New In ML @NeurIPS Workshop 2020-2021

Nature Scientific Reports

ECCB 2018

#### 2016-present **TechNights Volunteer**.

Created and led workshop on image processing for middle school girls Assisted teaching other STEM topics through interactive workshops

#### 2020 Graduate Application Support Program Mentor.

Provided feedback to potential CMU PhD program applicants

#### 2017-2020 CPCB Graduate Student Assembly.

President (2019-2020)

Vice President (2018-2019)

Senator (2017-2018)

## 2017-2019 TECBio REU (Training and Experimentation In Computational Biology Research Experiences for Undergraduates).

Co-led a journal club for undergraduate researchers (2019)

Ethics mentor for undergraduate researchers (2018)

Moderated a journal club for undergraduate researchers (2017 - 2018)

#### 2019 Convestro Pittsburgh Regional Science & Engineering Fair Judge.

Leader of a 5 judge team in Medicine, Health, and Microbiology category

#### 2018-2019 Admissions Committee Member CPCB PhD program.

Reviewed PhD program applications and assisted making admissions decisions

#### 2018 Intel International Science and Engineering Fair (ISEF) Grand Award Judge.

## Teaching and Mentoring

2019-2020 Mentored Undergraduate Student.

William Yang - CMU Computer Science

2017-2018 **Teaching Assistant**, Carnegie Mellon University.

Course: Lab Methods for Computational Biologists.

Planned and taught module on computational sequence analysis