Sahil Loomba

Contact 77 Massachusetts Avenue Information

Building E18-404 Cambridge, MA 02139, US +18572182159sloomba@mit.edu https://sloomba.github.io

Research Interests Design and analysis of experiments, causal inference, statistical network modeling, behavioral science, computational social science, public health, machine learning, Bayesian inference

EMPLOYMENT

Massachusetts Institute of Technology, US

Schmidt Science Fellow at the Institute for Data, Systems, and Society October 2023 -PIs: Prof Dean Eckles, MIT Sloan; Prof Elchanan Mossel, MIT Mathematics

Imperial College London, UK

EPSRC Doctoral Prize Fellow in Mathematics October 2022 – September 2023 October 2018 - December 2019 Graduate Teaching Assistant in Mathematics

Harvard University, US

Predoctoral Fellow in Systems Biology at the Wyss Institute August 2016 - July 2018

Xerox Research Center India, India

Research Intern in Multimedia Analytics

May - July 2015

EDUCATION

Imperial College London, UK

PhD in Mathematics PI: Prof Nick S Jones October 2018 - September 2022

Thesis: Sparse and partially observed large-scale networks: Analytic statistics, behaviour, and structural inference

Indian Institute of Technology Delhi, India

BTech in Computer Science and Engineering

2012 - 2016

PI: Prof Sumeet Agarwal

Thesis: Causal computational models for gene regulatory networks

Honors

Yael Naim Dowker Prize for the best Mathematics PhD thesis	2024
Schmidt Science Fellowship	
for the world's best emerging scientists to pursue a postdoctoral fellowship	2023
EPSRC Doctoral Prize Fellowship	
for the very best newly qualified PhDs receiving EPSRC doctoral funding	2022
Doris Chen Mobility Award	
for PhD students with exceptional potential to take their research abroad	2022
Imperial-TUM Global Fellows Programme	2019
Cargill Global Scholars Award for academic excellence & leadership	2014
IIT Delhi Institute Medal for obtaining highest overall GPA	2013
IIT Delhi Merit Semester Scholarship	2012, 2013

Publications

• Rubin, J., Loomba, S., & Jones, N. S. (2023, November). Geodesic Distributions Reveal How Heterophily and Bottlenecks Limit the Expressive Power of Message Passing Neural Networks. The Second Learning on Graphs Conference, https://openreview.net/forum? id=PEVln6psEH.

We provide a statistical perspective on MPNN performance by decomposing it into signal and noise sensitivity and showing that the former relates to higher-order homophily.

• Loomba, S.*, Maertens, R., Roozenbeek, J., Götz, F., van der Linden, S., & de Figueiredo, A. (2023). Ability to detect fake news predicts sub-national variation in COVID-19 vaccine uptake across the UK. medRxiv preprint, doi:10.1101/2023.05.10.23289764.

^{*}Corresponding author

Using a psychometrically-validated test of misinformation susceptibility, we show that belief in fake news is credibly associated with lower actual COVID-19 vaccine uptake.

 Götz, F., Maertens, R., Loomba, S., & van der Linden, S. (2023). Let the Algorithm Speak: How to Use Neural Networks for Automatic Item Generation in Psychological Scale Development. *Psychological Methods*, doi:10.1037/met0000540

We leverage recent advances in transformer-based neural language models that generate human-like text, like GPT, for automatic item generation for psychological scales.

Kite, K. A., Loomba, S., Elliott, T. J., Yongblah, F., Lightbown, S. L., Doyle, T. J., Gates, L., Alber, D., Downey, G. A., McCurdy, M. T., Hill, J. A., Super, M., Ingber, D. E., Klein, N., & Cloutman-Green, E. (2022). FcMBL magnetic bead-based MALDI-TOF MS rapidly identifies paediatric blood stream infections from positive blood cultures. PLOS ONE, doi:10.1371/journal.pone.0276777

We show that broad-range pathogen binding properties of an engineered FcMBL enable more accurate identification of patient positive cultures than the state-of-the-art.

• Loomba, S., & Jones, N. S. (2021). Geodesic statistics for random network families. arXiv preprint, arxiv:2111.02330. (74 pages)

We derive an analytic distribution of shortest path lengths in sparse conditionally independent edge models, yielding new results on percolation and path-based centralities.

• Loomba, S.[†], de Figueiredo, A.[†], Piatek, S. J., de Graaf, K., & Larson, H. J. (2021). Measuring the impact of COVID-19 vaccine misinformation on vaccination intent in the UK and USA. *Nature Human Behaviour*, 5(3), 337-348, doi:10.1038/s41562-021-01056-1.

Through a randomized study, we show that even brief exposure to COVID-19 vaccine misinformation can reduce the intention to definitely vaccinate by as much as 6.2%.

