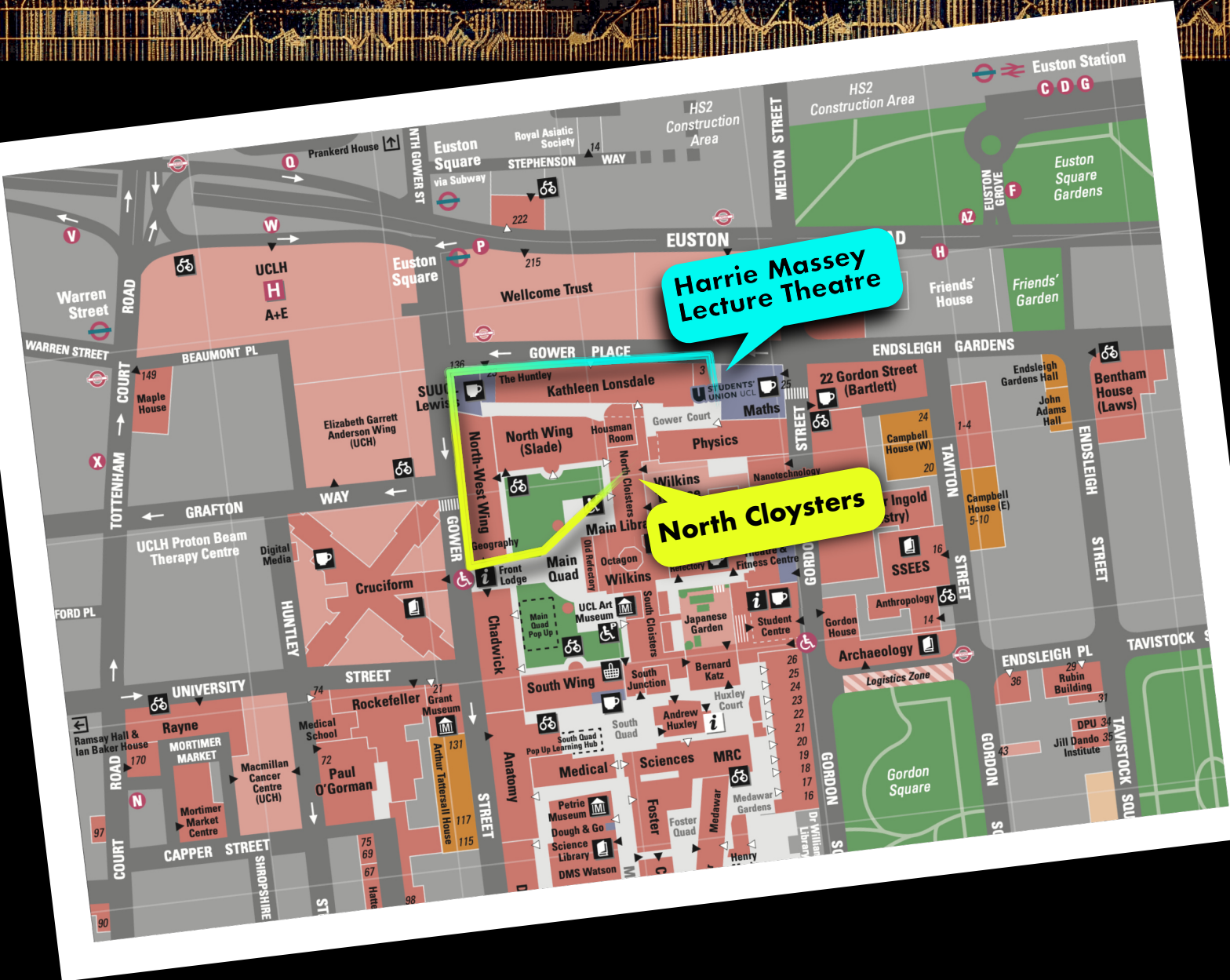


# LONDON MATHEMATICAL BIOLOGY CONFERENCE

UCL 18-19 September 2023



# LONDON MATHEMATICAL BIOLOGY 2023 SCHEDULE

All talks will be held in **Harrie Massey Lecture Theatre**

## Monday 18th September

- 10:30 - 10:40** Welcome
- 10:40 - 11:15** **Wenying Shou** - *University College London*  
1, 2, 3...
- 11:15 - 11:30** **Jessie Renton** - *Queen Mary University of London*  
A somatic genetic clock for clonal organisms
- 11:30 - 11:45** **Andrea Cairoli** - *The Francis Crick Institute*  
Data-driven modelling of abdominal tissue growth in *Drosophila* during metamorphosis
- 11:45 - 12:50** Lunch and Posters (North Cloisters)
- 12:50 - 13:25** **Omer Karin** - *Imperial College London*  
Self-tuning of long transients in cell biology
- 13:25 - 13:40** **Cara Neal** - *University of Liverpool*  
The role of non-Newtonian fluids in microswimmer propulsion
- 13:40 - 13:55** **Euan Smithers** - *Sainsbury Laboratory*  
How do plant leaf pavement cells pattern to form puzzle piece-like shapes?
- 13:55 - 14:30** Coffee Break and Posters (North Cloisters)

- 14:30 - 15:05**   **Berta Verd** - *University of Oxford*  
Evolving vertebral counts:  
a mathematical modelling approach
- 15:05 - 15:20**   **Santosh Manicka** - *Tufts University*  
Information integration during bioelectric regulation  
of morphogenesis of the embryonic frog brain
- 15:20 - 15:35**   **Kristian Kiradjev** - *University of Nottingham*  
Multiscale asymptotic analysis reveals how cell  
growth and subcellular compartments affect  
tissue-scale hormone transport
- 15:35 - 16:00**   Coffee Break and Posters (North Cloisters)
- 16:00 - 16:35**   **Carina Dunlop** - *University of Surrey*  
What do cells and tissues feel? Integrating cell  
contractility, adhesion and ECM stiffness in  
mechanosensing.
- 16:35 - 17:10**   **Andrew Krause** - *University of Durham*  
Rethinking Pattern Formation through Nonlinear  
Dynamics and Web Browsers
- 17:10 - 17:25**   **Martina Oliver** - *Imperial College London*  
