

Mathematics and Statistics Undergraduate Handbook

Supplement to the Handbook

Honour School of Mathematics and Statistics Syllabus and Synopses for Part C 2009–2010 for examination in 2010

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Every effort is made to ensure that the list of courses offered is accurate at the time of going online. However, students are advised to check the up-to-date version of this document on the Department of Statistics website.

Notice of misprints or errors of any kind, and suggestions for improvements in this booklet should be addressed to the Academic Administrator in the Department of Statistics.

Updated October 2009

1 Honour School of Mathematics and Statistics

[See the current edition of the Examination Regulations for the full regulations governing these examinations.]

Students staying on to take the four-year course will take 2 units from Part C in their fourth year, and will also offer a dissertation on a statistics project. Of the 2 units from Part C, at least half a unit will be from the schedule of 'Statistics' units for Part C.

This booklet describes the units and half-units available in Part C. Information about dissertations/ statistics projects is available on the Department of Statistics website at http://www.stats.ox.ac.uk/current_students/bammath/projects

We ask that you register by the end of week 9 Trinity Term 2009 for classes for the Mathematics/ Statistics courses that you wish to take. A registration form is attached to these synopses. Some combinations of subjects are not advised and lectures in these subjects may clash. However, when timetabling lectures we will aim to keep clashes to a minimum.

Language Classes: Mathematics and Statistics students are also invited to apply to take classes in a foreign language. In 2009-2010 classes are offered in French. Students' performance in these classes will not contribute to the degree classification in Mathematics and Statistics. However successful completion of the course may be recorded on student transcripts. See http://www.maths.ox.ac.uk/current_students/undergraduates/handbooks-synopses for further information.

Candidates leaving after four years who satisfy the Examiners will be awarded an MMath in Mathematics and Statistics with two associated classifications.

For the MMath in Mathematics and Statistics, one of the classifications will be based on Part C alone. Each candidate will receive a numerical mark on each paper in each Part of the examination in the University standardised range 0-100, such that

- a First Class performance (on that paper) is indicated by a mark of 70 to 100;
- an Upper Second Class performance (on that paper) is indicated by a mark of 60 to 69;
- a Lower Second Class performance (on that paper) is indicated by a mark of 50 to 59
- a Third Class performance (on that paper) is indicated by a mark of 40 to 49;
- a Pass performance (on that paper) is indicated by a mark of 30 to 39;
- a performance at the level of a Fail (on that paper) is indicated by a mark of 0 to 29.

1.1 Units and half-units and methods of examination

The Statistics lecture courses given are available as half-units apart from MS4b/C11.b *Graph Theory/ Probabilistic Combinatorics*. Half units are examined in an examination paper of 1 ½ hours or by mini-project.

All the units and half-units described in this booklet are “M-level”.

The Examination Papers in Part C

Each 16 hour lecture course is assessed as a half unit. On each half unit examination paper of 1 ½ hours, three questions will be set. Only the two best questions on that half unit paper count.

Most Mathematics papers including MS4b/C11.b *Graph Theory / Probabilistic Combinatorics* are whole unit papers, that is, they contain the questions for two half units. There are three questions set for each half unit. If you are taking a full unit, then you may attempt as many questions as you like but only your two best answers on each half unit will count towards your final mark for the paper.

Mini-project assignments can be collected from the Department of Statistics in week 8 of the relevant term and should be submitted to the Examination Schools by noon Monday week 1 of the following term.

For further details of examinations and assessment, please see the *Examination Conventions for Mathematics and Statistics*.

2 Statistics units and half units

2.1 MS1a: Graphical Models and Inference 16MT

Recommended Prerequisites

BS1 Applied statistics and BS2a Foundations of Statistical Inference would be helpful but not essential.

Aims & Objectives

Graphical models have become increasingly important in many areas where statistics play a role. They enable the description and analysis of complex stochastic systems via their natural modularity, expressed in terms of (mathematical) graphs which encode conditional independence structure. The modules correspond typically to well-understood, classical models. This course builds upon and develops the specific theory and computational tools needed in the analysis of graphical models for categorical and multivariate Gaussian data as well as Bayesian graphical models for complex stochastic systems.

Synopsis

Topics include:

1. Conditional independence and Markov properties.
2. Log-linear graphical models for categorical data.
3. Gaussian graphical models.
4. Graphical models for complex stochastic systems

Method of Examination

1 ½ hour written examination

Reading

1. D. Edwards, *Introduction to Graphical Models* (2nd ed.), Springer-Verlag, New York (2002).
2. S. L. Lauritzen, *Graphical Models*, Oxford University Press, Oxford (1996).
3. P. J. Green, N. L. Hjort and S. Richardson, eds. *Highly Structured Stochastic Systems*, Oxford University Press, Oxford (2003).