Novak, R., Lin, T., Kaushal, S., Sperry, M., Vigneault, F., Gardner, E., Loomba, S., Shcherbina, K., Keshari, V., Dinis, A., Vasan, A., Chandrasekhar, V., Takeda, T., Turner, J., Levin, M., & Ingber, D. (2022). Target-agnostic discovery of Rett Syndrome therapeutics by coupling computation and a CRISPR-enabled in vivo disease model. biorXiv preprint, doi:10.1101/2022.03.20.485056.

We combine computational modeling with *in vivo* screening in a CRISPR-edited *Xenopus* laevis tadpole model of Rett syndrome to carry out target-agnostic drug discovery.

• Gandhi, A., Biswas, A., Shrivastava, K., Kumar, R., **Loomba, S.**, & Deshmukh, O. (2016, March). Easy Navigation through Instructional Videos using Automatically Generated Table of Content. In *Companion Publication of the 21st International Conference on Intelligent User Interfaces* (pp. 92-96). ACM, doi:10.1145/2876456.2879472.

We demonstrate a pipeline to automatically generate a table of content for video lectures, using saliency scores of visual and spoken words for topic-based segmentation.

PATENTS

Cartwright, M. J., Super, M., Ingber, D. E., Grant, J., Scott, J., Duffy, S. C., & Loomba,
S. (2021). Methods for detection of microbes and microbe components. *International Publication No. WO 2021/201949 A2*.

Technology licensed to *BOA Biomedical*, with the aim of rapid detection of active infection from a blood sample within 1 hour, which is crucial for treating sepsis.

Barhate, S. S., Loomba, S., Gandhi, A., Biswas, A., Negi, S., & Deshmukh, O. D. (2019).
Method and system for generation of a table of content by processing multimedia content.
U.S. Patent No. US 10,296,533 B2.

Technology licensed to VideoKen for automated video content navigation

 Gandhi, A., Biswas, A., Deshmukh, O. D., & Loomba, S. (2018). Method and system for content processing to determine pre-requisite subject matters in multimedia content. U.S. Patent Publication No. US 2018/0060984 A1.

Technology licensed to VideoKen to create course structures for effective learning

Grants Syn

Synergistic Discovery and Design (SD2)

DARPA/I20, HR001117S0003, \$2M

PI: Prof James J Collins

Co-authored research proposal for data-driven discovery and design of biological circuits. Contributed about half of each of the writing, conceptual and methodological content.

September 2017

[†]Equal contribution

Designed core algorithmic pipeline to generate testable circuit designs: going from high-throughput omics data to inferred network structures, to functional network motifs, down to operable biological circuits in traditional and novel host organisms.

Developed sequence and circuit embedding models of E. coli and yeast for motif discovery.

Presentations

Oral

- Loomba, S. (2024, March). Social covariation and spillovers using geo-aggregated network data. Talk at *MIT Sloan Marketing Seminar*.
- Loomba, S.[†], Happenhofer, J.[†], Hoffmann, T., Agarwal, S., Jones, N. S. (2023, July). Inferring socio-physical networks and social Gini indices from aggregated data. Oral presentation at NetSci Satellite Symposium on Statistical Inference for Network Models.
- Loomba, S. (2023, June). Fake news and global health. Panel discussion at *The London University Diplomatic Summit*.
- Loomba, S.[†], Happenhofer, J.[†], Hoffmann, T., Agarwal, S., Jones, N. S. (2022, July). Global mapping of a "social Gini" index using socio-physical connectivity kernels inferred from large-scale social network data. Oral presentation at 8th International Conference on Computational Social Science.
- Loomba, S. (2021, November). Misinformation and the pandemic: a tale of two contagions. Invited talk at Students for Global Health Conference 2021.
- Loomba, S., & Jones, N. S. (2021, July). Geodesic statistics for random network families. Oral presentation at Networks 2021 (Joint Sunbelt and NetSci Conference).
- Loomba, S., & Jones, N. S. (2020, July). Scalable statistics for social networks of entire societies. Oral presentation at *Sunbelt XL Conference*.
- Loomba, S., & Jones, N. S. (2020, March). Stochastic block models as a social connectivity kernel. Talk at CMPH Seminar, Imperial College London.
- Loomba, S. (2018, August). Mathematical Representations For Biological Systems. Talk at *DAIR Seminar*, *Indian Institute of Technology Delhi*.

Poster

- Loomba, S. (2023, July). Causal inference in sparse partially observed large-scale networks. Poster Presentation at *International School and Conference on Network Science*.
- Loomba, S. (2023, July). Causal inference in sparse and partially observed large-scale social networks. Poster Presentation at 9th International Conference on Computational Social Science.
- Loomba, S., Hoffmann, T., & Jones, N. S. (2019, July). Social Access Statistic: Linking Social Connectivity to Health Outcomes. Poster presentation at 5th International Conference on Computational Social Science.
- Loomba, S., & Garrod, M. (2019, July). How Far Would You Go? Comparing Urban Access in 10 Global Cities. Poster presentation at *NetMob 2019*.

