

Amy X. Lu

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SUMMARY

I am a graduate student at the University of Toronto/Vector Institute for AI. I am interested in **developing biologically-principled machine learning methods**, with an emphasis on **self-supervised representation learning, generalizability, and biological interpretability**. I also serve on Research to the People, which organizes research collaborations of genomics/ML researchers for patients of understudied diseases.

EDUCATION

- **University of Waterloo** Waterloo, Canada
Bachelors of Science, Honours Science, Bioinformatics Option 2014 – 2019
- **University of Toronto, Vector Institute** Toronto, Canada
Master's in Computer Science Jan 2019 – Present
- **Stanford University** Palo Alto, USA
Visiting Student, Departments of Computer Science and Genetics Sept 2019 – Dec 2019

EXPERIENCE

- **Stanford University** Palo Alto, USA
Visiting Student Researcher — Domain Adaptation in Regulatory Genomics, Dr. Anshul Kundaje 2019 – Present
 - **Domain adaptation, ChIP-seq data, interpretability:** Using domain adaptation methods to improve transcription factor binding prediction when evaluating for a different cell line.
- **University of Toronto/Vector Institute** Toronto, Canada
Masters Student — Representation learning in genomics, Dr. Alan Moses & Dr. Marzyeh Ghassemi 2019 – Present
 - **Self-supervised learning, natural language processing:** Adapting attention-based NLP models to learn self-supervised representations of protein sequences.
 - **Generalizability, computer vision:** Benchmarked self-supervised and canonical computer vision methods in a microscopy image dataset with covariate shift to highlight generalization failures in machine learning. (*NeurIPS 2019; ICML 2019 Self-Supervised Learning Workshop*)
 - **Algorithmic fairness, clinical decision support:** Quantitative and qualitative evaluation of bias in “black box” contextual word embeddings on clinical notes. Explored debiasing methods using adversarial gradient-reversal. (*NeurIPS 2019 Workshop for ML for Healthcare, NeurIPS 2019 Workshop on Fair ML.*)
- **Harvard Medical School/Boston Children's Hospital** Boston, USA
Research Intern — Machine Learning in Clinical Genomics, Dr. Piotr Sliz 2018 – 2019
 - **Genotype-phenotype studies, clinical interpretability:** Understanding disease genotype-phenotype relationships using machine learning models. Interpreted important model features to seek novel disease-associated variants from whole exome (WES) data.
 - **Variant filtering, pathway analyses:** Applied standard filtering pipelines for false-positive variants. Explored classification from pathway- and variant-level features.
 - **Machine learning, dimensionality reduction:** Explored methods for capturing epistatic non-linearities and statistical dimensionality reduction.
- **University of Waterloo** Waterloo, Canada
Undergraduate Thesis Student — Deep Learning in Regulatory Genomics, Dr. Andrew Doxey 2017 – 2018
 - **Convolutional neural networks in genomics:** Implemented a convolutional neural network which classifies ATAC-seq accessible regions with 91% accuracy.
 - **Interpretability:** Interpreted the neural network by reconstructing the first-layer feature map as position-weighted matrix (PWM) motif. All motifs found statistical matches in JASPAR, a database of known motifs.
 - **Phylogenetics, metagenomic data mining:** Used various bioinformatics pipeline tools (HMMER, BLAST, etc.) to understand biochemical properties of potentially uncharacterized toxins in metagenomic data.
 - **Self-directed learning:** Self-learned deep learning concepts as a biology student.

- **École polytechnique fédérale de Lausanne** Lausanne, Switzerland
Research Intern — Molecular Dynamics Simulations, Dr. Matteo Dal Parero 2017
 - **Molecular dynamics:** Used molecular dynamics (MD) and GROMACS to simulate enzyme-membrane mechanisms of bacterial “superbugs”.
 - **Workflow automation:** Analyzed and automated simulations using R, tcl/tk, and Csh scripts.
- **University of Toronto** Toronto, Canada
Research Intern — Data Visualization in Pharmacoepidemiology, Dr. Suzanne Cadarette 2015 – 2017
 - **Data visualization, pharmacoepidemiology:** Used R to visualize the social diffusion of methodological innovation. Contributed to two co-authored publications and an international epidemiology conference.

AWARDS

- **NSERC Canada Graduate Scholarships-Master’s (CGS-M) Award:** NSERC federal research grant. Valued at \$17,500.
- **NSERC Michael Smith Foreign Supplement:** Supports high-calibre Canadian graduate students in pursuing research abroad. Valued at \$6,000.
- **Scholarship of Excellence in Research:** One of 13 students sponsored for research at EPFL. Valued at CHF 4,500.
- **University of Waterloo:** Various entrance awards totaling \$6,000.
- **Associate of The Royal Conservatory (ARCT):** Performer’s Diploma in Piano.

