Amy X. Lu

http://amyxlu.github.io
github.com/amyxlu

amyxlu@cs.toronto.edu
+1-647-968-6833
linkedin.com/in/amyxlu

SUMMARY

I am a graduate student at the Vector Institute for AI, applying **computer vision and natural language processing algorithms to biological sequence data**, with an emphasis on **self-supervised representation learning, robustness, and biological interpretability**. I also serve on SVAI, which organizes research collaborations of genomics/AI researchers and industry partners for understudied rare disease patients.

EDUCATION

- **University of Waterloo** — Waterloo, Canada
  Bachelors of Science, Honours Science, Bioinformatics Option — *2014 – 2019*

- **University of Toronto, Vector Institute** — Toronto, Canada
  Masters, Computer Science — *2019 – Present*

EXPERIENCE

- **Stanford University** — Palo Alto, USA
  Visiting Student Researcher — Domain Adaptation in Regulatory Genomics, Dr. Anshul Kundaje — *2019 – Present*
  - **Domain adaptation, ChIP-seq analyses:** Using domain adaptation to improve classifier performance and learned genomic sequence features for transcription factor binding when evaluating in a different cellular context.

- **University of Toronto/Vector Institute** — Toronto, Canada
  Masters Student — Representation learning in genomics, Dr. Alan Moses & Dr. Marzyeh Ghassemi — *2019 – Present*
  - **Natural language processing, self-supervised learning:** Applying contextual word embedding models to learn robust and interpretable representations for genomic and proteomic sequences.
  - **Algorithmic fairness:** Quantitative and quaitative evaluation of bias in "black box" contextual clinical word embeddings, and attempting to debias using adversarial gradient-reversal. (*Submitted to FAT* 2020*)
  - **Generalizability, computer vision:** Benchmarked a new "paired cell in-painting" method for out-of-sample generalization in microscopy against various common computer vision algorithms. (*NeurIPS 2019; ICML 2019 Self-Supervised Learning Workshop*)

- **Harvard Medical School/Boston Children's Hospital** — Boston, USA
  Research Intern — Deep Learning in Clinical Genomics, Sliz Lab — *2018 – Present*
  - **Variant analysis, genotype-phenotype studies:** Applying standard filtering pipelines to a whole exome (WES) dataset of epilepsy patients
  - **Machine learning, dimensionality reduction:** Building machine learning models to classify disease phenotype from high-dimensional exome variant data

- **University of Waterloo** — Waterloo, Canada
  Undergraduate Thesis Student — Deep Learning in Regulatory Genomics, Doxey Lab — *2017 – 2018*
  - **Convolutional neural networks in genomics:** Implemented a convolutional neural network which recognizes genomic enhancers regulating femur growth with 91% accuracy.
  - **Interpretability:** Distilled biological knowledge from the "black box" network by reconstructing first-layer features as position-weighted matrices (PWMs) of "motifs" (i.e. genomic patterns). All motifs found statistical matches in JASPAR (an experimentally-verified motifs database).
  - **Self-directed learning:** Self-learned deep learning concepts and Python as a biology student.

- **École polytechnique fédérale de Lausanne** — Lausanne, Switzerland
  Research Intern — Computational Biology, Laboratory for Biomolecular Modelling — *2017*
  - **Molecular dynamics:** Contributed three potential findings on enzyme-membrane mechanisms of bacterial "superbugs" using computational simulations based on molecular dynamics (MD) principles.
  - **Automating workflows:** Increased workflow efficiency by writing R, tcl/tk, and Csh scripts.

- **University of Toronto** — Toronto, Canada
  Research Intern — Data Visualization in Pharmacoepidemiology, Cadarette Lab — *2015 – 2017*

- ○ **Pharmacoepidemiology:** Took initiative to redesign manual workflows. Contributed two co-authored publications, one poster, and figures for an international epidemiology conference within two months.
- ○ **Data visualization:** Wrangled data using R and SQL to create animated and interactive co-authorship networks.

Service and Activities

- **SVAI:** Core Team of SVAI, a non-profit connecting patients of rare genomic diseases to the medical/AI research community and industry partners through collaborative research initiatives.
- **Residence Don, Velocity Incubator:** Leader for the residence program of Canada's most productive start-up incubator; fostered close relations with diverse students to support conduct and mental health crises
- **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 Assistant:** Taught weekly tutorial lectures for a 200-level genetics course.

Awards

- **NSERC Michael Smith Foreign Supplement**: Supports high-calibre Canadian graduate students in pursuing research abroad. Valued at $6,000.
- **Canada Graduate Scholarships-Master's (CGS-M) Award**: NSERC federal research grant. Valued at $17,500.
- **Scholarship of Excellence in Research**: One of 13 students sponsored for research at EPFL. Valued at CHF 4500.
- **University of Waterloo**: Various entrance awards totaling $6000.
- **Associate of The Royal Conservatory (ARCT)**: Performer's Diploma in Piano (highest academic standing offered).

Publications and Conference Proceedings

- **Accepted:** Lu AX, **Lu AXP**, Moses A. The Cells Out of Sample (COOS) dataset and benchmarks for measuring out-of-sample generalization of image classifiers. ***NeurIPS 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. ***Submitted to Pharmcoepi Drug Saf.***

Workshop Papers

- Lu AX, **Lu AXP**, Moses A. Paired Cell Inpainting: A Multiple-Instance Extension of Self-Supervised Learning for Bioimage Analysis. ***ICML 2019 Workshop on Self-Supervised Learning.***

Posters

- **Presented: Lu AX**, Rockowitz S, Poduri A, Sliz P. From data to precision medicine: predictive machine learning models to uncover disease-associated variants. Poster presentation, November 2019. ***Harvard Medical School BCMP Retreat 2019.***
- **Presented: Lu AX**, Consiglio GP, Cadarette SM. Dynamic Visualization in Co-Authorship Network Analysis. Poster presentation, August 2016. ***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, April 2015. ***Knowledge Integration Symposium.***
- **Submitted: Lu AX**, Tam ES. Effect of prophylactic brimonidine on subconjunctival hemorrhage in laser-assisted cataract surgery. Poster abstract. ***Submitted to American Academy of Ophthalmology 2017.***

Hackathons and Projects

- **NLP-seq:** Used n-grams to extract features for supervised learning of function prediction, trained on Protein Data Bank data.
- **Chatsense:** Mobile chat app using a sentiment analysis machine learning API on voice messages and mapping dominant emotions to emojis, designed for the autistic community. ***Team project. Winner at 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. Winner at ConUHacks III.***