Joshua Burton

Education

2018 - 2023	 Ph.D., University of Manchester Quantitative and Biophysical Biology. Thesis title: Bayesian inference and modelling of gene expression dynamics.
	• Derived a novel Kalman filtering algorithm for a delay differential equation model of gene expression
	• Developed Python and Julia packages hesdynamics and DelayedKalmanFilter.jl to enable Bayesian infer- ence on stochastic delay differential equation models
	• Contributed bug fixes and documentation to open-source Julia packages Pathfinder.jl and TuringGLM.jl
	• Learnt how to effectively communicate difficult mathematical concepts to biologists
2014 – 2018	 ◊ MMath (Hons), 1st Class in Mathematics. Thesis title: An exploration of multistationarity in chemical reaction networks.
	Overall average above 80%
	• Had a lead supervisor teaching role in my final year, teaching over 20 first-year students
Skills	

Math	 Bayesian inference, MCMC, dynamical systems, SDEs
Computing	g ◇ Julia, Python, R, 啓正X, Linux, Git, Excel
Other technical skill	 Academic research and teaching, writing publications, public speaking, independent remote working, strong organisational skills
Interest	◊ Drumming, jazz, volleyball, films

Awards and Achievements

- - ◊ Invited speaker at the Royal Statistical Society Invited Session: Statistical inference in stochastic biological systems with complex dynamics. Institute of Mathematical Statistics Annual Meeting (London, June)
 - ◊ Poster presentation at the EMBO | EMBL Symposium: Biological oscillators: Design, mechanism, function (Heidelberg, March)
 - ♦ **Speaker** at the Centre for Biological Timing Winter Symposium (Manchester, February)

Work Experience

2023 – Present \diamond Julia software developer and Bayesian inference researcher, PlantingSpace.

- Contributed to building a general knowledge system which is transparent, and incorporates uncertainty
- Research at the intersection of Bayesian statistics and Category theory
- Involved in multiple aspects of an early stage start up, including backend development (Julia), deployment and infrastructure (AWS, docker)
- Highly collaborative environment with a flat structure, which allowed for experience as both a team member and a team leader
- 2017 2019 \diamond **Graduate teaching assistant**, University of Manchester Mathematics department.
 - Taught foundational mathematical concepts to first-year university students
 - · Took a lead supervisor role and provided letters of reference for students

2017 \diamond **Intern**, Wellcome Trust summer studentship scheme.

- Worked with large data sets of high-throughput single-cell sequencing data
- Gained a mathematical understanding of multiple ML algorithms

Research Publications

Journal Articles

 C. E. Overton, L. Pellis, H. B. Stage, et al., "EpiBeds: Data informed modelling of the COVID-19 hospital burden in England," PLOS Computational Biology, vol. 18, no. 9, e1010406, Sep. 2022, Publisher: Public Library of Science, ISSN: 1553-7358. O DOI: 10.1371/journal.pcbi.1010406.

- 2 X. Soto, J. Burton, C. S. Manning, *et al.*, "Sequential and additive expression of miR-9 precursors control timing of neurogenesis," *Development*, vol. 149, no. 19, Oct. 2022, ISSN: 0950-1991. *9* DOI: 10.1242/dev.200474.
- J. Burton, C. S. Manning, M. Rattray, N. Papalopulu, and J. Kursawe, "Inferring kinetic parameters of oscillatory gene regulation from single cell time-series data," *Journal of The Royal Society Interface*, vol. 18, no. 182, Sep. 2021, ISSN: 1742-5662.
 DOI: 10.1098/rsif.2021.0393.

Pre-prints

S. Funk, S. Abbott, B. D. Atkins, et al., Short-term forecasts to inform the response to the Covid-19 epidemic in the UK, Dec. 2020. DOI: 10.1101/2020.11.11.20220962.

In preparation

J. Burton, M. Rattray, N. Papalopulu, and J. Kursawe, *Continuous time filtering and variational inference of combined single cell time-series data*, 2023.