

# Samuel E. Holtzen, Ph.D.

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## Education and Training

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<b>Postdoctoral Fellow, Biology and Biological Engineering</b> <i>California Institute of Technology</i>	2019-2024
<b>Doctor of Philosophy, Molecular Cell and Developmental Biology</b> <i>University of Colorado Boulder</i>	2019-2024
<b>Higher Education Research Experiences (HERE) Intern</b> <i>Oak Ridge National Laboratory</i>	2018-2019
<b>Bachelor of Science in Biochemistry, Pre-Health Concentration</b> <i>Georgia Institute of Technology, Cum Laude</i>	2013-2017

## Teaching and Mentorship Experience

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<b>Analytical and Quantitative Light Microscopy, Teaching Assistant</b> <i>Principal Instructor: Dr. Amy Palmer</i>	Spring 2023 and 2024
<b>Phage Discovery Laboratory, Teaching Assistant</b> <i>Principal Instructor: Dr. Nancy Guild and Megan Greening</i>	Spring 2020
<b>Principles of Genetics, Teaching Assistant</b> <i>Principal Instructor: Dr. Jennifer Knight</i>	Spring 2020
<b>Developmental Biology, Teaching Assistant</b> <i>Principal Instructor: Dr. Jennifer Knight</i>	Fall 2019

## Research Experience

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<b>Repetitive element-based spatial barcoding for rapid high-throughput screening</b> <i>Advisor: David Van Valen, Assistant Professor, California Institute of Technology</i>	2024-Present
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Cellular barcoding can allow researchers to identify single-cell perturbations in mixed populations. Spatial optical barcoding can accelerate these so-called pooled optical screens by accelerating the speed at which these barcodes can be read out. I am developing a method of spatial barcoding that uses dCas9 and gRNAs to target repetitive elements (REs) in the genome. The patterns of these gRNA-RE complexes can be read out with multicolor FISH. A deep learning model trained on these patterns can determine the identity of these gRNAs with 90-99% accuracy. Since the encoding space is only limited by the number of discernible patterns and FISH colors, this technique has a very large encoding space and requires only one multicolor FISH step for readout.

<b>Computational design and <i>in vitro</i> validation of novel metabolite biosensors</b> <i>Advisor: David Van Valen, Assistant Professor, California Institute of Technology</i>	2024-Present
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There is a distinct lack of well-characterized binding proteins for use in fluorescent biosensors designs that are specific to less common metabolites. The advent of computational protein design tools such as RoseTTAFold Diffusion and LigandMPNN, and heuristic-driven protein folding tools like AlphaFold has enabled our ability to build bespoke proteins on demand. I am currently designing and testing novel sensing domains for use in biosensors for less-studied

metabolites involved in immune metabolism. Through this, I hope to illuminate understudied aspects of metabolism in an array of cells as they grow, divide, and differentiate.

### **Mapping the zinc requirements of the mammalian cell cycle**

2020-2024

*Advisor: Amy Palmer, Professor of Biochemistry, University of Colorado Boulder*

Mild to moderate zinc deficiency causes a bifurcation in cell fate and increased DNA damage response in a cell culture model. This phenotype has been demonstrated by several studies, but to date, no mechanism has been defined for these hallmark phenotypes. My thesis research aims to decouple the causes, consequences, and sequence of events that cause zinc deficient cells to enter a quiescent state.

### **Cellular dynamics of free zinc throughout the mammalian cell cycle**

2020-2023

*Advisor: Amy Palmer, Professor of Biochemistry, University of Colorado Boulder*

The Palmer group has engineered a live-cell genetically encoded zinc sensor with high dynamic range within the physiological zinc range in mammalian cells. We have developed a method of monitoring cytosolic zinc using long-term live-cell imaging of single cells. Using this system, we have identified zinc dynamics in the mammalian cell cycle. Additionally, we monitor these dynamics after depletion of a transcription factor involved in the response to high zinc. I was responsible for data stewardship and analysis of terabytes of high-content live-cell imaging experiments.

### **Engineering *Pseudomonas putida* KT2440 for the valorization of lignin**

2018-2019

*Advisor: Adam Guss, Biosciences Division, Oak Ridge National Lab*

I worked with a graduate student to engineer the metabolism of *Pseudomonas putida* to catabolize lignin monomers into value-added chemicals. Specifically, I developed and optimized extraction techniques of fatty alcohols and polyhydroxyalkanoate monomers for analysis on our group's GC-MS. In addition, I was the caretaker of the GC-MS and performed routine maintenance and troubleshooting.

### **Biomarker analysis for validating Earth analogue sampling techniques**

2016-2018

*Advisor: Amanda Stockton, Assistant Professor of Chemistry & Biochemistry, Georgia Institute of Technology*

As an undergraduate research assistant, my task was the cataloging and curation of precious Icelandic tephra samples from field expeditions. In addition, I developed a pipeline for extraction of DNA from low-biomass samples and library preparation for 16S rRNA community analysis. Additionally, I aided in the development of an *in situ* co-culturing chip for unculturable microbes. I was afforded the opportunity to embark on a field expedition in June and July 2018 to the Icelandic highlands to sample volcanic tephra from the active Holuhraun lava field.