Modelling and parameter inference in synthetic  
reaction-diffusion patterns
- 18:00**   Conference dinner (Cabana)

## Tuesday 19th September

- 09:00 - 09:35** **Mohit Dalwadi** - *University College London*  
A universal spatiotemporal robustness in pattern formation
- 09:35 - 10:10** **James di Frisco** - *The Francis Crick Institute*  
The complementarity of mathematical and conceptual theoretical biology
- 10:10 - 10:25** **Svetlana Petrenko** - *University College London*  
Modelling of Silica Pattern Formation in Diatoms Using a Continuum Approach
- 10:30 - 11:00** Coffee Break and Posters (North Cloisters)
- 11:00 - 11:35** **Barbara Bravi** - *Imperial College London*  
Machine learning-guided modelling of immune protein interactions
- 11:35 - 11:50** **Mohit Kumar Jolly** - *Indian Institute of Science*  
Landscape of epithelial-mesenchymal plasticity as an emergent property of coordinated teams in regulatory networks
- 11:50 - 12:05** **Patricia Lamirande** - *University of Oxford*  
Mean First Passage Time and its Application in Ocular Drug Development
- 12:05 - 12:20** **Yaron Ben-Ami** - *University of Oxford*  
A probabilistic-continuum two-phase model for tumour cell migration induced by interstitial flow
- 12:20 - 13:20** Lunch (North Cloisters)
- 13:20 - 14:20** **Plenary talk: Philip Maini** *University of Oxford*  
Modelling collective cell movement: mathematical challenges and biological applications



# POSTER CONTRIBUTIONS

## Monday 18th September

**Matthew Asker** - *University of Leeds*

Coexistence of competing microbial strains under twofold environmental variability and demographic fluctuations

**Johannes Borgqvist** - *Wolfson Centre for Mathematical Biology, Mathematical Institute, University of Oxford*

Construction of travelling wave models of collective cell migration using Lie symmetries of the Fisher KPP model

**Joshua Bull** - *University of Oxford*

Your Space or Mine? Mathematical quantification of inter-patient tumour heterogeneity