2.2 **MS1b: Statistical Data Mining** - 12HT plus 4 1-hour computer practical classes

Recommended Prerequisites

Part A Probability and Statistics. BS1 *Applied Statistics* would be an advantage.

Aims & Objectives

'Data mining' is now widely used to find interesting patterns in large databases, for example in insurance, in marketing and in many scientific fields. With large amounts of data we can search for quite subtle patterns. This course concentrates on the statistical tools used to identify patterns, and then to identify those which are interesting not just the result of chance associations.

Synopsis

Fundamentals of pattern recognition, machine learning and data mining.

Exploratory methods: principal components analysis, biplots, independent component analysis, multidimensional scaling.

Cluster Analysis: K-means, hierarchical methods, vector quantisation, self-organising maps.

Linear discriminant analysis, logistic discrimination, linear separation.

Feed-forward neural networks, Classification trees, ensemble methods, V-fold cross-validation.

Method of Assessment

This course is assessed by mini-project.

Reading

C. Bishop, *Neural Networks for Pattern Recognition*, Oxford UP (1995).

D. Hand, H. Mannila, P. Smyth, *Principles of Data Mining*, MIT Press (2001).

I. H. Witten and E. Franke, *Data Mining. Practical Machine Learning Tools and Techniques with Java Implementations*, Morgan Kaufmann (2000).

Further Reading

B. D. Ripley, *Pattern Recognition and Neural Networks*, Cambridge UP (1996).

2.3 MS2a: Bioinformatics and Computational Biology- 16MT

Recommended Prerequisites

None. In particular, no previous knowledge of Genetics will be necessary.

Aims & Objectives

Modern molecular biology generates large amounts of data, such as sequences, structures and expression data, that needs different forms of statistical analysis and modelling to be properly interpreted. The fields of Bioinformatics and Computational Biology have this as their subject matter and there is no sharp boundary between them. Bioinformatics has an applied flavour while Computational Biology is viewed as the study of the models, statistical methodology and algorithms needed to do bioinformatics analysis. This course aims to present core topics of these fields with an emphasis on modelling and computation.

Synopsis

Fundamental Data Structures in Biology: Sequences, Genes and RNA secondary structure.

Stochastic Models of Sequence and Genome Evolution including models of single nucleotide/amino acid/codon evolution.

Phylogenies: enumerating phylogenies, the probability of sequences related by a specified phylogeny, the minimal number of events needed to explain a data set (Parsimony).

Likelihood and algorithms (Markov Chain Monte Carlo) for inference based on the likelihood.

Software packages for sample-based inference.

Alignment Algorithms. Comparing 2 strings, an arbitrary number of strings, find segments of high similarity in 2 strings.

Common Patterns in a set of Sequences.

Method of Assessment

This course is assessed by mini-project.

Reading

C. Semple and M. Steel, *Phylogenetics*, Oxford University Press (2003).

Durbin et al., *Biological Sequence Analysis*, Cambridge University Press (1998).

T. Jiang et al., (editors) *Current Topics in Computational Biology*, MIT Press, (2003).

M. S. Waterman et al., *Computational Genome Analysis: An Introduction*, Springer (2004).

2.4 MS2b: Stochastic Models in Mathematical Genetics - 16HT

Aims & Objectives

The aim of the lectures is to introduce modern Stochastic models in Mathematical Population Genetics and give examples of real world applications of these models. Stochastic and Graph theoretic properties of coalescent and gene trees are studied in the first eight lectures. Extensions to model additional key biological phenomena, and applications, are studied in the second eight lectures.

Synopsis

Evolutionary models in Mathematical Genetics:

The Wright-Fisher model. The Genealogical Markov chain describing the number ancestors back in time of a collection of genes.

The Coalescent process describing the stochastic behaviour of the ancestral tree of a collection of genes. Mutations on ancestral lineages in a coalescent tree. Inferring the time to the most recent common ancestor in a sample of genes from the number of mutations occurring to the genes. Models with a variable population size.

The frequency spectrum and age of a mutation. Ewens' sampling formula for the probability distribution of the allele configuration of genes in a sample in the infinitely-many-alleles model. Hoppe's urn model for the infinitely-many-alleles model.

The infinitely-many-sites model of mutations on DNA sequences. Gene trees as perfect phylogenies describing the mutation history of a sample of DNA sequences. Graph theoretic constructions and characterizations of gene trees from DNA sequence variation. Gusfield's construction algorithm of a tree from DNA sequences. Examples of gene trees from data. The probability distribution of a gene tree.