Impact

Generated viable drug candidate(s) for treatment of a monogenic disease

Created NeMoCAD, a Network Model for Causally Aware Discovery, that learns from generegulation and perturbative drug-gene interaction data to query for desirable therapeutic states; discovered a new therapeutic for Rett Syndrome which was validated in mouse models. $Software\ licensed\ to\ Unravel$ Biosciences

Impacted public health policy on health misinformation

Research on the impact of COVID-19 vaccine misinformation on vaccine uptake has seen considerable impact on pandemic research (1600+ citations), and on policymaking: the US Surgeon General's health misinformation advisory cites our work, and I delivered a video message for the UN on the impact of misinformation on our ability to end the pandemic.

POPULAR PRESS

UN Verified Is online misinformation threatening our ability to end the pandemic?

Reuters Misinformation could prompt people to turn against COVID-19 vaccines: study

BBC Covid vaccine: Rumours thrive amid trickle of pandemic facts

The Times Herd immunity put at risk by a few lies about Covid vaccine

The Quint Exposure to Misinformation Reduces COVID Vaccine Acceptance: Study

Hindustan Times Misinformation may hit vaccination drive: Study

Ongoing Research

Causal inference in partially observed networks

Collaborators: Dean Eckles (MIT Sloan), Elchanan Mossel (MIT Mathematics)

We are describing a statistical theory for inference of causal effects in the case of networked treatment units wherein inter-unit connections are only partially observed. Grounded in Boolean functional analysis, our goal is to allow applied researchers to design optimal experiments for inferring the causal estimand of their interest, like higher order spillover effects.

Disentangling dyadic and triadic social network effects

Collaborators: Dean Eckles (MIT Sloan), Arnab Sarker (MIT IDSS)

Higher-order interactions in social networks are often conflated with dyadic interactions but can produce drastically different network dynamics. Using a large-scale survey sample, we elicit data on triadic interactions between individuals to understand the social and spatial mechanisms that drive them, and study their qualitative differences from dyadic interactions.

Cross-cultural impact of misinformation on beliefs about democratic values, climate change, and vaccine acceptance

Collaborators: Simon J. Piatek & Heidi J. Larson (LSHTM), Martin Wiegand (UCL)

Through a large-scale randomized trial in Nigeria, Germany, and Poland, we have studied the impact of misinformation exposure on a variety of important attitudes. In particular, we find that misinformation negatively impacts belief in liberal democractic values across all three countries, with individual-level heterogeneity in the impact across political affiliations.

Mapping inequality in access to social connections via a "social Gini" index

Collaborators: Nick S. Jones & Johannes Happenhofer (Imperial), Till Hoffmann (Harvard), Sumeet Agarwal (IIT Delhi)

We have developed a rational theory for a "social Gini" index which quantifies the distance between people based on how surprising it is to see a social connection between them. Our aim is to map this index sub-nationally for the UK, US, and India. For that, we are building a Bayesian statistical model that learns from data on spatially aggregated friendship counts.

Teaching

Introduction to applied Bayesian inference (Imperial-X)	April 2023
Developed and delivered two lectures and coursework problems:	January – March 2023
Methods for Data Science (Years 3 & 4, MSc)	

Graduate teaching (tutorials)

Methods for Data Science (Years 3 & 4, MSc)	October – December 2019
Mathematical Computation in Python (Year 1)	January - March 2019
Probability and Statistics II (Year 2)	October – December 2018

SERVICES

Reviewing	
Physical Review E	2019 -
Scientific Reports	2019 -
New Journal of Physics	2021 -
Proceedings of the National Academy of Sciences	2021 -
Vaccine	2021 -
Social Network Analysis and Mining	2021 -
Healthcare in Low-resource Settings	2022 -
Mathematical Modelling of Natural Phenomena	2022 -
Science	2022 -
Epidemics	2023 -
Journal of Physics: Complexity	2023 -

PC Member

SFE-TSDM at IEEE International Conference on Data Mining 2021

Advisees	Jonathan Rubin Amelia French Kushal Kejriwal Siddharth Chandak Katie Collins	PhD in Computing, Imperial College London Year 4, MSci in Mathematics, Imperial College London Year 4, MSci in Mathematics, Imperial College London Year 2, Dual degree in Electrical Engineering, IIT Bombay Year 2, BTech in Electrical Engineering, IIT Bombay Year 1, Bachelors in Brain and Cognitive Sciences, MIT	2023 - 2022 - 2023 2021 - 2022 2021 2019 2017
Programming Skills	0	entific computing: Python, R, MATLAB, C++, C statistical inference frameworks: Stan, TensorFlow, PyTorch,	scikit-learn