**Jurjen Duintjer Tebbens** - *Faculty of Pharmacy, Charles University Prague*

Reaction-diffusion models for PXR-induced drug metabolism

**Giulia Laura Celora** - *University College London*

Self-organised patterning in Dictyostelium group migration

**Calum Gabbutt** - *Institute of Cancer Research*

Lineage tracing in patients with blood cancer using fluctuating DNA methylation

**Poulami Somanya Ganguly** - *Queen Mary University of London*

Models of extrachromosomal DNA replication during cancer cell division

**Magnus Haughey** - *Barts Cancer Institute, Queen Mary University of London*

Investigating spatial signatures of extracellular DNA with agent-based computational modelling

**Diana Khoromskaia** - *The Francis Crick Institute*

Morphogen transport in tissues as active porous media

**Nandakishor Krishnan** - *Centre for Ecological Research, Budapest*

The evolution of symbiosis in the context of eukaryogenesis

**Jamie Lee** - *Imperial College London*

In silico-guided treatment design for skin damaging *S. aureus*-*S. epidermis* colonisation in atopic dermatitis lesions

**Ves Manojlovic** - *City, University of London*

Fluctuating methylation clocks for inferring the evolutionary history of human cancers

**Taniya Mandal** - *The Francis Crick Institute*

Exploring evolutionary variations in forebrain development using patterned organoids

**Charlotte Manser** - *Imperial College London*

A Mathematical Framework for Tuning Tempo in Developmental Gene Regulatory Networks

**Andela Markovic** - *University College London*

Communication is the key: positional information transmission in neural tube patterning

**Nathaniel Mon Pere** - *Barts Cancer Institute*

Clonal interference in aging haematopoiesis

**Roozbeh Pazuki** - *Imperial College London*

Upper limits on the robustness of Turing models and other multi-parametric dynamical systems

**Domènec Ruiz-Balet** - *Imperial College London*

The tragedy of the commons in mean-field games

**Hugh Selway-Clarke** - *Imperial College London*

In Silico Testing of Hypotheses for the Effect of Smoking on Somatic Evolution in the Healthy Human Lung

**Kimberley Verity** - *City, University of London*

Improved tree indices under the Yule and Uniform models.

# POSTER CONTRIBUTIONS

## Tuesday 19th September

**Andrew Bate** - *University of Leeds*

Modelling respiratory virus exposure on local buses

**Amy Bowen** - *The Francis Crick Institute*

TBC

**Prakrati Dangarh** - *Imperial College London*

Mechanistic modelling of pre-school wheezing and progression to school-age asthma

**Luke Davis** - *University College London*

Understanding biomolecular condensates through statistical mechanical and minimal modelling approaches

**Anqi Huang** - *The Francis Crick Institute*

Brk feedback motif underlies precise decoding of the morphogen tail

**Ferdinando Insalata & Daniel Kornai** - *Imperial College London*

Stochastic survival of the densest can solve the enigma of the expansion of mitochondrial mutations in the ageing of skeletal muscle fibres

**Jack Jennings** - *Sheffield University*

Understanding self-organised tissue patterning across scales

**Jacob Jepson** - *University of Nottingham*

Modelling phloem transport and sucrose delivery within the plant seedling

**Juan David Marmolejo Lozano** - *University of Los Andes*

A generalised model for noise propagation in transcriptional genetic cascades

**Xiaoyuan Liu** - *University of York*

Parthenogenesis, sexual conflict and the evolution of oogamy

**Norberto Lucero Azuara** - *Queen Mary University of London*

Modeling movement in two dimensions by fractional Brownian motion

**Antonio Matas Gil** - *Imperial College London*

Unraveling Biochemical Spatial Patterns: Physics Informed Neural Networks for solving the inverse problem in Turing systems

**Christo Morison** - *Queen Mary University of London*

Single-cell mutational burden distributions in birth-death processes

**Lewis Mosby** - *Lewis Mosby*

Evolving Tissue Pattern Scaling and Robustness Through Spatially Heterogeneous Feedback

**Frixos Papadopoulos** - *University of Southampton*

Simpler sequence Histograms outperform word2vec-based Representations across Protein Inference tasks

**Paul Piho** - *Imperial College London*

Quantifying the fitness effects of stochastic gene expression

**Elisa Scanu** - *Queen Mary University of London*

Coordinated inheritance of multiple extrachromosomal DNA species in human cancer cells

**Alan Scaramangas** - *Queen Mary University of London*

Evolutionary and eco-evolutionary stability of Batesian mimicry systems

**Dimitris Volterras** - *Imperial College London*

Understanding cell-cycle dependent transcription dynamics using single-cell RNA-sequencing data