# **Boyang Fu**

• Email: boyang1995@g.ucla.edu • Phone: (+1) 908-294-4423

• Linkedin: https://www.linkedin.com/in/boyang-fu • GitHub: https://github.com/FBoyang

## **Education**

University of California, Los Angeles

Ph.D., Computer Science (Advisor: Prof Sriram Sankararaman)

Los Angeles, CA September 2019 - now

Rutgers, The State University of New Jersey

B.S (Summa cum laude). Computer Science & Mathematics

New Brunswick, NJ January 2016 – May 2019

## **Relevant Courses**

Linear Algebra Statistics Theory Principles Prog. Language Operating System Convex Optimization
Database Management
System Programming
Machine Learning

Graph Theory Algorithms Machine Learning Bioinformatics

## **Skills**

o Programming Languages: Python, R, C++, C, Java, MATLAB, Bash

o Frameworks & Softwares: TensorFlow, Keras, MySQL, Docker

o Operating Systems: Windows, Linux (Ubuntu), Unix.

# Research and Internship

#### Machine Learning and Genomics Lab

Los Angeles, CA

<sup>o</sup> UCLA, Advisor - Prof. Sriram Sanakararaman

Auguest 2019 - Now

- Developing interpretable machine learning algorithms and leverage them to find meaningful biological signals (Emphasis on counterfactual analysis and feature importance explanation).
- Developing highly scalable statistical tools to model genetic architecture of complex traits (Emphasis on hypothesis testing, variance component estimation, kernel learning)

#### **Independent Study & Research Assistant**

New Brunswick, NJ

Rutgers University CS Department, Advisor - Prof. Desheng Zhang

May 2018 - May 2019

Our research focuses on designing a new road construction algorithm that takes important road features (i.e. road type and speed limit) into account to generate a high-quality real-time map

- Constructing the raw roadmap based on sparse GPS points and vehicle trajectory regeneration
- Designing road construction algorithm through roadmap segmentation, then applying kernel density estimation on each segment to filter outliers and then use supervised learning to perform road type classification
- Analyzing the influence of anomalies level and category to the passengers' waiting time based on spatio-temporal information.

Research Assistant, Bruins-In-Genomics (B.I.G.) Summer Program

University of California, Los Angeles , Advisor - Prof. Sriram Sankararaman

June 2018 – August 2018

We developed a comprehensive benchmarking tool during the summer to compare 4 representative genome-wide association study algorithms under multiplexed simulated genetic architectures.

- Designed data simulation algorithm based on multiple underlying genetic structural assumptions and parameters
- Developed software to statistically compare the performance of different algorithms
- Designed theoretical performance threshold to detect the statistical power of each SNP by estimating the non-centralized parameter under the linear model assumption

## **Aresty Research Assistant**

New Brunswick, NJ

Rutgers University Genetics Department, Advisor - Prof. Kevin Chen July 2017 - December 2017
The objective of this research is to apply novel machine learning algorithms to the field of genetic association study

- Processed raw data, including DNA binary transformation, normalization, and sparse matrix data processing
- Researched performance comparison of dimension reduction algorithms (PCA and autoencoder) and performed feature analysis on SNPs dataset
- Applied DNNregressor to predict the numerical phenotype value based on SNPs sparse matrix and studied the best hyperparameters combination with the help of computer cluster

## **Publications**

- o "Marginal Contribution Feature Importance—an Axiomatic Approach for The Natural Case" Authors: A Catav, **B Fu**, J Ernst, S Sankararaman, R Gilad-Bachrach. *arXiv*
- o "A Statistical Model for Quantifying the Needed Duration of Social Distancing for the COVID-19 Pandemic"

Authors: N Rakocz, B Fu, E Halperin, S Sankararaman. medRxiv

- o "PrivateBus: Privacy Identification and Protection in Large-Scale Bus WiFi Systems"

  Authors: Zhihan Fang, **Boyang Fu**, Zhou Qin, Fan Zhang, Desheng Zhang *In ACM UbiComp 2020*
- o "MAC: Measuring the Impacts of Anomalies on Travel Time of Multiple Transportation Systems" Authors: Zhihan Fang, Yu Yang, Shuai Wang, Boyang Fu, Zixing Song, Fan Zhang, Desheng Zhang. In ACM UbiComp 2019

## **Teaching**

Web Applications

Los Angeles, CA

UCLA Computer Science Department

March 2021 - Now

Introduction to Formal Languages and Automata Theory

Los Angeles, CA September 2020 – December 2020

UCLA Computer Science Department

New Brunswick, NJ

**Learning Assistant of Calculus II**Rutgers University Mathematics Department

September 2018 – May 2019

Grader of Calculus III

New Brunswick, NJ

Rutgers University Mathematics Department

September 2017 – December 2017

## **Poster & Presentation**

#### 1. Non-linear set-based association tests for Biobank-scale data

Boyang Fu, Mukund Sudarshan, Ali Pazokitoroudi, Kathryn Burch, Bogdan Pasaniuc, Lakshminarayanan Subramanian, Sriram Sankararaman; Annual meeting of the American Society of Human Genetics, Oct 2020, Virtual

## 2. Pandemic Prediction Modeling Tutorial

Boyang Fu; Focus on variation SIR models and IHME model and the implementation; Tutorial prensentation on Bruins-In-Genomics Hackathon, July 2020, Virtual

#### 3. Non-linear set-based association tests for Biobank-scale data

Boyang Fu, Mukund Sudarshan, Ali Pazokitoroudi, Kathryn Burch, Bogdan Pasaniuc, Lakshminarayanan Subramanian, Sriram Sankararaman; Probabilistic Modeling In Genomics, April 15 2021, Virtual

## **Award**

o Graduate Division Fellowship Award	November 2020
o Most Unique Hack Award in HackHer Competition, Rutgers University	February 2018
o Best Use of Amazon Web Services in HackRU	October 2017
o Academic Excellence Scholarship	2016-2018
o Rutgers University School of Arts and Sciences Dean's List	2016-2018
o Member of Phi Beta Kappa Society	from 2018
o Member of Hall of Fame Data 101	from 2016

# **Mentoring Experience**

o Kevin Delao (Master Student) & Maya Singh (Undergrad student), Bruins in Genomics Summer Program.

Project title: An Analysis on the Performance of SHAP at Predicting Phenotypes in Complex Machine Learning Models