

# Louis Cammarata

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## EDUCATION

### Harvard University

*PhD, Statistics*

Dissertation topic: Network modeling and machine learning with applications to genomics

Advised by Professor Tracy Ke (Harvard Statistics) and Professor Caroline Uhler (MIT EECS)

Cambridge, MA  
Expected May 2023

### Massachusetts Institute of Technology, Institute for Data, Systems, and Society

*MSc, Technology and Policy*, 2018 Best Thesis Award

Cambridge, MA

May 2018

### École Polytechnique

*MSc, Biological Engineering* (ranked 34<sup>th</sup> out of 506)

*BSc, Mathematics and Sciences*

Palaiseau, FRANCE

May 2016

May 2015

## FELLOWSHIPS & AWARDS

**Bok Center Statistics Pedagogy Fellowship**, Harvard University

2021

**Bok Center Certificate of Distinction in Teaching Award**, Harvard University

2019, 2020, 2021

**Keystone Symposia Travel Fellowship**, Keystone Symposia

2019

**Technology & Policy Program Best Thesis Award**, Massachusetts Institute of Technology

2018

**Outstanding Investment Award**, École Polytechnique

2015

## PUBLICATIONS & PREPRINTS

### Statistical Theory & Methods

**Cammarata, L.**, Jiang, K., Jin, J. and Ke, Z.T. (2022). Estimating Dynamic Mixed Memberships by Trajectory Embedding. Manuscript in preparation.

- Develop dynamic mixed-membership estimation algorithm using spectral methods in the degree-corrected mixed membership stochastic block model and apply algorithm to international trade network dynamics
- Manuscript 80% complete (available upon request)

**Cammarata, L.** and Ke, Z.T. (2022). Power Enhancement and Phase Transitions for Global Testing of the Mixed Membership Stochastic Block Model. *Bernoulli*.

- Derived fundamental statistical limit of global detection of network communities (i.e., testing whether network has more than one community) and designed a novel statistically optimal hypothesis test
- 79-page article with proofs

### Applied Statistics & Machine Learning

**Cammarata, L.**, Shivashankar, G.V. and Uhler, C. (2022). Adhesome Receptor Clustering is Accompanied by the Co-localization of the Associated Genes in the Cell Nucleus. Manuscript in preparation.

- Characterize map between protein assemblies on cell membrane and 3D organization of genome in nucleus, specifically for adhesome in IMR90 fibroblasts using multiple genomic/epigenomic data modalities (RNA-seq, ChIP-seq, in situ Hi-C)
- Project 90% complete (available upon request)

Belyaeva\*, A., **Cammarata\***, L., Radhakrishnan\*, A., Squires, C., Yang, K.D., Shivashankar, G.V. and Uhler, C. (2021). Causal network models of SARS-CoV-2 expression and aging to identify candidates for drug repurposing. *Nature Communications*.

- Developed drug repurposing pipeline using overparametrized representation learning, network analysis and causal structure learning and applied it to identify *in silico* candidate drug targets for treatment of COVID-19
- Media coverage in [MIT News Office](#) and other specialized online media
- Co-first authors marked with \*

Zhang, J., **Cammarata\***, L., Squires\*, C., Sapsis, T.P. and Uhler, C. (2022). Active Learning for Optimal Intervention Design in Causal Models. *arXiv*.

- Designed active learning procedure to push system from source state to target state when this system can be described with a structural causal model (relevant applications in cell reprogramming)
- Equal contribution marked with \*

Delaney\*, C., Schnell\*, A., **Cammarata\*, L.**, Yao-Smith, A., Regev, A., Kuchroo, V.K. and Singer, M. (2019). COMET: a tool for marker-panel selection from single-cell transcriptomic data. *Molecular Systems Biology*.

- Developed COMET, a computational tool for combinatorial marker detection from single-cell transcriptomic data available as an [online platform](#) and [Python package](#) and leveraged tool to identify B cell subpopulations
- Co-first authors marked with \*

## RELATED PROFESSIONAL EXPERIENCE

### PathAI

*Biomedical Data Science PhD Intern*

*Boston, MA*

*May-Aug 2022*

- Executed project to characterize novel biomarkers of tumor micro-environment using deep learning model in Python in collaboration with team of computational biologists and expert pathologists and used markers for prognosis prediction

### Google

*Data Science PhD Intern (gTrade)*

*Virtual/Mountain View, CA*

*May-Aug 2020*

- Developed statistical model to predict competition price on Google Ads Exchange using R and SQL on massive data
- Implemented model and sent it for experiments on small percentage of display adds traffic to assess conversion performance

