Samuel Christensen

Education	University of California, Los Angeles (UCLA)	Los Angeles, CA
	- Ph.D . In Diomathematics	Sep 2017 – Present
	- BS . in Honors Mathematics & Biophysics	Sep 2013 – June 2017
Skills	Numerical Analysis, Image Analysis, Finite Element Analysis, Linear Algebra, Machine Learning, Python, Pandas, Scikit, MATLAB, Julia, C++, Comsol, SolidWorks, Probability Theory, Stochastic Processes, Partial Differential Equations, and Data Analysis.	
Publications	 Christensen S., Huang Y., Walch O. J., Forger D. B. Optimal adjustment of the human circadian clock in the real world. <i>PLOS Computational Biology</i> 2020. Christensen S., Chu R., Anderson C., Roper M. Fast Asymptotic-Numerical Method For Coarse Mesh Particle Simulation In Channel Of Arbitrary Cross Section. <i>Journal of Computational Physics</i> 2022. Christensen S., Roper M. Inertial Migration in Micro-Centrifuge Devices. Submitted to <i>Physical Review Fluids</i> 2023. Christensen S., Roper M. Quasi-Steady Inertial Migration in Arbitrary Geometries. In preparation for submission to <i>Physical Review Fluids</i> 2023. 	
Awards	– Ursula Mandel Medical Science Research Scholarship, UCLA	2018
	- NSF Systems and Integrative Biology Training Grant, NSF	2018, 2019
Research Experience	 Physics of Microfluidic Devices, UCLA Created fast and accurate models for cell movement in models symbolic and numerical techniques, published in <i>The Journal</i> Optimized design of cell sorting chambers within microfluid increase sorting precision, published in <i>Physical Review Fluids</i> Developed custom cell tracking software from video data that deeper understanding of cell motion within a microfluidic device Optimal Sleep Schedules for Overcoming Jet Lag in Minimum T Leveraged data from wearable devices, mathematical models, sleep schedules for overcoming jet lag under realistic human called 'Entrain' with more than 80,000 users and published in Genetic Heritability in Fungal Spores, UCLA Developed hierarchical Bayesian classifier to distinguish fluor Machine Learning Analysis of Website Data for Fingerhut.com Guided teams of undergraduate students to train a classifier th result in a purchase. 	2020-Present derate Reynolds number flows using of Computational Physics. idic devices using custom models to tracked a cell's 3D position, enabling vice. Time, University of Michigan 2016-2018 and control theory, I created optimal n constraints. Resulted in an iOS app PLOS Computational Biology. 2022 rescent probes in spore histones. n, UCLA 2023 and predicts which visits were likely to
Coding Projects	 3-D Fluid Flow Solver in Matlab, MATLAB, COMSOL Wrote custom Finite Element Navier-Stokes Solver that coubetween a cell and the surrounding fluid. Exploited sparse matrix and parallelized structure to accelerate Developed custom preconditioner application which reduced restructure and implemented 3-D cell tracking algorithm from computational solve of the fluid flow, and Kalman filter. Enabled engineers to see cell movements in additional dimense Nonlinear Optimal Control Algorithm, MATLAB Implemented optimal control algorithm to create optimal scircadian rhythm in minimum time. Adapted algorithm to ensure sleep schedules were customized. Approximate Bayesian Inference to Classify Microscopy Data for Created classifier that identifies nucleus type in fungal spores. Feature Analysis, Identification, and Classification of Website users by other states. 	ald quickly calculate the interaction te code by a factor of 10. matrix inversion times by a factor of 2. <i>ATLAB</i> n 2-D data by using template images, sion. sleeping schedules for changing the d to user set conditions. <i>Python</i> fluorescently tagged histones. based on fluorescence data. Data , <i>Python</i> commercial potential.