

# Elliot MacKrell

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## Education

<b>California Institute of Technology</b> Ph.D. Chemical Engineering Thesis Committee: David A. Tirrell, Dianne K. Newman, Mikhail Shapiro, Rustem Ismagilov (Chair)	Pasadena, CA 2016-Present
<b>University of Florida</b> B.S. Chemical Engineering Minor: Biomolecular Engineering	Gainesville, FL 2016

## Research Experience

<b>California Institute of Technology</b> , Division of Chemistry and Chemical Engineering <i>NSF Graduate Fellow; Advisor: Prof. David A. Tirrell</i>	2016-Present
<ul style="list-style-type: none"><li>- Engineered the degradation of a mutant methionyl-tRNA synthetase to establish a method for time-resolved proteomic analysis of transient physiological states in <i>E. coli</i>.</li><li>- Identified 152 novel endogenous protease substrates in growing and growth-arrested <i>E. coli</i> cells by establishing a quantitative proteomic screen.</li><li>- Discovered global regulatory roles of proteolysis by combining linear modeling, independent component analysis, customized bioinformatic searches against the EcoCyc database, and ontological and regulatory network enrichment.</li></ul>	
<i>Graduate Rotation Student; Advisor: Prof. Frances H. Arnold</i>	
<ul style="list-style-type: none"><li>- Expanded the substrate scope of the <i>P. furiosus</i> tryptophan synthase <math>\beta</math>-subunit by assisting in the construction and screening of a directed evolution-based library.</li></ul>	
<b>EMD Serono</b> , Department of Translational Medicine and Global Clinical Development <i>Bioinformatics Co-op; Supervisor: Dr. Irina Kalatskaya</i>	2022
<ul style="list-style-type: none"><li>- Developed Shiny application for visualizing cell type-specific gene expression, identifying differentially expressed genes across cell types and disease states, and conducting gene set enrichment in real time.</li><li>- Established the semantic association of text-mined ontology terms in PubMed publications through information-theoretic and statistical approaches.</li><li>- Developed Shiny application for the real-time analysis of text mining data from PubMed to identify genes, pathways, and diseases associated with EMD Serono queries.</li></ul>	
<b>Harvard Medical School</b> , Wellman Center for Photomedicine, Massachusetts General Hospital <i>NSF REU Fellow; Advisor: Prof. Robert Redmond</i>	2015
<ul style="list-style-type: none"><li>- Investigated the effects of photocrosslinking on ex vivo tissue biomechanics via cyclic uniaxial tensile testing.</li></ul>	
<b>Johns Hopkins University School of Medicine</b> , Department of Radiation Oncology <i>NSF REU Fellow; Advisor: Prof. Robert Ivkov</i>	2014
<ul style="list-style-type: none"><li>- Evaluated the anti-tumor immune response generated by magnetic nanoparticle hyperthermia and ionizing radiation treatments in a mouse model for cancer.</li></ul>	
<b>University of Florida</b> , Department of Chemistry <i>Undergraduate Research Assistant; Advisor: Prof. Brent S. Sumerlin</i>	2012-2015
<ul style="list-style-type: none"><li>- Developed an amphiphilic and pH-responsive copolymer for phloem-specific drug delivery in plants by functionalizing polysuccinimide (PSI) with various hydrophilic molecules.</li></ul>	

## Publications

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Jones, J.J., **MacKrell, E.J.**, Wang, T.Y., Lomenick, B., Roukes, L.M., and Chou, T.F. “tidyproteomics: An open-source R package and data object for quantitative proteomics post analysis and visualization.” **BMC Bioinformatics**, **2023**, *24*, 239. <https://doi.org/10.1186/s12859-023-05360-7>

Hill, M.R., **MacKrell, E.J.**, Forsthoefel, C.P., Jensen, S., Chen, M., Moore, G., He, Z.L. and Sumerlin, B.S. “Biodegradable and pH-Responsive Nanoparticles Designed for Site-Specific Delivery in Agriculture.” **Biomacromolecules**, **2015**, *16*, 1276-1282. <https://doi.org/10.1021/acs.biomac.5b00069>

