

# JACQUELINE R. M. A. MAASCH

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EDUCATION	05.2026	<b>Cornell Tech   New York, NY</b> Doctor of Philosophy in Computer Science (anticipated)
	05.2024	MS in Computer Science, conferred on PhD candidacy   GPA 4.0 Areas: AI / ML, Scientific Computing, Applied Probability & Statistics <i>NSF Graduate Research Fellow   Presidential Life Science Fellow</i>
	05.2021	<b>University of Pennsylvania   Philadelphia, PA</b> Master of Computer & Information Technology   GPA 3.97 <i>Interdisciplinary Innovation Fellow   Reproducible Research Fellow</i>
	05.2016	<b>Smith College   Northampton, MA</b> BA Anthropology (Biological, Medical), Environmental Science   GPA 3.97 <i>Summa Cum Laude   Phi Beta Kappa   Sigma Xi</i>
EXPERIENCE	05.2024 – 08.2024	<b>Research Intern</b> <i>Microsoft Research (MSR) Machine Intelligence Core   Cambridge, UK</i> <i>PI: <a href="#">Dr. Aditya Nori</a>, <a href="#">Dr. Javier González</a>.</i> Novel methods in AI reasoning.
	05.2022 – 08.2022	<b>Clinical Data Science Intern</b> <i>Boehringer Ingelheim Global Biostatistics &amp; Data Sciences   Ridgefield, CT</i> <i>PI: <a href="#">Dr. Yi Liu</a>.</i> Multimodal deep learning methods for survival analysis in pharmaceutical development.
	08.2021 – Present	<b>PhD Student Researcher</b> <i>Weill Cornell Medicine <a href="#">Institute of AI for Digital Health</a>   New York, NY</i> <i>PI: <a href="#">Dr. Fei Wang</a>.</i> AI for clinical risk modeling, causal inference, target trial emulation, and computational biomedicine.
		<i>Cornell Tech Operations Research   New York, NY</i> <i>PI: <a href="#">Dr. Kyra Gan</a>.</i> Robust and efficient statistical inference, scalable causal discovery, and causal fairness in healthcare.
		<i>Cornell Tech Computer Science   New York, NY</i> <i>PI: <a href="#">Dr. Volodymyr Kuleshov</a>.</i> Core problems in generative and probabilistic modeling with applications to genomics and biomedicine.
	05.2020 – 07.2021	<b>Master's Student Researcher</b> <i>University of Pennsylvania Bioengineering   Philadelphia, PA</i> <i>PI: <a href="#">Dr. César de la Fuente</a>.</i> DOD-funded research on discriminative and generative ML for antibiotic discovery.
INTERESTS	Probabilistic graphical models; reasoning in generative models; causal discovery / graph structure learning; causal inference; graph theory; applied probability.	
LANGUAGES	<i>Proficient:</i> Python; R; $\text{\LaTeX}$ . <i>Prior experience:</i> Java; C; JavaScript; MATLAB.	
TOOLS	<i>Frequently using:</i> sklearn; numpy; tidyverse; git; high-performance computing. <i>Prior experience:</i> PyTorch; TensorFlow; Stan.	

SELECT FELLOWSHIPS & AWARDS	2023	Cornell Tech Service and Community Award
	2021	NSF Graduate Research Fellowship
	2021	Presidential Life Science Fellowship   Cornell
	2021	Reproducible Research Fellowship   OKFN, Alfred P. Sloan Foundation
	2020	Grace Hopper Celebration Scholarship   UPenn
INVITED TALKS	10.24	INFORMS Annual Meeting   Seattle, WA (forthcoming) <i>Local Causal Discovery for Structural Evidence of Direct Discrimination</i>
	07.24	Microsoft Research Machine Intelligence Core   Cambridge, UK
	06.24	University of Cambridge Statistical Laboratory   Cambridge, UK <i>Local causal discovery for effect estimation</i>
	04.24	34th Annual POMS Conference   Minneapolis, MN <i>Local Discovery by Partitioning</i>
WORKSHOP PRESENTATIONS	2023	<b>Maasch, J;</b> et al. Local Discovery by Partitioning: Polynomial-Time Causal Discovery Around Exposure-Outcome Pairs. <i>NeurIPS Causal Representation Learning Workshop</i> . [ARXIV]
	2023	<b>Maasch, J;</b> et al. Regularized Data Programming with Automated Bayesian Prior Selection. <i>ICML Workshop on Structured Probabilistic Inference &amp; Generative Modeling</i> . [ARXIV]
IN PREPARATION & UNDER REVIEW	2024	Kuleshov, V; <b>Maasch, J;</b> Ermon, S. <i>Probabilistic Graphical Models: A Concise Tutorial</i> . In preparation for <i>Foundations &amp; Trends in Machine Learning</i> .
	2024	<b>Maasch, J;</b> et al. <i>Local Causal Discovery for Structural Evidence of Direct Discrimination</i> . Under review. [ARXIV]
	2024	Hiremath, S; <b>Maasch, J;</b> et al. <i>Hybrid Global Causal Discovery with Local Search</i> . Under review. [ARXIV]
SELECT PEER-REVIEWED PUBLICATIONS (GOOGLE SCHOLAR)	2024	<b>Maasch, J;</b> et al. <i>Local Discovery by Partitioning: Polynomial-Time Causal Discovery Around Exposure-Outcome Pairs</i> . Proceedings of the 40th Conference on Uncertainty in Artificial Intelligence. [ARXIV] [SLIDES] [POSTER]
	2024	Pan, W; Su, C; <b>Maasch, J;</b> et al. <i>Learning Phenotypic Associations for Parkinson's Disease with Longitudinal Clinical Records</i> . AMIA Summits on Translational Science Proc. American Medical Informatics Assoc. [NCBI]
	2023	<b>Maasch, J*</b> ; Torres, M*; et al. <i>Molecular de-extinction of ancient antimicrobial peptides enabled by machine learning</i> . Cell Host & Microbe. 31. 8. 1260-1274. e6. 2023. *Equal contribution. [CELL]
	2022	Melo, M*; <b>Maasch, J*</b> ; et al. <i>Machine Learning for Drug Discovery</i> . 2022. American Chemical Society. *Equal contribution. [GOOGLE BOOKS]
PROFESSIONAL ACTIVITIES	2024	Co-organizer, NYC Learning on Graphs Conference
	2024	Student leader, Cornell CS PhD Visit Days; Reviewer, PhD Admissions
	2023	Co-developer, Cornell CS 6006: Succeeding in the Graduate Environment
	2023	Founder / organizer, Cornell Causal Reading Group
PATENTS	2022	Co-Inventors: de la Fuente-Nunez C, Torres M, Melo M, <b>Maasch J</b> . Title: <i>Identification of antimicrobial peptides</i> . Docket no: 104377.000299 / 23-10289. Application no: 63/383,761. Pending.
PEER REVIEW	AI/ML	AISTATS; ACL Rolling Review; ICML SPIGM; NeurIPS WiML.
	Bio	Communications Biology (Nature Portfolio); Bioinformatics (Oxford Academic); ACS Infectious Diseases.