Trov A. Brier

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PROFESSIONAL SUMMARY

Chemist and software developer with a background in biological scientific computing. Experienced solving diverse problems in computational science settings applying novel techniques across domains. Skilled in the design and management of exploratory multi-year projects requiring flexibility, self-assessment mechanisms, and documentation suitable for publication. Proficient in presenting technical material to audiences with varying subject-matter expertise.

TECHNICAL SKILLS

Programming: Python, MATLAB, Bash, R, Tcl, ET_EX, C/C++, CUDA, HTML

Tools: Unix/Linux, MacOS, Windows, AWS, SVN, Git, SSH, Docker, Apptainer, slurm, Jupyter, VScode, MARTINI CG, NVIDIA DGX Software: pandas, matplotlib, seaborn, numpy, scipy, biopython, cobra, HTSeq, Guppy, minimap2, PyTorch, sklearn, HDF5, VMD, MAFFT, Clustal, BLAST, deepTools, SAMtools, GROMACS, AMBERMD, NAMD, AlphaFold2

Theoretical Training: Mathematics — Probability Theory, Stochastic Processes, Information Theory, Numerical Analysis, Linear Algebra, ODEs | Science & Engineering — Biophysics, Systems Biology, Bioinformatics, Molecular Dynamics, Equil. and Nonequil. Statistical Physics and Thermodynamics, Quantum Mechanics, Chemical Kinetics

EXPERIENCE

Graduate/Postdoctoral Research Assistant | The Luthey-Schulten Group, Dept. of Chemistry at UIUC

Jan 2018 - present

- Constructed a stochastic kinetic model using data collected from super-resolution imaging experiments to simulate the sugar-stress response in Escherichia coli.
- Collaborated in the use of GPU-accelerated, hybrid stochastic-deterministic methods to simulate whole-cell models of bacteria that capture the reaction and diffusion of \sim 10.000 biochemical components at the cell-scale.
 - * Constructed a kinetic model of the nucleotide metabolism for a bacteria cell.
 - * Curated genomics, transcriptomics, proteomics, metabolomics, and kinetic rates data.
 - * Updated software/code for use on high performance GPU cluster.
- Developed a methodology to profile the bacterial transcriptome coupling bioinformatics and experiment.
 - * Built a sequence alignment based pipeline to identify genetic motifs within a bacterial genome and predict transcriptional events.
 - * Analyzed Illumina, Pacific Biosciences, and Oxford Nanopore Technologies RNA sequencing experiments.
- * Created tools to predict transcription units and visualize the RNA isoforms.

 Assisted PI in preparation of grant materials to acquire federal funding: NSF MCB 2221237 \$2.00M (proposal, annual reports) | NSF MCB 1818344 - \$1.50M (annual reports) | NSF MCB 1840320 - \$1.18M (annual reports).
- Maintained group's GPU cluster and website.
- Collaborated with industrial and academic partners, both domestic and international.
- Presented research results to broader scientific community at domestic and international conferences.
- Mentored junior graduate and undergraduate student researchers.

Seminar/Workshop Teaching Assistant | The Luthey-Schulten Group, Dept. of Chemistry at UIUC May 2024, October 2021, July 2019

- NSF STC-QCB Advanced Computational Workshop 2024—Mentored junior graduate students to construct teaching material, installed software on supercomputer, and troubleshot errors during live tutorials.
- Online Hands-on Workshop on Computational Biophysics—Held virtual lecture for graduate students and post-doctoral and prepared tutorials run on AWS instances.
- Center for the Physics of Living Cells (CPLC) Summer School—Held lectures for graduate students and post-doctoral associates and prepared tutorials run on AWS instances.

Undergraduate Research Assistant | The Baker Group, Dept. of Chemistry at TCNJ

Dec 2014 - May 2017

- Explored atomistic and coarse grained molecular dynamics simulations of membrane bound proteins
- Presented research results to broader scientific community at domestic conferences.
- Mentored junior undergraduate student researchers.

EDUCATION

University of Illinois at Urbana-Champaign (UIUC) The College of New Jersey (TCNJ)

PhD in Chemistry **BS in Chemistry (ACS-certified)** Aua 2017 - Dec 2024 Aug 2013 - May 2017

- T. A. Brier, J. E. Cournoyer, B. R. Gilbert, S. A. Glass, Y. Gao, Z. R. Thornburg, K. Goglin, G. John, T. Mamaghani, S. Shivakumar, Y. Yu, C. Fields, J. I. Glass, A. P. Mehta, Z. Luthey-Schulten, Unraveling the Transcriptional Landscape within a Minimized **Bacterium via Comparative Analysis**, in preparation
- J. A. Stevens, F. Grünewald, P.A. Marco van Tilburg, M. König, B. R. Gilbert, T. A. Brier, Z. R. Thornburg, Z. Luthey-Schulten, S. J. Marrink, Molecular Dynamics Simulation of an Entire Cell, Frontiers in Chemistry 2023
- Z. R. Thornburg, D. M. Bianchi, T. A. Brier, B. R. Gilbert, T. M. Earnest, M. C.R. Melo, N. Safronova, J. P. Sáenz, A. T. Cook, K. S. Wise, C. A. Hutchison III, H. O. Smith, J. I. Glass, Z. Luthey-Schulten, Fundamental Behaviors Emerge from Simulations of a Living Minimal Cell. Cell 2022
- D. M. Bianchi, T. A. Brier, A. Poddar, M. S. Azam, C. K. Vanderpool, T. Ha, Z. Luthey-Schulten, Stochastic Analysis Demonstrates the Dual Role of Hfq in Chaperoning E. coli Sugar Shock, Frontiers in Molecular Biosciences 2020