

# ELIJAH KARVELIS

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Thorough and thoughtful computational biophysicist with a passion for combining physics-based and data-driven approaches to address critical problems in biomolecular engineering; currently seeking a role in this area

## EDUCATION

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<b>Massachusetts Institute of Technology, Cambridge</b> Doctorate, Biological Engineering (Advisor: Bruce Tidor)	<i>2018 - Present</i> GPA: 5.0/5.0
<b>University of Illinois, Urbana-Champaign</b> Bachelor of Science, Chemical Engineering	<i>2014 - 2018</i> GPA: 4.0/4.0

## TECHNICAL SKILLS AND EXPERIENCE

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<b>Methods</b>	Molecular dynamics, machine learning, data science, protein design
<b>Programming</b>	Python, Unix/Linux and familiarity with Bash and MATLAB
<b>Software &amp; packages</b>	CHARMM, Gaussian, PyTorch, TensorFlow, Scikit-learn, Pandas, OSPREY

## RELEVANT RESEARCH EXPERIENCE

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<b>Computational Modeling and Engineering of Enzyme Catalysis</b> <i>Graduate Research Assistant</i>	MIT 2019 - present
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- Implemented a platform to apply a rare event sampling technique to flexibly collect and process hundreds of thousands of hybrid QM/MM molecular dynamics simulations of attempted enzyme-catalyzed reactions. Used this platform to (i) compute energies of activation, (ii) collect simulations of thousands of turnover events, and (iii) calculate rate constants using kinetic data for 55 different enzyme variants.
- Trained machine learning models to characterize key structural drivers of catalysis in a model enzyme.
- Developed an approach, combining machine learning models and simulations of turnover dynamics, for rational enzyme redesign toward increased activity with an over 50% success rate (based on calculated results) for a natural enzyme-substrate system.
- Identified three unique mechanisms by which designed mutants achieved increased specific activity.
- Developed a pipeline for interfacing with the dynamics data collected across 55 different enzyme variants.
- Evaluated different deep learning models for mapping from turnover dynamics to overall activity, demonstrating that activity can be predicted from pre-reaction structural dynamics.
- Utilized and helped maintain the Tidor lab's private cluster, managing basic upkeep and troubleshooting tasks in a high-performance Unix-based computing environment.

<b>In vitro Modeling of Glioblastoma Invasion</b> <i>Undergraduate Research Assistant</i>	UIUC 2016 - 2018
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- Developed a microfluidic platform for modeling glioblastoma invasion in the context of a vascularized brain-mimetic hydrogel with tunable biochemical and biophysical properties.

## Teaching and Training

*Teaching Assistant and Graduate Research Assistant*

- TA'ed Principles of Molecular Bioengineering (20.420) at MIT in Fall 2019, a graduate level course covering methods in biomolecular analysis and engineering. TA'ed for four different engineering courses at UIUC.
- Mentored and trained two undergraduate researchers and four rotating graduate students.
- Kaufman Teaching Certificate from MIT (in progress).

## SELECTED AWARDS

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<b>NSF GRFP Fellow:</b> doctoral studies funding	2018 - 2021
<b>MIT Viterbi Fellow:</b> doctoral studies funding	2018
<b>Stamps Leadership Scholar:</b> UIUC's most generous and selective award	2014 - 2018
<b>Amgen Scholar - UC Berkeley:</b> research experience scholarship	2017
<b>Goldwater Scholar:</b> undergraduate award in STEM	2016
<b>Phi Beta Kappa:</b> honor society	2016
<b>Bronze Tablet Recipient:</b> designates top 3% of UIUC graduates	2018
<b>Chancellor's Scholar:</b> UIUC's highest honors program (125/7000 students)	2014 - 2018
<b>James Scholar:</b> UIUC honors program (top 20% of students)	2014 - 2018

## PEER-REVIEWED PUBLICATIONS

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Karvelis, E., Swanson, C., Tidor, B. Substrate turnover dynamics guide ketol-acid reductoisomerase redesign for increased specific activity. *In preparation.*

Ngo, M. T., Karvelis, E., Harley, B. A. C. Multidimensional hydrogel models reveal endothelial network angiocrine signals increase glioblastoma cell number, invasion, and temozolamide resistance. *Integrative Biology*, 12(6):139-149. 2020. <https://doi.org/10.1093/intbio/zyaa010>

## SELECTED PRESENTATIONS

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37th Annual Symposium of the Protein Society; Boston, MA: Karvelis, E., Tidor, B. *Relating Turnover Dynamics to Catalytic Efficiency Identifies Enzyme Mutants with Increased Activity.* Poster. July 14, 2023.

ACS Spring 2023; Indianapolis, IN: Karvelis, E., Tidor, B. *Turnover dynamics and machine learning guide redesigning KARI for increased catalytic activity.* Oral. March 29, 2023.

<https://acs.digitellinc.com/sessions/535607/view>

67th Biophysical Society Annual Meeting; San Diego, CA: Karvelis, E., Tidor, B. *Using turnover dynamics to redesign KARI for improved catalytic activity.* Oral. Feb. 21, 2023.

<https://doi.org/10.1016/j.bpj.2022.11.1754>

Molecular Machine Learning Conference; Cambridge, MA: Karvelis, E. *Modeling substrate turnover dynamics to guide the redesign of a natural enzyme for increased activity.* Poster. Oct. 21, 2022.

Bioengineering and Toxicology Seminar; Cambridge, MA: Karvelis, E. *Understanding and Re-engineering Enzymes Using Molecular Dynamics.* Oral. February 12, 2021.

BMES National Conference; Phoenix, AZ: Karvelis, E. *Modeling Glioblastoma Invasion with Microfluidics.* Oral. Oct. 14, 2017.

Energy Biosciences Building Colloquium; Berkeley, CA: Karvelis, E. *High-throughput Identification of Genetic Interactions Using Barcoded Transposon Mutants.* Oral. Aug. 4, 2017.

BMES National Conference; Minneapolis, MN: Karvelis, E., Ngo, M., Gilchrist, A., Kamm, R., Harley, B. *Utilizing Microfluidics to Recapitulate the Microenvironment of Glioblastoma.* Poster. Oct. 8, 2016.

## HOBBIES AND SERVICE

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I enjoy playing guitar, basketball, and strength training. During my time at MIT, I served as the BE Social Chair, a member of CSAIL's Coffee Committee, tutored, and was an instructor for Momentum AI. In the past, I volunteered for several years at an exotic pet sanctuary and am still passionate about animal welfare.