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RESEARCH INTERESTS Statistical network modeling, asymptotic network properties, Bayesian inference, computational social science, systems biology, public health statistics, spatial statistics, survey methods

EXPERIENCE

Imperial College London, United Kingdom

EPSRC Doctoral Prize Fellow in Mathematics

October 2022 – present

Wyss Institute at Harvard University, United States

Post-baccalaureate Fellow in Computational Biology

August 2016 - July 2018

Advisor: Prof James J. Collins, MIT

Xerox Research Center India, India

Research Intern in Multimedia Analytics

May - July 2015

EDUCATION

Imperial College London, United Kingdom

PhD in Applied Mathematics

October 2018 – September 2022

Advisor: Prof Nick S. Jones

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

Advisor: Dr Sumeet Agarwal

Thesis: Causal computational models for gene regulatory networks

ACADEMIC AND COMPETITIVE HONORS 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

• Loomba, S.¹, 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.

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

¹First and corresponding author

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.², 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. *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 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. (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

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.

Presentations

Oral presentations

- Jones, N. S., Loomba, S., Marshall, A., Happenhofer, J. (2023, April). Applied Bayesian inference. Panel discussion at *Imperial-X*.
- 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

²Co-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

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 presentations

- 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 (1000+ 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

Online informational interventions for improved offline vaccine uptake

Collaborators: Amit Bahl (Meta CDS), Heidi Larson & Alexandre de Figueiredo (LSHTM), Nick S. Jones (Imperial), Sumeet Agarwal (IIT Delhi)

We are designing interventions to improve vaccine uptake on a national scale via a large-scale randomized trial to gauge their causal impact, and gather evidence to motivate the study of network spillover effects using privacy-preserving spatially aggregated data. This precedes the goal of closing the loop on monitor-design-intervene for building vaccine confidence.

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

	*	cal model that learns from data on spatially	,	0
TEACHING	Developed and delive	ed Bayesian inference (Imperial-X) red two lectures and coursework problems: ience (Years 3 & 4, MSc)	April 2023 January – Marc	h 2023
Graduate Teaching (Tutorials)		ience (Years 3 & 4, MSc) utation in Python (Year 1) istics II (Year 2)	October – Decen January – Marc October – Decen	h 2019
SERVICES	Vaccine Social Network Analy Healthcare in Low-re Mathematical Model Science PC Member	ational Academy of Sciences ysis and Mining	2019 2021 2021 2021 2021 2022 2022	 present
MENTORING	Jonathan Rubin Amelia French Kushal Kejriwal Siddharth Chandak Katie Collins	Year 4, MSci in Mathematics, Imperial C Year 4, MSci in Mathematics, Imperial C Year 2, Dual degree in Electrical Engineer Year 2, BTech in Electrical Engineering, I Year 1, Bachelors in Brain and Cognitive	ollege London ing, IIT Bombay IIT Bombay	2022 - 2023 2021 - 2022 2021 2019 2017

Programming and scientific computing: Python, R, MATLAB, C++, C

Machine learning and statistical inference frameworks: Stan, TensorFlow, PyTorch, scikit-learn

Programming Strengths

aim is to map this index sub-nationally for the UK, US, and India. For that, we are building