## Awards and Honors

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<b>Center for Environmental Microbial Interactions (CEMI) Pilot Grant</b> , California Institute of Technology	2018
<b>CEMI Fellow</b> , California Institute of Technology	2016-
<b>NSF Graduate Research Fellowship</b> , California Institute of Technology	2016-2021
<b>Yao Su Student Research Prize</b> , HST Summer Institute	2015
<b>NSF REU Fellowship</b> , Harvard-MIT Division of Health Sciences and Technology (HST)	2015
<b>Collins Engineering Scholarship</b> , University of Florida	2015
<b>NSF REU Fellowship</b> , Johns Hopkins University Institute for NanoBioTechnology (INBT)	2014
<b>University Scholars Program</b> , University of Florida	2013

## Presentations

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**MacKrell, E.J.**, Sweredoski, M.J., Moradian, A., Garbos, S.D., Tirrell, D.A. “Engineered Protein Degradation for Time-Resolved Proteomics in Bacteria.” Poster presentation delivered at ASM Microbe, San Francisco, CA, June 2019.

Liu, X., **MacKrell, E.J.**, Tirrell, D.A. “Selective Protein Labeling of Nutrient-Limited Cells in Heterogeneous Biofilms with Bio-Orthogonal Non-Canonical Amino Acid Tagging (BONCAT)” Poster presentation delivered at the Institute for Collaborative Biotechnologies Summer Meeting, Santa Barbara, CA, July 2017.

**MacKrell, E.J.**, McCormack, M., Goldstone, R. and Redmond, R. “Effects of Photocrosslinking on Tissue Biomechanics.” Poster presentation delivered at the Harvard-MGH Summer Institute Presentations, Cambridge, MA, August 2015.

Hill, M.R., **MacKrell, E.J.**, Forsthoefel, C.P., Jensen, S., Moore, G., He, Z.L. and Sumerlin, B.S. “Biodegradable and pH-Responsive Nanoparticles Designed for Site-Specific Delivery in Agriculture.” Poster presentation delivered at the AIChE Annual Student Conference, Atlanta, GA, November 2014.

**MacKrell, E.J.**, Attaluri, A.C., Zhou, H., Hedayati, M. and Ivkov, R. “Immunomodulation via Nanoparticle-Mediated Hyperthermia and Ionizing Radiation to Enhance Resistance against Distal Tumors.” Oral presentation delivered at the Johns Hopkins University Nano-Bio Symposium, Baltimore, MD, August 2014.

Hill, M.R., **MacKrell, E.J.** and Sumerlin, B.S. “pH-Responsive and Biodegradable Nanoparticles Capable of Loading and Releasing Small-Molecule Cargo.” Poster presentation delivered at the Florida Undergraduate Research Conference, Miami, FL, February 2014.

## Skills and Coursework

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**Computational skills:** Bioinformatic and statistical analysis of mass spectrometry-based proteomics data, Text mining, Machine learning, Shiny application development, Linear modeling, Regulatory network analysis, Gene set enrichment, Modeling and parameter estimation through Bayesian inference and maximum likelihood estimation, Stochastic simulation of biochemical systems

**Software environments:** Python (Pandas, NumPy, Scikit-learn, Matplotlib, Bokeh, Altair, Seaborn, Jupyter Notebook), R (Shiny, Bioconductor, Clarivate CBDD, RStudio), ImageJ, Mathematica, GeneData Profiler, Visual Studio Code, Git, LaTeX,

Proteomics analysis suites (MaxQuant, ProteomeDiscoverer, Cytoscape, STRING, Perseus with SILAC and TMT quantitation and normalization)

**Relevant elective coursework:** CS155: Machine Learning & Data Mining, CS156a: Learning Systems, Bi/BE103: Data Analysis in the Biological Sciences, Bi/BE/CS 183: Introduction to Computational Biology and Bioinformatics, ChE/BE 163: Biomolecular Engineering, BE 150: Design Principles of Genetic Circuits, APh 161: Physical Biology of the Cell

## Teaching Experience

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### California Institute of Technology

- **Undergraduate Senior Thesis Mentor**, Cindy Cao 2019-2021
- **Graduate Teaching Assistant**, ChE 141: Data Science for Chemical Systems 2020
- **Graduate Teaching Assistant**, ChE 103c: Transport Phenomena (Mass Transport) 2018-2019
- **Certificate of Practice in University Teaching**, Caltech Center for Learning and Teaching Outreach 2018-2020
- **Undergraduate Research Mentor**, Samuel Owens 2017

### University of Florida

- **Society for Biomaterials Education Challenge**, Society for Biomaterials, University of Florida Chapter 2015
- **Tutoring for Engineering Excellence (TEE) Chair**, Tau Beta Pi, Florida Alpha Chapter 2014-2015