Modelling biological forces in Population Genetics:

Recombination. The effect of recombination on genealogies. Detecting recombination events under the infinitely-many-sites model. Hudson's algorithm. Haplotype bounds on recombination events.

Modelling recombination in the Wright-Fisher model. The coalescent process with recombination: the ancestral recombination graph. Properties of the ancestral recombination graph. Applications of coalescent-based methods to the estimation of historical recombination rate.

Introduction to diffusion theory. Tracking mutations forward in time in the Wright-Fisher model. Modelling the frequency of a neutral mutation in the population via a diffusion process limit. The generator of a diffusion process with two allelic types. The probability of fixation and expected time to loss or fixation of a mutation. The frequency spectrum of a mutation.

Genic selection. Extension of results from neutral to selection case. Behaviour of selected mutations. Brief discussion of modern approaches to detecting selection from variation data.

Method of Assessment

1 ½ hour written examination

Reading

R. Durrett, *Probability Models for DNA Sequence Evolution*, Springer (2008).

W. J. Ewens, *Mathematical Population Genetics*, 2nd ed, Springer (2004).

J. R. Norris, *Markov Chains*, Cambridge University Press (1999).

M. Slatkin and M. Veuille, *Modern Developments in Theoretical Population Genetics*, Oxford Biology (2002).

S. Tavaré and O. Zeitouni, *Lectures on Probability Theory and Statistics, Ecole d'Eté de Probabilités de Saint-Flour XXXI - 2001*, Lecture Notes in Mathematics 1837.

Springer (2004).

2.5 MS3b: Lévy Processes and Finance - 16HT

Prerequisites

Part A Probability is a prerequisite. BS3a/OBS3a Applied Probability or B10 Martingales and Financial Mathematics would be useful, but are by no means essential; some material from these courses will be reviewed without proof.

Aims

Lévy processes form a central class of stochastic processes, contain both Brownian motion and the Poisson process, and are prototypes of Markov processes and semimartingales. Like Brownian motion, they are used in a multitude of applications ranging from biology and physics to insurance and finance. Like the Poisson process, they allow to model abrupt moves by jumps, which is an important feature for many applications. In the last ten years Lévy processes have seen a hugely increased attention as is reflected on the academic side by a number of excellent graduate texts and on the industrial side realising that they provide versatile stochastic models of financial markets. This continues to stimulate further research in both theoretical and applied directions. This course will give a solid introduction to some of the theory of Lévy processes as needed for financial and other applications.

Synopsis

Review of (compound) Poisson processes, Brownian motion (informal), Markov property. Connection with random walks, [Donsker's theorem], Poisson limit theorem. Spatial Poisson processes, construction of Lévy processes.

Special cases of increasing Lévy processes (subordinators) and processes with only positive jumps. Subordination. Examples and applications. Financial models driven by Lévy processes. Stochastic volatility. Level passage problems. Applications: option pricing, insurance ruin, dams.

Simulation: via increments, via simulation of jumps, via subordination. Applications: option pricing, branching processes.

Method of assessment

1 ½ hour written examination

Reading

J.F.C. Kingman: *Poisson processes*, Oxford University Press (1993), Ch.1-5, 8.
A.E. Kyprianou: *Introductory lectures on fluctuations of Lévy processes with Applications*, Springer (2006), Ch. 1-3, 8-9.
W. Schoutens: *Lévy processes in finance: pricing financial derivatives*, Wiley (2003).

Further reading

J. Bertoin: *Lévy processes*, Cambridge University Press (1996), Sect. 0.1-0.6, I.1, III.1-2, VII.1.
K. Sato: *Lévy processes and infinite divisibility*, Cambridge University Press (1999), Ch. 1-2, 4, 6, 9.

2.6 MS4b/C11.1b: **Probabilistic Combinatorics** - 16HT

[In the synopses booklet for Mathematics Part C, this course appears in the Mathematics Department Units section; in this booklet it is in the Statistics section. For any Mathematics and Statistics student taking this half-unit, it will count as a Statistics half-unit. Note that the prerequisite is C11.1a Graph Theory, which is available to Mathematics and Statistics students in Section 3 – C11.1a counts as a Mathematics half-unit for Mathematics and Statistics students.]

Prerequisites

C11.1a Graph Theory. Part A Probability.

Learning outcomes

To develop an appreciation of probabilistic methods in discrete mathematics.

Aims and objectives

Probabilistic combinatorics is a very active field of mathematics, with connections to other areas such as computer science and statistical physics. Probabilistic methods are essential for the study of random discrete structures and for the analysis of algorithms, but they can also provide a powerful and beautiful approach for answering deterministic questions. The aim of this course is to introduce some fundamental probabilistic tools and present a few applications.

