

DANIEL LI

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EDUCATION

UNIVERSITY OF CALIFORNIA, BERKELEY

Aug 2014 – May 2017

B.S. Electrical Engineering and Computer Sciences

Berkeley, CA

- **Academics:** 3.5 cumulative GPA
- **Coursework: Computer Science:** Efficient Algorithms and Intractable Problems, Human Computer Interaction, Algorithms in Computational Biology, Structures and Interpretations of Programs (SICP), Data Structures and Algorithms, Computer Architectures | **Mathematics:** Integral, Differential, Vector, Multivariable, Lambda Calculi, (Partial) Differential Equations, Linear Algebra, Discrete Mathematics, Probability Theory and Statistics, (some) Algebra | **Electrical Engineering & Physics:** Mechanics, Electricity, Magnetism, Designing Information Devices and Systems I,II
- **Skills:** Python, Java, C, SQLite, Lisp - Scheme, HTML, LaTeX, R, MIPS, Assembly, Data Analysis, Digital Signal Processing (DSP), Android Studio app development

LA CANADA HIGH SCHOOL

Fall 2011 – Spring 2014

- **Academics:** 4.7/4.0 weighted GPA, 4.0/4.0 GPA, 2310 SAT
- Graduated in 3 years, ranked 2nd out of 400

RESEARCH EXPERIENCE

Pachter Group – *Principal Investigator: Lior Pachter*

Fall 2015 – Present

Computational Biology (Mathematics and Genomics)

Berkeley, CA

- Research in novel approaches to RNA-sequencing with the features in abundance estimation, transcript annotation difficulties, differential expression
- Current investigation on the improvement of bulk cell and single cell RNA-seq analysis

Rao Group – *Principal Investigator: Satish Rao*

Spring 2016 – Present

Algorithms, Theory, Optimization on Computational Biology

Berkeley, CA

- Investigating phylogenetic algorithms
- Optimization of paths for algorithms (? TBD)

INDUSTRY EXPERIENCE

Factual

Summer 2016

Research and Development (R&D) Intern

Los Angeles, CA

- Operating the data extraction pipeline through quality evaluation, localespecific extraction, canonicalization, and decanonicalization
- Developing, maintaining, and documenting new processes for data extraction and presentation

PROJECTS

KALLISTO – C++ / R (Pachter Group)

- Novel approach decreasing analysis time of 30 million human reads in less than 3 minutes
- Optimization of statistical likelihood model through non uniform distribution analysis for more accurate projections onto correct subspaces

SLEUTH – R (Pachter Group)

- Implement statistical algorithms for differential analysis for pseudo-alignment of RNA transcripts with interactive plots for real-time exploratory analysis
- Visualization of bias weights of RNA through integration of bias weights and hexamer indices

AWARDS AND HONORS

MIT THINK

Spring 2014

Designed a Novel Approach to Mitigate Earthquakes

Cambridge, MA

- Awarded \$2,000