## **Skills and Expertise**

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**Laboratory Skills:** Mammalian cell culture, high-content live-cell imaging, molecular cloning, Western blot analysis, next-generation sequencing library preparation and analysis, fluorescence microscopy, single- and multi-color flow cytometry, ATAC-seq, RNA-seq, quantitative PCR, high-content microscopy, microscope maintenance

**Data Skills:** image analysis, data pipeline development, machine learning, statistics, high performance computing

**Programming Languages:** Python, MATLAB, Bash, R, SQL

**Languages:** English (native speaker), French (limited working proficiency)

## Publications and Other Works

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### In Press:

Ocampo, D., Damon, L., Sanford, L., **Holtzen, S.**, Jones, T., Allen, M., Dowell, R., Palmer, A. (2024). Cellular zinc status alters chromatin accessibility and binding of TP53 to DNA. *Life Science Alliance*, e202402638

**Holtzen, S.**, Rakshit, A., Palmer, A. (2024). Measuring cell cycle  $Zn^{2+}$  dynamics using a FRET-based biosensor. *STAR Protocols*.

**Holtzen, S.**, Rakshit, A., Palmer, A. (2024). Measuring labile cytosolic  $Zn^{2+}$  using an in-situ calibration of a genetically encoded FRET sensor. *STAR Protocols*.

**Holtzen, S.**, Navid, E., Kainov, J., Palmer, A. (2024). Transient  $Zn^{2+}$  deficiency induces replication stress and compromises daughter cell proliferation. *Proceedings of the National Academy of Sciences*, 121(19), e2321216121.

Rakshit, A.\*, **Holtzen, S. E.\***, Lo, M. N., Conway, K. A., & Palmer, A. E. (2023). Human cells experience a  $Zn^{2+}$  pulse in early G1. *Cell Reports*, 42(6).

Tan, G. K., Simpson, A., **Holtzen, S.**, Amador, E., Cable, M. L., Cantrell, T., ... & Stockton, A. M. (2022). Spatial Variation in Results of Biosignature Analyses of Apparently Homogeneous Samples from Mars Analogue Environments in Iceland. *ACS Earth and Space Chemistry*, 6(6), 1472-1481.

Rader, E., Simpson, A., Amador, E., Fraser, J. M., **Holtzen, S.**, Hanna, A., ... & Stockton, A. (2020). Preferably Plinian and Pumaceous: implications of microbial activity in modern volcanic deposits at Askja Volcano, Iceland, and relevancy for Mars exploration. *ACS Earth and Space Chemistry*, 4(9), 1500-1514.

\*These authors contributed equally

### Conference Abstracts:

*Untangling the Role of Zinc in the Mammalian Cell Cycle and DNA Damage Response*, **Samuel Holtzen**, Joseph Kainov, Elnaz Navid, Lottie Steward, Amy Palmer, Chemical Biology and Physiology, 2022, Conference Abstract

*Production of medium chain length alcohols from lignin-derived aromatic compounds in Pseudomonas putida*, Jay Huenemann, Joshua Elmore, **Samuel Holtzen**, Adam Guss, SIMB Annual Meeting and Exhibition, 2019, Conference Abstract

*Biomarker and Geochemical Assay Validation in Mars Analog Sites: Lessons from the FELDSPAR (Field Exploration and Life Detection Sampling for Planetary Analog Research)*, Diana Gentry, Elena Sophia Amador, Morgan L Cable, Thomas Cantrell, Nosheen Chaudry, Thomas Cullen, Zachary A Duca, **Samuel Holtzen**, Malene B Jacobsen, David King, Jessica Kirby, Heather Catherine McCaig, Gayathri Murukesan, Erika Rader, Adrienne Reeder, Vincent Rennie, Edward Schwieterman, Alexander Michael Sessa, Adam Stevens, Scot M Sutton, George Kenneth Tan, Chang Yin, David Cullen, Wolf Geppert, Amanda M Stockton, Project, 2019, Conference Abstract

*Evaluation of spatial variation in life-detection assay results in apparently homogeneous samples from Mars analog environments in Iceland*, George Kenneth Tan, Amanda M Stockton, Morgan L Cable, Wolf Geppert, David Cullen, Diana Gentry, Elena Sophia Amador, Adam Stevens, Thomas Cantrell, Zachary A Duca, **Samuel Holtzen**, Heather Catherine McCaig, Jarah Whitehead, Gayathri Murukesan, Vincent Rennie, Jessica Kirby, Thomas Cullen, Astrobiology Science Conference, 2019, Conference Abstract

*A Recent Volcanic Eruption, Holuhraun, in the Central Highlands of Iceland as a Mars analog: The 2018 Field Campaign of FELDSPAR*, Amanda M Stockton, Elena Amador, Morgan L Cable, David Cullen, Julia Fraser, Diana Gentry, Wolf Geppert, **Samuel Holtzen**, Gayathri Murukesan, Erica Raider, Adrienne Reeder, Alexander Michael Sessa, Astrobiology Science Conference, 2019, Conference. Abstract