### Goldman Sachs

*Summer Quantitative Analyst (Franchise Analytics team)*

*New York City, NY*

*Jun-Aug 2017*

- Designed statistical similarity model to score US corporate bonds and support relative value trading operations

## TEACHING EXPERIENCE

### Harvard University

*Statistics Teaching Fellow*

*Cambridge, MA*

*Sep 2019 – May 2022*

- Taught STAT303 – The Art and Practice of Teaching and Communicating Statistics (with Prof. Joe Blitzstein and Prof. Morgane Austern) to first-year Statistics PhD students in Fall 2021 and Spring 2022
- Taught STAT210 – Probability I (Fall 2019, Fall 2020 with Prof. Joe Blitzstein), STAT212 – Probability II (Spring 2020 with Prof. Subhabrata Sen) and STAT131 – Time Series (Spring 2020 with Prof. Tracy Ke) to classes of 30-80 students

**The Jubilee Institute** | MIT Science and Technology Initiative Global Teaching Lab

*Amman, JORDAN*

*High School Mathematics Teacher*

*Jan 2017*

- Designed and taught 5-session introductory "Randomness & Probability" course to 6 cohorts of 25 students (10<sup>th</sup> grade)

## ON-CAMPUS LEADERSHIP

**Harvard GSAS Consulting Club** | Harvard University

*Cambridge, MA*

*Vice President of 2022 Harvard/MIT Case Competition*

*Sep 2021–Present*

- Organize 12-day flagship event involving 100+ participants in cooperation with consulting firms and industry experts

**Harvard Statistics Graduate Committee** | Harvard University

*Cambridge, MA*

*Vice President*

*Sep 2019–Present*

- Coordinate yearly Statistics PhD students retreat for ~70 students, faculty and affiliates in Harvard's Statistics Department
- Organize mentoring of ~10 first-year Statistics PhD students by senior graduate students

**PolitiX (student non-partisan policy club)** | École Polytechnique

*Palaiseau, FRANCE*

*President*

*Jan 2015–Mar 2016*

- Organized monthly conferences with politicians, CEOs and academics for 500+ students – "Outstanding Investment Award"

## SELECTED CONFERENCE PRESENTATIONS

**Cammarata, L.** Adhesome receptor clustering is accompanied by the co-localization of adhesome genes in the cell nucleus. Oral presentation delivered at the Broad Institute Cell Circuits and Epigenomics Seminar Series, Cambridge, MA, June 2022.

**Cammarata, L.** Network Global Testing: Phase Transitions and Power Enhancement. Oral presentation delivered at the Harvard Mini-Workshop on Algebraic Statistics, Cambridge, MA, April 2022.

**Cammarata, L.** Drug Repurposing Pipeline Using Overparametrized Representation Learning and Causality. Oral presentation delivered at Purdue University Data Science Week, West Lafayette, IN (virtual), November 2021.

**Cammarata, L.** Drug Repurposing Pipeline Using Overparametrized Representation Learning and Causality. Oral presentation delivered at Barcelona Supercomputing Center (BSC), Barcelona, Spain (virtual), September 2021.

**Cammarata, L.** Phase Transitions in Network Global Testing. Oral presentation delivered at Joint Statistical Meetings (JSM), virtual, August 2021.

**Cammarata, L.** and Belyaeva, A. Causal Network Models of SARS-CoV-2 Expression and Aging to Identify Candidates for Drug Repurposing. Oral presentation delivered at Machine Learning in Computational Biology (MLCB), virtual, November 2020.

**Cammarata, L.** COMET: Combinatorial Marker Detection from Single-Cell Transcriptomic Data. Spotlight oral presentation delivered at Machine Learning in Computational Biology (MLCB), Vancouver, Canada, December 2019.

**Cammarata, L.** COMET: Combinatorial Marker Detection from Single-Cell Transcriptomic Data. Poster presented at NeurIPS, Vancouver, Canada, December 2019.

**Cammarata, L.** COMET: Combinatorial Marker Detection from Single-Cell Transcriptomic Data. Poster presented at Keystone Symposia on Cancer Immunotherapy: Mechanistic Insights to Improve Clinical Benefits (C2), Whistler, Canada, March 2019.

**Cammarata, L.** Inferring Total Queueing Time Using Only Elapsed Time to Date. Oral presentation delivered at INFORMS Annual Meeting, Houston, TX, October 2017.