

# Kian Faizi

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## EDUCATION

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### California Institute of Technology

*Ph.D. in Systems Biology*

Sep. 2021 – present

*Pasadena, CA*

### University of California, San Diego

*B.S. in Molecular Biology, Minor in Mathematics (GPA: 3.75)*

Aug. 2017 – June 2021

*La Jolla, CA*

- Selected coursework: Bioinformatics, Biophysics, Computational Linear Algebra, Dynamical Systems, Gene Regulation, Gene Editing, Statistics, Stochastic Processes

### International School of Kuala Lumpur

*I.B. Diploma, Earth Club President, Varsity Basketball Team Captain*

Aug. 2013 – June 2017

*Kuala Lumpur, Malaysia*

## EXPERIENCE

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### Lab Technician

*Lab of Wolfgang Busch, Salk Institute for Biological Studies*

Nov. 2019 – Sep. 2021

*La Jolla, CA*

- Investigated cost-performance trade-offs in the *Arabidopsis* root system using high-throughput phenotyping and graph-theoretic modeling
- Created a Python GUI for segmenting time-series images of root growth
- Helped develop an algorithm for plant phenotyping from noisy LiDAR scans
- Built a pipeline for co-expression network analysis of scRNA-seq data to identify genetic targets for future crop engineering

### Volunteer Research Assistant

*Lab of Patrick Hsu, Salk Institute for Biological Studies*

Nov. 2018 – Nov. 2019

*La Jolla, CA*

- Worked on characterizing Cas13d, a novel RNA-targeting CRISPR effector
- Created an automated pipeline to mine metagenomes for new Cas13d orthologs, and searched over 20TB of sequence data
- Assisted in conducting a pooled 150,000-guide CRISPR-Cas13d screen in K562s to optimize gRNA design

## PUBLICATIONS

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### Branch-Pipe: Improving graph skeletonization around branch points in 3D point clouds.

- Illia Ziamtsov, **Kian Faizi**, and Saket Navlakha. *To appear in Remote Sensing*.

### Network design principles in the *Arabidopsis* root system.

- **Kian Faizi**, Matthieu Platre, Arjun Chandrasekhar, Saket Navlakha, and Wolfgang Busch. *In prep.*

### A pooled CRISPR-Cas13d screen reveals guide RNA efficiency rules.

- Silvana Konermann, **Kian Faizi**, Peter Lotfy, and Patrick Hsu. *In prep.*

## TEACHING

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### Undergraduate Instructional Apprentice | UCSD

Aug. 2020 – Dec. 2020

- For *Genetic Inquiry*, supervised by Stanley Lo

## POSTERS AND PRESENTATIONS

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### Co-expression analysis of single-cell RNA-seq data | Talk

Oct. 2020

- HDSI Research Conference

### Mining Genomes for RNA-Targeting CRISPR Effectors | Talk

Aug. 2019

- UCSD Summer Research Conference

### Metagenomic Discovery of Type VI-D CRISPR Effectors | Poster

June 2019

- UCSD Biology Student Research Showcase

## HONORS AND AWARDS

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<b>Halicioglu Data Science Institute Scholarship Project Award</b>   <i>UCSD</i>	May 2021
<b>DOE CSGF Honorable Mention</b>   <i>Krell Institute</i>	Apr. 2021
<b>Halicioglu Data Science Institute Scholarship</b>   \$2,500   <i>UCSD</i> <ul style="list-style-type: none"><li>Project: <i>Single-cell transcriptomics and web mining for rapid reverse genetics in plants</i>, proposed under Wolfgang Busch</li></ul>	Dec. 2019
<b>Eureka! Research Scholarship for Biological Sciences</b>   \$5,000   <i>UCSD</i> <ul style="list-style-type: none"><li>Project: <i>Discovery and development of Type VI-D CRISPR effectors for transcriptome engineering applications</i>, proposed under Patrick Hsu</li></ul>	June 2019
<b>Provost Honors</b>   <i>UCSD</i>	quarterly

## PROFESSIONAL ACTIVITIES

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<b>Undergraduate Bioinformatics Club</b> Member   <i>UCSD</i>	Nov. 2017 – June 2021
<ul style="list-style-type: none"><li>Collaborated with Illumina to develop digital resources for high school students interested in bioinformatics</li><li>Helped organize the 2018 Faculty &amp; Industry Bioinformatics Symposium</li><li>Volunteered at the SD Science &amp; Engineering Festival to teach the community about DNA sequencing technology</li></ul>	

## SKILLS

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**Laboratory:** Cell/tissue culture, molecular cloning, CRISPR screens, optical microscopy  
**Computational:** Image analysis, biologically-inspired algorithms, point clouds, GUI development  
**Languages:** Python, bash, HTML/CSS/JS  
**Organizational:** Git, L<sup>A</sup>T<sub>E</sub>X, Linux/Unix systems  
**Libraries:** NetworkX, matplotlib, seaborn, numpy, pandas, BLAST+