# JACOB KELLY

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### **EDUCATION**

# University of Toronto

Toronto, ON

Computer Science, Math, Stats · cGPA:3.93/4 (90%)

Sep 2017 — Jun 2022

Coursework: Machine Learning (Graduate-Level)  $\cdot$  Advanced Algorithms & Data Structures  $\cdot$ 

Advanced Differential Equations · Stochastic Processes · Molecular Biology

### EXPERIENCE

#### Machine Learning Researcher · Python · JAX

Toronto, ON

Vector Institute for AI · Supervisor: David Duvenaud

Sep 2019 — Present

• Improving efficiency of Neural Ordinary Differential Equations (Neural ODEs).

## Computational Biology Researcher $\cdot R \cdot Bash \cdot MATLAB$

Toronto, ON

Princess Margaret Cancer Research  $\cdot$  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.
- Trained elastic net regularized regression with pathway-based feature selection to infer drug response of cell lines from gene expression for patients with acute myeloid leukemia (AML).

Computer Vision Software Developer · SPEL+ (internal C++ wrapper) Epson Research and Development Lab Markham, ON

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%.
- Designed and implemented anchor point based motor-camera calibration method for comparison of 2D object detection and pose estimation algorithm performance on a wide-range of objects.

Android Developer · Java · Android SDK · Estimote API

Toronto, ON

Cossette Health Lab

Jul 2016 — Aug 2016

• Led two team members in reducing noise in bluetooth beacon signal to improve localization and pathfinding algorithms for indoor navigation system for SickKids Hospital.

# PROJECTS

Machine Learning for Health  $\cdot$  PostgreSQL  $\cdot$  Python  $\cdot$  pandas  $\cdot$  scikit-learn  $\cdot$  Keras  $\cdot$  NLTK  $\cdot$  gensim

- Queried Postgres database and summarized data from over 50,000 admissions of nearly 40,000 patients to Beth Israel Deaconess Medical Center.
- Predicted mortality and hypertension from clinical notes and vital signs data using logistic regression and recurrent neural network (RNN) machine learning models.

Genomic Sequencing  $\cdot C++ \cdot Make \cdot Bash$ 

github.com/jacobjinkelly/sequencing

• Implemented Boyer-Moore for genomic sequence alignment with linear time construction of indexes using Z algorithm, achieving 3.5x speedup over naive algorithm.

#### 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, UWaterloo	2017
1st Place, ECOO Central Ontario Programming Contest	2017

#### SKILLS

**Languages & Frameworks:** Python  $\cdot$  C/C++  $\cdot$  R  $\cdot$  Java  $\cdot$  JAX  $\cdot$  PyTorch  $\cdot$  TensorFlow  $\cdot$  NumPy

### Interests