

# Rosalind Pan

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## EDUCATION

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- California Institute of Technology** Expected, 2028  
Pasadena, CA  
Ph.D in Biochemistry and Molecular Biophysics  
Cumulative GPA: 3.90/4.00
- The University of Chicago** June, 2022  
Chicago, IL  
B.S. *summa cum laude* in Biology & Computational and Applied Mathematics  
Cumulative GPA: 3.97/4.00  
Thesis: “Evolutionary analysis of unalignable protein sequences”

## EXPERIENCE

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- California Institute of Technology** June 2022 - Present  
Pasadena, CA  
*Graduate Researcher, Rob Phillips Group*
- Build thermodynamic models to model transcriptional regulation in microbial organisms
  - Map transcription factors to corresponding regulatory regions using gene knockout and RNA sequencing
- The University of Chicago** October 2019 – June 2022  
Chicago, IL  
*Undergraduate Researcher, D. Allan Drummond Laboratory*
- Research sequence-function relationships of low-complexity regions through alignment-free evolutionary analysis
  - Develop computational methods to quantify sequence conservation and identify signatures of selection
  - Perform biochemical and biophysical assays to investigate the effects of perturbing evolutionarily conserved sequence features in low-complexity regions
  - Carried out proteome-wide conservation analysis of highly charged protein regions
- The Marine Biological Laboratory** June – August 2019  
Woods Hole, MA  
*Undergraduate Researcher, Joshua Rosenthal Laboratory*
- Researched a new system of site-directed RNA editing that uses the bacterial enzyme TadA
  - Designed and purified recombinant TadA-based enzymes and novel RNA substrates
  - Carried out in cellula and in vitro editing assays using the TadA editing system
- The University of Chicago** February – June 2019  
Chicago, IL  
*Undergraduate Researcher, Margaret Gardel Laboratory*
- Studied the dynamics and regulation of three-dimensional cell shape in epithelial tissues
  - Imaged live cells on collagen gel using confocal fluorescence microscopy
  - Performed computational analysis of time-lapse microscopy images

## PUBLICATIONS

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- [3] **Rosalind W. Pan**, Tom Röschinger, Kian Faizi, and Rob Phillips. 2024. **Dissecting Endogeneous Genetic Circuits from First Principles**. bioRxiv. <https://doi.org/10.1101/2024.01.28.577658>.

[2] Shichen Liu, **Rosalind W. Pan**, Heun Jin Lee, Shahriar Shadkhoo, Fan Yang, Chunhe Li, Zijie Qu, and Matt Thomson. 2024. **Force Propagation in Active Cytoskeletal Networks**. arXiv [cond-Mat.soft]. <http://arxiv.org/abs/2401.04217>.

[1] Cat Triandafillou\*, **Rosalind W. Pan\***, Aaron Dinner, D. Allan Drummond (\*contributed equally). 2023. **Pervasive, conserved secondary structure in highly charged protein regions**. PLOS Computational Biology 19(10): e1011565. <https://doi.org/10.1371/journal.pcbi.1011565>.

## PRESENTATIONS

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[9] “Dissecting endogeneous genetic circuits from first principles.” 11th Annual Winter q-bio meeting, Feb 19-23, 2024. O’ahu, HI. **Oral and poster presentation**.

[8] “Dissecting endogeneous genetic circuits from first principles.” Mechanisms of Microbial Transcription Gordon Research Conference, June 11-16, 2023. Hooksett, NH. **Poster presentation**.

[7] “Dissecting endogeneous genetic circuits from first principles.” Mechanisms of Microbial Transcription Gordon Research Seminar, June 10-11, 2023. Hooksett, NH. **Oral and poster presentation**.

[6] “Alignment-free evolutionary analysis of unalignable protein sequences.” Chicago Area Undergraduate Research Symposium, April 10, 2021. Remote. **Poster presentation; won Top Poster in Biological Sciences**.

[5] “Alignment-free evolutionary analysis of unalignable protein sequences.” UChicago Computational and Applied Mathematics Student Seminar, February 23, 2021. Remote. **Oral presentation**.

[4] “Alignment-free evolutionary analysis of unalignable protein sequences.” 26th Annual Midwest Stress Response and Molecular Chaperone Virtual Meeting, January 16, 2021. Remote. **Poster presentation**.

[3] “Quantifying natural selection in the low-complexity regions of poly(A)-binding proteins.” Midstates Consortium for Math and Science Undergraduate Research Symposium in the Biological Sciences and Psychology, October 31, 2020. Remote. **Oral presentation**.

[2] “Site-Directed RNA Editing Using TadAs.” UChicago Careers in STEM 6th Annual Undergraduate Research Symposium, October 11, 2019. Chicago, IL. **Poster presentation**.

[1] “Site-Directed RNA Editing Using TadAs.” Marine Biological Laboratory Undergraduate Poster Session, August 15, 2019. Woods Hole, MA. **Poster presentation**.

## TEACHING

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### **Bi 1: The Great Ideas of Biology**

*Head Teaching Assistant*

April – June 2023

*Pasadena, CA*

- Led computational tutorials, weekly recitations, and office hours
- Prepared and graded homework assignments

### **BIOS 26120: An Introduction to Bioinformatics and Proteomics**

*Teaching Assistant*

September – December 2021

*Chicago, IL*

- Led weekly computer lab sessions and graded coding assignments

## HONORS AND AWARDS

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UChicago Quantitative Biology Summer Fellowship	2021
Enrico Fermi Scholar, UChicago Physical Sciences Collegiate Divisional Honors	2021
Janet Rowley Scholar, UChicago Biological Sciences Collegiate Divisional Honors	2021
Elected to UChicago Beta of Illinois Chapter of Phi Beta Kappa	2021
Liew Family College Research Fellows Fund	2021
UChicago Robert Maynard Hutchins Scholar	2020
UChicago Biological Sciences Collegiate Division Summer Research Fellowship	2020
Jeff Metcalf Summer for Undergraduate Research Fellowship	2019
Chicago Materials Research Center Research Exploration Fellowship	2019

## SKILLS

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<b>Software</b>	Python, R, bash, Adobe Illustrator, $\LaTeX$
<b>Wet Lab</b>	Molecular cloning, RNA-Seq, cell culture, optical microscopy