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EDUCATION **Carnegie Mellon University**, Pittsburgh, PA

Ph.D. in Machine Learning

August 2013 - May 2019

▷ Advisor: Seyoung Kim

GPA: 3.80

▷ Selected Courses: Probabilistic Graphical Models, Convex Optimization, Foundations of Machine Learning Theory, Graduate Molecular Biology

University of Michigan, Ann Arbor, MI

Bachelor of Science in Engineering

August 2009 - May 2013

▷ Major: Computer Science, Minor: Mathematics

GPA: 3.98

▷ Selected Courses: Operating Systems, Computer Architecture, Database Systems, Numerical Methods, Linear Algebra, Theoretical Statistics

EXPERIENCE **Lightmatter**, Boston, MA

January 2021 - Present

Researching ways to accelerate deep learning inference on custom photonic hardware, and to make model accuracy robust to noise. Helping guide development of next generation of hardware based on current results.

Tempus Labs, Chicago, IL

June 2019 - January 2021

Created a new domain adaptation method that accounts for confounders, which was deployed on the Tempus RNA-seq pipeline as the source-of-truth for all clinical AI models and pharma data deliveries. Developed a new topic model for gene expression deconvolution in metastatic cancers. Analyzed network learning methods and graph neural nets for gene expression networks and chromosomal rearrangement graphs.

Carnegie Mellon University, Pittsburgh, PA

August 2013 - May 2019

Worked on sparse graphical model learning problems and scalable optimization algorithms for tasks in systems biology. My focus was on using statistical machine learning to discover the gene regulatory networks which explain the effect of genetic variation on clinical traits.

Van Andel Research Institute, Grand Rapids, MI

Summer 2013

Worked under the supervision of Brian Haab to apply feature selection method to pancreatic cancer biomarker discovery and to validate method on proteomics database.

Google, Mountain View, CA

Summer 2012

Worked on server backend for Google Flight Search, developing functionality to improve quality of results for live Flight Search queries.

University of Michigan, Ann Arbor, MI

Winter - Fall 2011

Worked under the supervision of Valeria Bertacco and Debapriya Chatterjee to develop post-silicon validation method. Designed and implemented parallel algorithm in CUDA.

Arbor Networks, Ann Arbor, MI

Summer 2011

Implemented instrumentation in deep packet inspection system and prepared performance analysis tools geared to IPv6 transition.

University of Michigan REU Program

Summer 2010

Analyzed data from simulated advertising auctions under the supervision of Michael Wellman to understand impact of bidding strategies on advertiser profitability.

PEER-REVIEWED PUBLICATIONS	<p>R Hanson, D Martin, <u>C McCarter</u>, J Paulson, “If Loud Aliens Explain Human Earliness, Quiet Aliens Are Also Rare.” <i>The Astrophysical Journal (APJ)</i>, 2021.</p> <p>LE Fernandes, et al., “Real-world Evidence of Diagnostic Testing and Treatment Patterns in US Breast Cancer Patients with Implications for Treatment Biomarkers from RNA-sequencing Data.” <i>Clinical Breast Cancer</i>, 2020.</p> <p><u>C McCarter</u>, J Howrylak, S Kim, “Learning Gene Networks Underlying Clinical Phenotypes Using SNP Perturbations”, <i>PLOS Computational Biology</i>, 2020.</p> <p><u>C McCarter</u> and S Kim, “Large-Scale Optimization Algorithms for Sparse Conditional Gaussian Graphical Models”, <i>International Conference on Artificial Intelligence and Statistics (AISTATS)</i>, 2016.</p> <p><u>C McCarter</u> and S Kim, “On Sparse Gaussian Chain Graph Models”, <i>Advances in Neural Information Processing Systems (NeurIPS)</i>, 2014.</p> <p>S. Moon, <u>C McCarter</u>, YH Kuo, “Active learning with partially featured data”, <i>Proceedings of the 23rd International Conference on World Wide Web</i>, 2014.</p> <p><u>C McCarter</u>, D Kletter, H Tang, K Partyka, Y Ma, S Singh, J Yadav, M Bern, B Haab, “Prediction of Glycan Motifs Using Quantitative Analysis of Multi-lectin Binding”, <i>Proteomics Clinical Applications</i>, vol: 7, issue: 9-10, 2013.</p> <p>D Chatterjee, <u>C McCarter</u>, V Bertacco, “Simulation-based Signal Selection for State Restoration in Silicon Debug”, <i>International Conference on Computer-Aided Design (ICCAD)</i>, 2011</p>	
PATENTS	<p>Systems and methods for multi-label cancer classification. US Patent App. 17/150,992. Publication date: 2021/5/13.</p>	
OPEN-SOURCE CONTRIBUTIONS	<p>onnx2pytorch https://github.com/ToriML/onnx2pytorch Converts ONNX models to PyTorch. [main contributor]</p> <p>PerturbNet https://github.com/SeyoungKimLab/PerturbNet Learns multi-omic gene regulatory networks. [main contributor]</p> <p>MLPerf Inference Benchmark. https://github.com/mlcommons/inference Deep learning benchmark. [memory-efficient pyramidal encoder for RNN-Transducer]</p> <p>matrix-completion. https://github.com/tonyduan/matrix-completion Classical matrix completion. [incremental singular-vector thresholding]</p>	
PRESENTATIONS	<p><i>Transcriptome background tissue correction in metastatic cancers using a correlated composition admixture model.</i></p> <p>American Association for Cancer Research (AACR), Annual Meeting 2020.</p> <p><i>An efficient algorithm for learning a gene network underlying clinical phenotypes under SNP perturbations.</i></p> <p>Genome Informatics meeting at Cold Spring Harbor Labs, November 2017.</p>	
TEACHING	<p><i>Probabilistic Graphical Models</i> (Teaching Assistant) Spring 2016</p> <p><i>Introduction to Machine Learning</i> (Teaching Assistant) Fall 2015</p>	
ACTIVITIES AND PROFESSIONAL SERVICE	<p><i>Paper reviewing</i> June 2016 - Present</p> <p>Reviewer for <i>NeurIPS</i>, <i>IEEE Internet of Things</i>, <i>Statistics and Computing</i>, and <i>SciPy</i>.</p> <p><i>Machine Learning Department Admissions Committee</i> 2015</p> <p>Reviewed application materials of prospective graduate students.</p> <p><i>Machine Learning Department Student Research Symposium</i> 2014</p> <p>Member of organizing committee. Created website and helped plan symposium.</p> <p><i>University of Michigan Robocup (Robot Soccer) Team</i> 2009 - 2012</p> <p>Member and team leader (2010-2011). Developed computer vision subsystem.</p>	

LANGUAGES Python (pandas, numpy, numba, PyTorch), Matlab, C++, C, CUDA, R, Shell, L^AT_EX