

Statistics in Medicine (3 units)

Statistics in Medical Practice (M12)

Lecturers from the MRC Biostatistics Unit

This part of the course includes three modules covering a range of statistical methods and their application in three areas of biostatistics.

A. Multi-State Models in Health [4 Lectures] (C. Jackson, D. De Angelis, P. Birrell)

(i) Continuous-time multi-state models, applied to modelling the incidence and progression of chronic diseases. Transition intensities, transition probabilities, and describing quantities of applied interest as functions of these. Constructing likelihoods for parametric models given various forms of individual-level data. Interpreting and comparing estimates from fitted models.

(ii) Multi-state modelling to estimate incidence of infectious diseases from population-level data. Backcalculation methods for the estimation of incidence of disease with long incubation periods. Dynamic modelling of infectious disease transmission.

B. Bayesian nonparametric models in molecular medicine [4 Lectures] (P. Kirk, J. Vd-molen, L. Ronneberg)

Molecular medicine is a rapidly evolving field that relies on accurate modelling and analysis of complex biological data. Much of the work undertaken in molecular medicine and statistical omics relies on the identification of latent structures, ranging from discrete subgroups (such as molecular subtypes of cancer) to continuous latent functions (such as drug dose-response relationships). Bayesian nonparametric models, such as Dirichlet process mixture (DPM) models and Gaussian processes (GPs), allow for flexible and data-driven modelling and have the potential to provide deeper insights into gene expression, disease subtypes, and drug responses. In this module we will introduce, and explore the principles of Bayesian nonparametric modelling approaches for latent structure discovery. We will present DPM models for uncovering population subgroups and the use of GP regression for modelling latent functions. Throughout, we will illustrate with specific examples from molecular medicine, but would emphasise that no previous experience of molecular biology is required for this module.

C. Design and Analysis of Clinical Trials [4 Lectures] (M. Law, D. Couturier, S. Villar, D. Robertson)

Introductory concepts in clinical trial design and analysis, including sample size estimation. Early phase dose-finding clinical trials using the continual reassessment method and Bayesian parametric models. Types of randomisation procedures. Optimal response-adaptive procedures. Adaptive and multi-stage designs (including group-sequential designs). Treatment effect estimation following a group-sequential trial.

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Analysis of Survival Data (L12)

P. Treasure

This part of the course includes three modules covering the fundamentals of time-to-event analysis with applications to cancer survival.

D. Time-to-Event Analysis [4 Lectures]

‘Survival analysis’ is generalised to *time-to-event* analysis. The implications of event times which are unknown or in the future (*censored* data) are discussed. Time-to-event distributions are introduced and their parametric (maximum likelihood) and non-parametric (*Kaplan-Meier*) characterisations are described. Methods for comparing two time-to-event distributions (as in a clinical trial of an active treatment versus a placebo) are derived (*log-rank* test).

E. Modelling Hazard [4 Lectures]

The *hazard* function (instantaneous event rate as a function of time) is defined. It is shown how the hazard function can naturally be used to model the effect of explanatory variables (such as age, gender, treatment, blood pressure, tumour location and size...) on the time-to-event distribution (*proportional hazards* modelling). Model checking procedures are introduced with an emphasis on excess event (*Martingale*) plots.

F. Population Cancer Survival Analysis [4 Lectures]

Analysis of survival data from real-world cancer studies is complicated by patients also being at risk from other causes of death. Methods of dealing with more than one cause of death are presented for the cases (i) the cause of death is known (*competing risk* analysis) and (ii) the cause of death is unknown (*net survival*). The conceptual difficulties inherent in the notion of a cancer survival distribution relevant to a particular calendar year (e.g. 2019) are addressed: *period* survival analysis.

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Additional Information

Prerequisites

Undergraduate-level statistics and probability: including analysis and interpretation of data, maximum likelihood estimation, hypothesis testing, basic stochastic processes.

Literature

There are no course books, but relevant medical papers may be made available before some of the lectures for prior reading. Some literature to complement the course material is listed below.

1. Armitage P, Berry G, Matthews JNS, *Statistical Methods in Medical Research*. Wiley-Blackwell, 2001. [A good introductory companion to the whole course]
2. van den Hout, A, *Multi-State Survival Models for Interval-Censored Data*. Chapman and Hall, 2016 [Module A]
3. Keeling, MJ, & Rohani, P *Modeling Infectious Diseases in Humans and Animals*. Princeton University Press, 2008 [Module A]
4. Rasmussen, CE and Williams, CKI *Gaussian Processes for Machine Learning* <https://gaussianprocess.org/gpml/> [Module B]

5. Neal, RM (2000). Markov Chain Sampling Methods for Dirichlet Process Mixture Models. *Journal of Computational and Graphical Statistics*, 9(2), 249–265. (Section 2) Available from: <https://glizen.com/radfordneal/ftp/mixmc.pdf> [Module B]
6. Teh, YW (2010). Dirichlet Process. *Encyclopedia of Machine Learning*, 1063, 280-287. Available from: <https://www.stats.ox.ac.uk/~teh/research/npbayes/Teh2010a.pdf> [Module B]
7. Senn, S. *Statistical Issues in Drug Development*. Wiley, 2021. [Module C]
8. Jennison C, Turnbull B, *Group Sequential Methods with Applications to Clinical Trials*. Chapman and Hall, 2000. [Module C]
9. Rosenberger, William F., and John M. Lachin. *Randomization in clinical trials: theory and practice*. John Wiley & Sons, 2015. [Module C]
10. Cox DR, Oakes D, *Analysis of Survival Data*. Chapman and Hall, 1984 [Modules D, E, F: the classic text]
11. Collett D, *Modelling Survival Data in Medical Research*. CRC Press, 2023 [Modules D, E, F: modern, applied, supports and extends lectures.]
12. Aalen OO, Borgen Ø, Gjessing HK, *Survival and Event History Analysis: A Process Point of View*. Springer, 2008 [Modules D, E, F: excellent modern approach]

Additional support

[Modules A, B, C] One-hour example classes for each of the three modules, supported by question sheets and solutions, will be given in Michaelmas Term.

[Modules D, E, F] A 90-minute example class, supported by question sheets and solutions, will be given in each of the Lent and Easter Terms. A two-hour revision class will be held just before the examination. There will be regular office hours by Zoom.