Modern Techniques in Modelling



Who we are



Course organisers

- Nicholas Davies, PhD <u>Nicholas.Davies@lshtm.ac.uk</u>
- Yang Liu, PhD <u>Yang.Liu@lshtm.ac.uk</u>
- Oli Brady, PhD <u>Oliver.Brady@lshtm.ac.uk</u>

Course administration

• Francesco Grisolia <u>Francesco.Grisolia@lshtm.ac.uk</u>

Lecturers and Demonstrators

• Billy Quilty, Kath O'Reilly, Seb Funk, Johnny Filipe, Alexis Robert, Alex Richards, Kaja Abbas (All LSHTM / CMMID-based)





A certificate of attendance will be issued automatically by LSHTM's short courses team (look out for an e-mail next week).

Please complete the feedback form on Moodle after the course — tell us what we did well and what we could improve.

Feel free to contact us if you have any questions on the course material or about your modelling work!

Recommended textbooks









Model Fitting and Inference for Infectious Disease Dynamics

Overview Course objectives How to apply

Overview

Course dates: Feb 2025

The course will take place in London, UK.

A short course taught by members of the Centre for the Mathematical Modelling of Infectious Diseases.

There is a growing demand for mathematical modellers in public health to explain observed disease trends and predict the outcome of interventions, often by synthesising information from different data sources. At the same time, increasing computational power and methodological advances are providing exciting opportunities to fit ever more complex mechanistic models to data. In light of the speed of methodological advances and the broad nature of the field, the task of choosing from the available methods and packages, as well as putting them into practice, can be daunting.





ℜ Course organisers

Which models did we see in the course?



Difference equations Tracks the number of individuals in each epidemiological "compartment" (e.g. Infected or Susceptible) at each e.g. day or week timestep	Ordinary Differential Equations (ODEs) Same as 'difference equations' but instead of calculating at each timestep, we move to continuous time	Metapopulation Add in structure to ODE model by creating multiple subpopulations that can transmit infections within and between each subpopulation
Individual-based model	Network model	Stochastic compartment model
Tracks each individual, each with their own epidemiological	Adds structure to the individual-based model, where each individual is	A stochastic implementation of our compartment ODE



Main Question: how do we choose a model type and a model structure?

Key principle: build with parsimony ("as simple as necessary")

- What is the research question?
- How big is the population?
- Are there stochastic fluctuations in the data that cannot be mechanistically accounted for?
- Do we need to track every individual?
- What type of events are we modelling and how do we parameterise them?
- What type of data do we have?





Any final questions?

Photo time!

