Jacob Kelly

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EDUCATION

University of Toronto Toronto, ON Computer Science, Math, Stats · cGPA: 3.93/4 Course Average: 90% Sep 2017 — Jun 2022 Recipient of more than \$11,000 in scholarships and grant funds. Graduate Courses: Machine Learning I & II \cdot Deep Learning \cdot ML for Health \cdot Stochastic Processes Advanced Algorithms & Data Structures · Advanced Differential Equations Undergrad Courses: C & Systems Programming · Enriched Calculus I & II · Intro. Molecular Biology Teaching Assistant: STA414/2104 (Grad. Machine Learning II) · Office hours and assignment grading Experience Machine Learning Research Intern \cdot Python \cdot TensorFlow \cdot pandas \cdot Bash \cdot Git Toronto, ON Sep 2020 — Apr 2021 Deep Genomics • Developed framework for compressing deep convolutional splicing models with neural distillation. Resulting models matched performance across tasks and metrics while 3.7x smaller and 3.3x faster. • Integrated knockdown and peak data with deep learning model for automated oligonucleotide design using a new biological mechanism. Developed statistical benchmarks of performance and significance. Machine Learning Researcher \cdot Python \cdot JAX \cdot PyTorch \cdot Bash \cdot Git \cdot SLURM Toronto, ON Vector Institute for AI · Supervisor: David Duvenaud Sep 2019 — Aug 2020 • Analyzed bias of estimator for scalable entropy-regularized training of Energy-Based Models (EBMs). Cleaned data and tuned EBM performance on semi-supervised classification for 3 tabular datasets. • Derived and implemented numerically-stable Taylor-mode automatic differentiation rules in JAX. Implemented and numerically tested 12 ODE solvers of various orders. Aggregated results for sweeps of >100 jobs on SLURM cluster measuring regularization effect on Neural ODE models across tasks. Computational Biology Researcher · R · MATLAB · Bash · Git Princess Margaret Cancer Research · Supervisor: Benjamin Haibe-Kains Apr 2019 — Sep 2019 • Developed R package for benchmarking machine learning methods for inferring sample-specific gene regulatory networks from single-cell RNA sequencing (scRNA-Seq) data. Computer Vision Software Developer · SPEL+ (internal C++ wrapper) · SVN Markham, ON Epson Research and Development Lab May 2018 — Aug 2018 • Optimized motored stage movements and performed image capture and evaluation asynchronously, supporting researchers by improving speed of data collection by 58%. Android Developer · Java · Android SDK · Estimote API · Git Toronto, ON Jul 2016 — Aug 2016 Cossette Health • Led two team members in reducing noise in bluetooth beacon signal to improve localization and developing pathfinding algorithms for indoor navigation system for The Hospital for Sick Children. Publications 1. W. Grathwohl*, J. Kelly*, M. Hashemi, M. Norouzi, K. Swersky, D. Duvenaud, "No MCMC for me: Amortized sampling for fast and stable training of energy-based models". International Conference on Learning Representations (ICLR) 2021 2. J. Kelly*, J. Bettencourt*, M. J. Johnson, D. Duvenaud, "Learning Differential Equations that are Easy to Solve". Neural Information Processing Systems (NeurIPS) 2020 Projects **JAX** (Open-source contributor) · Python · Git github.com/google/jax • Top 10% of contributors (25 commits, ~ 1000 lines of code). Derived and implemented numerically stable Taylor-mode automatic differentiation rules. Wrote numerical tests and fixes for ODE solvers. AWARDS Undergraduate Student Research Award, NSERC Canada 2020 Dorothy Helen McRobb Scholarship 2019 David L. Squires Foundation Scholarship 2019 Margaret Ronald & Thomas Paxton Taylor Scholarship in Mathematics 2019 Distinction (Top 15%), Euclid National Mathematics Contest, Univ. of Waterloo 2017

Python · Bash · Git · \LaTeX · C/C++ · R · Java Languages: Frameworks: $PyTorch \cdot JAX \cdot TensorFlow \cdot Keras \cdot NumPy \cdot pandas \cdot scikit-learn$ 2017

Top 10, ECOO Central Ontario Programming Contest

SKILLS