Kristy Mualim

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EDUCATION

McGill University

Montreal, QC

B.Sc in Biochemistry

Sept. 2015 – Dec. 2018

Email: kmualim@stanford.edu

- o BL21 Research Scholar: Research Grant for Independent Student Researchers
- Yale Hackathon Award: Developed a machine learning classifier to rate building accessibility for people with disabilities.
- T.E.A.M Scholar: Grant for Exceptional Students in Chemistry

RESEARCH EXPERIENCE

Carnegie Institute of Science

Palo Alto, CA

Research Assistant, Advisor: Professor Moisés Expósito-Alonso

April. 2021 - Present

• Disentangling genotype-phenotype differences (such as drought tolerance and root growth) in the Brassicae family using deep learning algorithms.

Stanford University

Palo Alto, CA

Research Assistant, Advisor: Professor Jesse Engreitz

April. 2019 - Present

- Curated CRISPR-FlowFISH, GWAS and eQTL datasets to benchmark the performance of existing cCRE-gene linking prediction methods.
- Organized multiple bioinformatics challenge days to foster community collaboration and analysis in the field of Distal Regulation.

Stanford University

Palo Alto, CA

Research Assistant, Advisor: Professor Anshul Kundaje

April. 2019 - Present

- Developed and optimized a data mining and analysis pipeline for processing DNase, ATAC and Histone data on ENCODE to generate an atlas of cCRE-gene linking maps across 200 celltypes and tissues
- Developed deep learning algorithms to investigate the transcription factor landscape and mechanistic behavior of chromatin modifiers

University of California, Berkeley

Berkeley, CA

Research Assistant, Advisor: Professor Montgomery Slatkin

April. 2019 - April 2020

• Developed an analytical solution to estimating coalescence probabilities and population divergence times from SNP data in Neanderthals.

McGill University

Montreal, QC

Undergraduate Researcher, Advisor: Professor Jerome Waldispühl

May. 2018 - Jan. 2019

- Improved and developed upon a citizen-science interface to solve the multiple sequence alignment problem across biological kingdoms - Phylo; a citizen-science platform for solving the multiple sequence alignment problem
- Utilized Phylo to curate a syllabus focused on introducing kids to biological topics.

McGill University

Montreal, QC

Undergraduate Researcher, Advisor: Professor Kalle Gehring

Sept. 2016 – June. 2018

- Developed and improved protein purification experimental protocols.
- Used size-exclusion chromatography and Nuclear Magnetic Resonance (NMR) to discover an unexpected secondary structure within a linker domain of a mitochrondrial protein involved in autophagy (DENND3).

PUBLICATIONS

- 1. **Mualim**, **K.**, Theunert, C. & Slatkin, M. Estimation of coalescence probabilities and population divergence times from SNP data. *Heredity* **127** (2021).
- 2. Nasser, J. et al. Genome-wide enhancer maps link risk variants to disease genes. Nature **593**, 1–6 (2021).
- 3. **Mualim**, **K.** et al. A Computational Validation of Enhancer-Gene Linking. ENCODE Consortium (2019).