

MATHEMATICAL MODELS IN EPIDEMIOLOGY (MME)

MTB Group

March 17th – May 5th, 2023
Friday (6 sessions)
April 7 and April 14, 2023 – Easter Break
11:45-17:15 (a total of 24 hours)

Epidemiological models are important tools to understand the dynamics of infectious diseases, contributing to public health authorities' capacity to implement the available intervention measures to control disease transmission. As the COVID-19 pandemic progressed, research on mathematical modeling became imperative and very influential, to understand the epidemiological dynamics of disease spreading and control under different scenarios. This course provides an introduction to mathematical epidemiology, from mathematical concepts and applications up to discussion of real world case studies, including the COVID-19 pandemic.

PROGRAMME

1. Mathematical concepts and applications

Intro deterministic and stochastic epidemiological models

Stability analysis, complex eigenvalues, bifurcation diagrams, state space plots

Basic concepts on epidemiology: reproduction number, growth rates, relative risk and vaccine efficacy

Data on epidemiology

Model validation with empirical data

Model projections

Infectious disease epidemiology

Multi-strain models

Chaotic dynamics

Lyapunov exponents, predictability

Model validation and data analysis

Control measures (vaccination and other)

Epidemic processes in complex networks

2. Case studies

a. Epidemiological concepts

b. Dengue fever: multi-strain, within-host, vaccine

- c. COVID-19 pandemic: data analysis, epidemiological measures, projections, prediction
- d. Other public health threads

March 17		March 24		March 31	
Introduction to MME	The SIR epidemic model	The SHAR epidemic model	Case study: Modeling COVID-19	Stochastic systems	Dengue, HIV, etc
The SIS epidemic Model		Concepts in Epidemiology: R_0 and growth rates	SIR with seasonality and introduction to stochastic processes		Mathematical models applied to infectious disease dynamics: case studies
April 21		April 28		May 5	
Network theory	Bifurcation analysis	Parameter estimation	Optimal control (vectorborne diseases)	Multi-strain dengue models	COVID-19 models
	Modeling chagas disease	Model selection	Optimal control (HIV)		

PREREQUISITES:

Basic knowledge of ordinary differential equations, linear algebra, and probabilities/statistics

REFERENCES:

[1] Maíra Aguiar, Nico Stollenwerk and Bob W. Kooi (April 20th 2012). Modeling Infectious Diseases Dynamics: Dengue Fever, a Case Study, Epidemiology Insights, Maria de Lourdes Ribeiro de Souza da Cunha, IntechOpen, DOI: 10.5772/31920. Available from: <https://www.intechopen.com/chapters/35765>

[2] Stollenwerk, N., Jansen, V.: Population Biology and Criticality: From Critical Birth–Death Processes to Self-Organized Criticality in Mutation Pathogen Systems. World Scientific, London (2011)

[3] Aguiar, M., Kooi, B., Stollenwerk, N. Epidemiology of Dengue Fever: A model with temporary cross immunity and possibly secondary infection shows bifurcations and chaotic behaviors in wide parameter region. Math. Model. Nat. Phenom. Vol 3 (4) 48-70 (2008).

[4] Aguiar, M., Ballesteros, S., Kooi, B.W., Stollenwerk, N. The role of seasonality and import in a minimalistic multi-strain dengue model capturing differences between primary and secondary infections: complex dynamics and its implications for data analysis. Journal of Theoretical Biology, 289, 181-196 (2011).

[5] Aguiar, M., Stollenwerk, N., Kooi, W.B. Scaling of stochasticity in dengue hemorrhagic fever epidemics. Mathematical Modelling of Natural Phenomena. 7, 1-11, (2012)

- [6] L. Mateus, N. Stollenwerk and J.C. Zambrini, Stochastic Models in Population Biology: From Dynamic Noise to Bayesian Description and Model Comparison for Given Data Sets, *Int. Journal. Computer Math.*, 90, 2161-2173, (2013).
- [7] Kooi W. B., Aguiar, M. and Stollenwerk, N. Analysis of an asymmetric two-strain dengue model. *Mathematical Biosciences*. 248, 128-139, (2014).
- [8] Aguiar, M., Paul, R., Sakuntabhai, A., Stollenwerk, N.: Are we modeling the correct data set? Minimizing false predictions for dengue fever in Thailand. *Epidemiol. Infect.* 142, 2447--59 (2014).
- [9] Aguiar, M.; Stollenwerk, N.; and Halstead. S.B. Modeling the impact of the newly licensed dengue vaccine in endemic countries. *PLoS Neglected Tropical Diseases* 10(12), e0005179 (2016)
- [10] Aguiar, M., Stollenwerk, N. The impact of serotype cross-protection on vaccine trials: DENVax as a case study (2020). *Vaccines*. 8(4), 674 (2020).
- [11] Aguiar, M., Millán Ortuondo, E., Bidaurrazaga Van-Dierdonck, J., Mar, J., Stollenwerk, N.: Modelling COVID 19 in the Basque Country from introduction to control measure response. *Sci. Rep.* 10, 17306 (2020)
- [12] Aguiar, M., Bidaurrazaga Van-Dierdonck, J., Stollenwerk, N.: Reproduction ratio and growth rates: measures for an unfolding pandemic. *PLoS ONE* 15, e0236620 (2020)
- [13] Aguiar, M., Bidaurrazaga Van-Dierdonck, J., Mar, J., Cusimano, N., Knopoff, D., Anam, V., Stollenwerk, N.: Critical fluctuations in epidemic models explain COVID-19 post-lockdown dynamics. *Sci. Rep.* 11, 13839 (2021)
- [14] Pastor Satorras, R., Castellano, C., Van Mieghem, P., Vespignani, A. Epidemic processes in complex networks. *Reviews of Modern Physics* 87, 925 (2015).
- [15] Maíra Aguiar, Joseba Bidaurrazaga Van-Dierdonck, Javier Mar, Nico Stollenwerk (2021). The role of mild and asymptomatic infections on COVID-19 vaccines performance: a modeling study. *Journal of Advanced Research* (In Press). <https://doi.org/10.1016/j.jare.2021.10.012>
- [16] Maíra Aguiar, Vizda Anama, Konstantin B. Blyuss, Carlo Delfin S. Estadilla, Bruno V. Guerrero, Damián Knopoff, Bob W. Kooi, Akhil Kumar Srivastava, Vanessa Steindorf, Nico Stollenwerk. (2022). Mathematical models for dengue fever epidemiology: a 10 year systematic review. *Physics of Life Review*, Volume 40, March 2022, Pages 65-92. <https://doi.org/10.1016/j.plrev.2022.02.001>
- [17] Steindorf, V.; Srivastav, A.K, Stollenwerk, N., Kooi, B.W., Aguiar, M. (2022). Modelling secondary infections with temporary immunity and disease enhancement factor: mechanics for complex dynamics in simple epidemiological models. *Chaos, Solitons & Fractals* 164:112709.

***Registration is free, but inscription is required before March 10th, 2023:** So as to inscribe go to <https://forms.gle/cpA3B7zcGnFBB7FX6> and fill the registration form.