

Sahil Loomba

CONTACT INFORMATION	301A Sir Ernst Chain Building Imperial College London, South Kensington London SW7 2AZ, UK	+44 7721 538305 s.loomba18@imperial.ac.uk https://sloomba.github.io
RESEARCH INTERESTS	Network science, Bayesian inference, statistical machine learning, computational social science, computational & systems biology, public health statistics	
EXPERIENCE	Wyss Institute for Biologically Inspired Engineering at Harvard University Post-baccalaureate Fellow in Computational Biology August 2016 – July 2018 Advisor: Prof. James J. Collins, MIT	
	Xerox Research Center India Research Intern in Multimedia Analytics May – July 2015	
EDUCATION	Imperial College London, United Kingdom PhD in Applied Mathematics 2018 – August 2022 (expected) Advisor: Prof. Nick S. Jones	
	Indian Institute of Technology Delhi, India BTech in Computer Science and Engineering 2012 – 2016 Advisor: Dr. Sumeet Agarwal Thesis: Causal Computational Models for Gene Regulatory Networks	
ACADEMIC AND COMPETITIVE HONORS	Imperial-TUM Global Fellows Programme Summer Undergraduate Research Award, IIT Delhi Cargill Global Scholar Award for academic excellence & leadership IIT Delhi Institute Medal for obtaining highest overall GPA IIT Delhi Merit Semester Scholarship	2019 2014 2014 2013 2012, 2013
PUBLICATIONS	<ul style="list-style-type: none">• Loomba, S., & Jones, N. S. (2021). Geodesic statistics for random network families. <i>arXiv preprint</i>, 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.¹, de Figueiredo, A.¹, 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. <i>Nature Human Behaviour</i>, 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%.• 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 <i>Companion Publication of the 21st International Conference on Intelligent User Interfaces</i> (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. <p><i>Forthcoming</i></p> <ul style="list-style-type: none">• Götz, F., Maertens, R., Loomba, S., & van der Linden, S. (<i>Revise & resubmit at Psychological Methods</i>). Development and Validation of an Algorithmically-Derived Personality Scale.	

¹Co-first authors: SL contributed to survey design, did RCT experiment design, designed all statistical models and performed all inference, generated publication figures, co-drafted the paper and responded to referees

We leverage recent advances in transformer-based neural language models that generate human-like text, like GPT-2, for automatic item generation for psychometric scales.

- Duffy, S., **Loomba, S.**, Cartwright, M., Dimitrakakis, N., Scott, J., McCarty, A., Shapiro, N. I., Super, M., & Ingber, D. (*In prep*). Application of MALDI-TOF mass spectrometry for the rapid identification of PAMPs in blood culture negative septic patients using FcMBL-coated magnetic beads.

We develop a rapid diagnostic test from the mass spectra of patient blood/urine samples, using a probabilistic model that leverages the mass spectra of a microbial library.

PATENTS

- Cartwright, M. J., Duffy, S. C., Grant, J., Ingber, D. E., **Loomba, S.**, Scott, J., & Super, M. (2021). Methods for detection of microbes and microbe components. *U.S. Patent Application No. PCT/US21/12717*.

Technology has been 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. (2018). Method and system for generation of a table of content by processing multimedia content. *U.S. Patent Application No. 15/203,868*.
- 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 Application No. 15/250,958*.

GRANTS

Synergistic Discovery and Design (SD2)

DARPA/I20, HR001117S0003, \$2M

September 2017

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.

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.

CONFERENCE PRESENTATIONS

Invited talks

- **Loomba, S.** (2021, November). Misinformation and the pandemic: a tale of two contagions. Invited talk at *Students for Global Health Conference 2021*.

Accepted oral presentations

- **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*.

Accepted poster presentations

- **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 gene-regulation 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 (340+ citations in less than a year), and on policymaking (the U.S. Surgeon General’s health misinformation advisory cites our work).

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

Online informational interventions for improved offline vaccine uptake

Collaborators: Amit Bahl (Facebook CDS), Heidi Larson & Alexandre de Figueiredo (LSHTM)

As part of an internship with Facebook CDS (2022), we will design online interventions likely to improve COVID-19 vaccine uptake, run a large-scale randomized trial on Facebook to gauge their causal impact, and study “spillover” effects on peer outcomes. This precedes the goal of closing the loop on monitor-design-intervene for building vaccine confidence online.

Regional relationship of misinformation susceptibility and vaccine uptake

Collaborators: Friedrich Götz (UBC), Rakoén Maertens, Jon Roozenbeek & Sander van der Linden (Cambridge), Alexandre de Figueiredo (LSHTM)

Building on the misinformation susceptibility test (MIST), we have collected data on 16k survey respondents across the UK, and are building a Bayesian spatial model of MIST scores that accounts for spillover between “well-connected” regions. Post-stratification will help identify regional “hotspots” of susceptibility, and study the relationship to regional uptake.

Mapping inequality in access to social connections via a “social Gini” index

Collaborators: Nick S. Jones, Johannes Happenhofer & Till Hoffmann (Imperial), 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.

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 – present

Nature Scientific Reports

2019 – present

New Journal of Physics

2021 – present

Proceedings of the National Academy of Sciences

2021 – present

Vaccine

2021 – present

Social Network Analysis and Mining

2021 – present

Healthcare in Low-resource Settings

2022 – present

PC Member

SFE-TSDM at IEEE International Conference on Data Mining

2021

MENTORING	Amelia French	Year 4, MSci in Mathematics, Imperial College London	2021
	Kushal Kejriwal	Year 2, Dual degree in Electrical Engineering, IIT Bombay	2021
	Siddharth Chandak	Year 2, BTech in Electrical Engineering, IIT Bombay	2019
	Katie Collins	Year 1, Bachelors in Brain and Cognitive Sciences, MIT	2017

PROGRAMMING	Programming: Python, R, MATLAB, C, C++		
STRENGTHS	Machine learning and statistical tools: Stan, TensorFlow, PyTorch, scikit-learn		