

## EDUCATION

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<b>University of Utah</b> Honors BS Biomedical Engineering with Chemistry Minor, GPA: 3.99	Salt Lake City, UT 2019–Present
<b>University of Utah</b> Honors BS Applied Mathematics, GPA: 3.99	Salt Lake City, UT 2019–Present
<b>Riverton High School</b> High School Diploma, GPA: 3.99	Riverton, UT 2015–2019

## EXPERIENCE

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<b>Visiting Researcher, MIT</b> Weiss Lab Research Aide	Cambridge, MA May 2022–August 2022
<ul style="list-style-type: none"><li>– Constructed biophysical models of neuromorphic genetic circuits</li><li>– Cleaned and generated statistical models of</li></ul>	scRNA-Seq Data  – Constructed neuromorphic genetic circuits <i>in vivo</i>
<b>Undergraduate Researcher, University of Utah</b> Bidone Lab Research Aid	Salt Lake City, UT August 2021–June 2022
<ul style="list-style-type: none"><li>– Analyzed simulations of E-Cadherin dynamics</li></ul>	
<b>Undergraduate Researcher, CU Boulder</b> Genetic Logic Lab Research Aide	Boulder, CO May 2020–Present
<ul style="list-style-type: none"><li>– Developed simulation tools for the computer-aided design of genetic circuits</li><li>– Adapted variance-reduction techniques for</li></ul>	stochastic simulation of genetic circuit dynamics – Developed novel stochastic simulation algorithms for the analysis of rare genetic events
<b>Undergraduate Researcher, University of Utah</b> Yellepeddi Research Group Research Aide	Salt Lake City, UT April 2020–December 2020
<ul style="list-style-type: none"><li>– Developed a Python library for rapid physiologically-based pharmacokinetics modelling of</li></ul>	nanoparticle biodistribution
<b>Tutor, High Performance Tutoring</b> Private Tutor	Salt Lake City, UT August 2018–Present
<ul style="list-style-type: none"><li>– Tutored Mathematics, Physics, Chemistry, Biology, and English at the high school and undergraduate</li></ul>	levels
<b>Research Intern, InnovaBio</b> SKIP Project Research Aide	West Jordan, UT August 2017–August 2019
<ul style="list-style-type: none"><li>– Utilized genetic engineering techniques to produce bioactive, recombinant human skeletal muscle and</li></ul>	kidney-enriched inositol phosphatase in <i>E. coli</i>
<b>Tutoring Assistant, Kumon</b> Private Tutor	West Jordan, UT August 2017–May 2018

## AWARDS AND HONORS

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- **SLCC SME Symposium First Place: Team Technical Poster, 2018** First place prize at the annual Salt Lake Community College Science, Math, and Engineering Symposium in the Team Technical Poster Category. Awarded for my team's research on the expression and solubility of SKIP in E. Coli.
- **National AP Scholar, 2018 and 2019** The highest level of distinction given by AP, granted to students in the United States who receive an average score of at least 4 on all AP Exams taken, and scores of 4 or higher on eight or more of these exams. Won twice.
- **National Merit Scholar, 2019** The highest level of distinction given by NMSC, given to students who have been judged to have the strongest combination of academic skills and achievements, extracurricular accomplishments, and potential for success in rigorous university studies.
- **Sterling Scholar Finalist: Science, 2019** A Sterling Scholar is a high school senior who is publicly recognized and awarded for the pursuit of excellence in scholarship, leadership, and citizenship in the State of Utah. The title of finalist is given to 14 students in each subject annually.
- **Goldwater Scholar Nominee, 2021** The Goldwater Scholarship is the most prestigious undergraduate scholarship in the natural sciences, mathematics, and engineering in America. Four students are nominated for the award by participating institutions each year.
- **UROP Recipient, 2021** UROP provides funding for students who assist with a faculty member's research. I was granted a UROP award for my work simulating E-Cadherin dynamics with Tamara Bidone.
- **SB<sup>3</sup>C Student Paper Contest Winner, 2022** Each year, the Summer Bioengineering, Biomechanics, Biotransport Conference (SB<sup>3</sup>C) selects 2 undergraduate papers from its international applicant pool as 'winners' for an undergraduate paper competition.

## PUBLICATIONS

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- [1] B. Shaikh, L. P. Smith, D. Vasilescu, G. Marupilla, M. Wilson, E. Agmon, H. Agnew, S. S. Andrews, A. Anwar, M. E. Beber, F. T. Bergmann, D. Brooks, L. Bruschi, L. Calzone, K. Choi, J. Cooper, J. Detloff, B. Drawert, M. Dumontier, G. B. Ermentrout, J. R. Faeder, A. P. Freiburger, F. Fröhlich, A. Funahashi, A. Garny, J. H. Gennari, P. Gleeson, A. Goelzer, Z. Haiman, J. Hasenauer, J. L. Hellerstein, H. Hermjakob, S. Hoops, J. C. Ison, D. Jahn, H. V. Jakubowski, R. Jordan, M. Kalaš, M. König, W. Liebermeister, R. S. M. Sheriff, S. Mandal, R. McDougal, J. K. Medley, P. Mendes, R. Müller, C. J. Myers, A. Naldi, T. V. N. Nguyen, D. P. Nickerson, B. G. Olivier, D. Patoliya, L. Paulevé, L. R. Petzold, A. Priya, A. K. Rampadarath, J. M. Rohwer, A. S. Saglam, D. Singh, A. Sinha, J. Snoep, H. Sorby, R. Spangler, J. Starrau, P. J. Thomas, D. van Niekerk, D. Weindl, F. Zhang, A. Zhukova, A. P. Goldberg, J. C. Schaff, M. L. Blinov, H. M. Sauro, I. I. Moraru, and J. R. Karr, "BioSimulators: a central registry of simulation engines and services for recommending specific tools", *Nucleic Acids Research*, May 2022, gkac331, ISSN: 0305-1048. eprint: <https://academic.oup.com/nar/advance-article-pdf/doi/10.1093/nar/gkac331/43614498/gkac331.pdf>.
- [2] T. Stoughton, L. Buecherl, P. J. Thomas, P. Fontanarrosa, and C. J. Myers, "Ibiosim server: A tool for improving the workflow for genetic design and modeling", *ACS Synthetic Biology*, 2022, [Submitted].

- [3] P. J. Thomas, M. Ahamdi, L. Buecherl, C. Winstead, C. J. Myers, and H. Zheng, “A comparison of weighted stochastic simulation methods for the analysis of genetic circuits”, *ACS Synthetic Biology*, 2022, [Submitted].
- [4] L. Buecherl, R. Roberts, P. Fontanarrosa, P. J. Thomas, J. Mante, Z. Zhang, and C. J. Myers, “Stochastic hazard analysis of genetic circuits in ibiosim and stamina”, *ACS Synthetic Biology*, vol. 10, no. 10, pp. 2532–2540, 2021, PMID: 34606710. eprint: <https://doi.org/10.1021/acssynbio.1c00159>.
- [5] T. Stoughton, L. Buecherl, P. J. Thomas, P. Fontanarrosa, and C. J. Myers, “Ibiosim server: A tool for improving the workflow for genetic design and modeling”, *IWBDA Conference 2021*, 2021.
- [6] P. J. Thomas, M. Ahmadi, H. Zheng, and C. J. Myers, “A comparison of weighted stochastic simulation methods”, *IWBDA Conference 2021*, 2021.