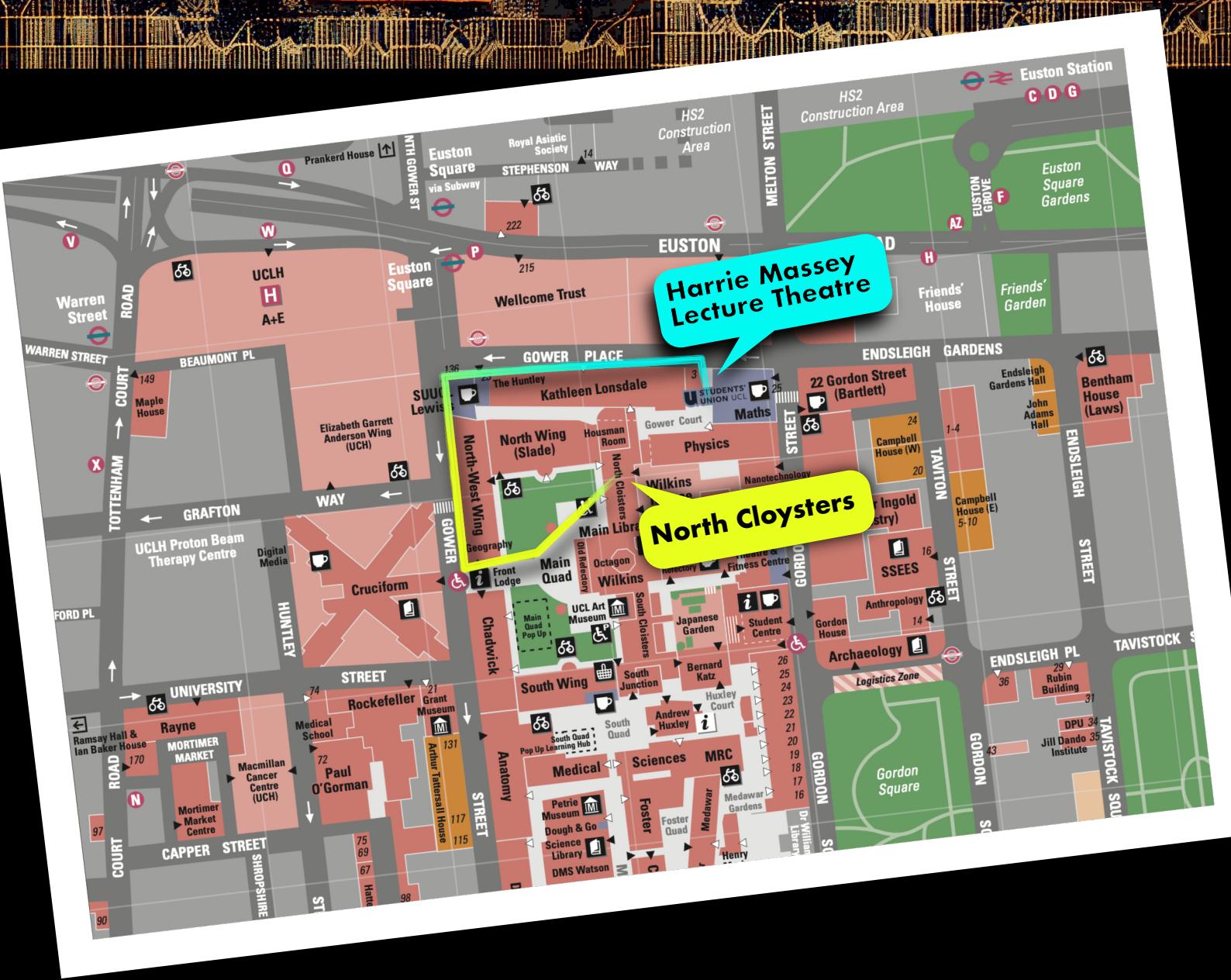


# LONDON MATHEMATICAL BIOLOGY CONFERENCE

# 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 Kiradjiev** - *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 Gabbett - *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. epidermidis 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 Pihó** - *Imperial College London*

Quantifying the fitness effects of stochastic gene expression

**Elisa Scaru** - *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 Volteras** - *Imperial College London*

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