PUBLICATIONS AND CONFERENCE PROCEEDINGS

- **Accepted:** Zhang H*, **Lu AX***, Abdalla M, McDermott M, Ghassemi M. Hurtful Words: Quantifying Biases in Clinical Contextual Word Embeddings. *Spotlight Talk, ACM Conference on Health, Inference, and Learning (CHIL) 2020. *Equal contribution.*
- **Published:** Lu AX, **Lu AXP**, Schormann W, Ghassemi M, Andrews DW, Moses AM. The Cells Out of Sample (COOS) dataset and benchmarks for measuring out-of-sample generalization of image classifiers. *Poster, Neural Information Processing Systems (NeurIPS) 2019.*
- **Presented:** Moses A, Lu AX, **Lu AXP**, Ghassemi M. Transfer Learning vs. Batch Effects: what can we expect from neural networks in computational biology? *Poster, Machine Learning for Computational Biology (MLCB) 2019.*
- **Published:** Ban J, Tadrous M, **Lu AX**, Cicinelli EA, Cadarette SM. Diffusion of indirect comparison meta-analytic methods to study drugs: a systematic review and co-authorship network analysis. *BMJ Open.*
- **Under revision:** Consiglio GP, Maclure M, **Lu AX**, Cicinelli EA, Ban JK, McCarthy L, Cadarette SM. Guidance documents for the application of Self-controlled Crossover Observational PharmacoEpidemiology (SCOPE) designs are needed: systematic review and animated co-authorship networks. *Pharmcoepi Drug Saf.*

WORKSHOP PAPERS AND POSTERS

- **Presented:** **Lu AX***, Zhang H*, Abdalla M, McDermott M, Ghassemi M. Hurtful Words: Quantifying Biases in Clinical Contextual Word Embeddings. Hurtful Words: Quantifying Biases in Clinical Contextual Word Embeddings. *Poster presentation, NeurIPS 2019 Workshop on Fair ML for Health. *Equal contribution.*
- **Presented:** Zhang H*, **Lu AX***, Abdalla M, McDermott M, Ghassemi M. Hurtful Words: Quantifying Biases in Clinical Contextual Word Embeddings. Hurtful Words: Quantifying Biases in Clinical Contextual Word Embeddings. *Extended Abstract, NeurIPS 2019 Workshop on Machine Learning for Healthcare. *Equal contribution.*
- **Presented:** Abdalla M, Zhang H, **Lu AX**, Chen I, Ghassemi M. Quantifying Fairness in a Multi-Group Setting and its Impact in the Clinical Setting. *Poster presentation, NeurIPS 2019 Workshop on Fair ML for Health.*
- **Presented:** Lu AX, **Lu AXP**, Moses A. Paired Cell Inpainting: A Multiple-Instance Extension of Self-Supervised Learning for Bioimage Analysis. *Poster presentation, ICML 2019 Workshop on Self-Supervised Learning.*
- **Presented:** **Lu AX**, Rockowitz S, Poduri A, Sliz P. From data to precision medicine: predictive machine learning models to uncover disease-associated variants. *Poster. Harvard Medical School BCMP Retreat 2019.*
- **Submitted:** **Lu AX**, Tam ES. Effect of prophylactic brimonidine on subconjunctival hemorrhage in laser-assisted cataract surgery. *Poster abstract.*
- **Presented:** **Lu AX**, Consiglio GP, Cadarette SM. Dynamic Visualization in Co-Authorship Network Analysis. *Poster presentation. Leslie Dan Faculty of Pharmacy Undergraduate Research Symposium.*
- **Presented:** McIlroy-Young R, **Lu AX**, Guenther N, Olarnyk A. A Network Analysis: Did the Arab Spring Impact the Academic Network of Tunisia Between 2010 and 2015? *Poster presentation. Knowledge Integration Symposium.*

SERVICE AND ACTIVITIES

- **Research to the People** (*formerly SVAI*): Core Team of Research to the People, a non-profit connecting patients of rare genomic diseases to the medical/AI research community and industry partners through collaborative research initiatives.
- **Residence Don:** Organized events and established rapport with diverse students. Responded to mental health and conduct crises. Leader for the Velocity Residence, a spin-off for the Velocity start-up incubator.
- **Tosamaganga Hospital, Tanzania:** Supported operations at a rural Tanzanian hospital and shadowed surgical procedures. Expenses were covered by scholarships, fundraising, and part-time tutoring.

TEACHING

- **Teaching Assistant, Genetics:** Taught weekly tutorial lectures for BIOL 239 at the University of Waterloo.
- **Piano, music theory:** Taught piano performance (up to RCM 7), ear training, RCM music history, and RCM Intermediate Rudiments.
- **Tutoring:** Tutor for IB Math and IB Chemistry.

TALKS

- **Vector NLP Talks:** Quantifying and Removing Biases in Clinical Contextual Word Embeddings. *Co-presenter.*
- **Harvard Medical School BCMP Retreat 2019:** From data to precision medicine: predictive machine learning models to uncover disease-associated variants. *Lightning talk.*

SELECTED HACKATHON PROJECTS

- **Chatsense:** Mobile chat app using a sentiment analysis machine learning API on voice messages and mapping dominant emotions to emojis, designed for the autism community. *Team project. “Can a Computer Hear How You Feel?” prize: YHacks at Yale.*
- **Alice:** Web login page using a trained facial recognition model to login to online banking without passwords, designed for dementia patients. *Team project. Desjardin prize: ConUHacks III.*