Synopsis

Spaces of random graphs. Threshold functions.

First and second moment methods. Chernoff bounds. Applications to Ramsey numbers and random graphs.

Lovasz Local Lemma. Property B.

Poisson approximation, and application to the distribution of small subgraphs. Janson's inequality.

Concentration of measure. Martingales and the Azuma-Hoeffding inequality.

Chromatic number of random graphs.

Talagrand's inequality.

Method of Assessment

3 hour written examination also covering C11.1a Graph Theory

Reading

N. Alon and J.H. Spencer. *The Probabilistic Method*, Second edition, Wiley, 2000.

Further reading:

B. Bollobas, *Random Graphs*, second edition, CUP, 2001.

M. Habib, C. McDiarmid, J. Ramirez-Alfonsin, B. Reed, ed., *Probabilistic Methods for Algorithmic Discrete Mathematics* (Springer, 1998).

S.Janson, T. Luczak and A.Rucinski, *Random Graphs*, John Wiley and Sons, 2000.

M. Mitzenmacher and E. Upfal. *Probability and Computing : Randomized Algorithms and Probabilistic Analysis*, Cambridge University Press, New York (NY), 2005.

M. Molloy and B. Reed, *Graph Colouring and the Probabilistic Method* (Springer, 2002).

R. Motwani and P. Raghavan, *Randomized Algorithms* (CUP, 1995).

2.7 MS5b: High-throughput Data Analysis – 16 HT

Prerequisites: Part A (second-year) courses in Statistics and Probability.

Learning outcomes: Understanding the problems of working with high-throughput, high-frequency data-streams in applications such as finance, engineering and environmental monitoring. Understanding the limitations of established statistical techniques in dealing with such data. Development of working knowledge and experience with conventional and simulation-based methods for online statistical inference.

Understanding how to construct informative ‘data sketches’ from transiently observed data-streams. Experience in the use of such sketches.

Synopsis:

Illustrations of the problems of high-throughput, high-frequency data analysis in finance, internet monitoring and object-tracking. Defining the objective, namely the construction of efficient statistical algorithms that operate with limited storage and process with one pass over the data. Online data analysis. Introduction to dynamic statistical models. Gaussian and non-Gaussian models; Kalman filter, extended Kalman filter. Estimating volatility and cross-correlation in high-frequency financial time series. Problems of asynchronicity and scale.

Simulation based online filtering: concept of particle filters, design, optimisation, performance characteristics, connections with the Kalman filter. Applications to object tracking and parameter estimation in mathematical models of finance. Data-sketching. Probabilistic counting. Estimation of frequency moments by stable-law projection. Comparison of data streams via data sketches. Estimation of entropy and cross-entropy. Application to MCMC convergence diagnostics and monitoring of internet traffic.

Method of assessment

1 ½ hour written examination

Reading

- A. Doucet, N. De Freitas and N.J. Gordon, *Sequential Monte Carlo Methods in Practice*, Springer, 2001.
- A. Doucet and A.M. Johansen, *A tutorial on particle filtering and smoothing: fifteen years later*, Technical report, Department of Statistics, University of British Columbia, http://www.cs.ubc.ca/~arnaud/doucet_johansen_tutorialPF.pdf, 2008.
- J.P. Nolan, Information on Stable Distributions, <http://academic2.american.edu/~jpnolan/stable/stable.html>
- P. Indyk, Selected readings of his papers on data sketching: as indicated in the lectures.

Background reading: financial applications

P. Lequeux (Editor), *Financial Markets Tick by Tick*, Wiley, 1999.

2.8 MS6a: **The Analysis of Biological Networks** - 16MT

Recommended Prerequisites

None. In particular, no previous knowledge of Genetics will be necessary.

Aims & Objectives

Modern molecular biology generates large amounts of data, that allows the characterisation of biological systems. In the describing the dynamics and relationships of biological systems, network is the most central concept. At the cellular level there are 4 main classes: metabolic, regulatory, signalling and protein interaction networks. Such networks pose a set of common problems, such as inference, evolution, enumeration, comparison and dynamics. This course aims to present core techniques and problems of these topics with an emphasis on modelling and computation.

Synopsis

Dynamic High Throughput Data in Biology
Statistical analysis of Dynamic High Throughput Data
Biological Networks and their Temporal Dynamics
Evolution of Networks
Network Algorithms
Network Inference
Combinatorial Methods of Network Comparison
Probability Theory of Networks
Reconstruction Problems in Networks
Sampling Networks
Network Robustness

Method of Assessment

This course is assessed by mini-project.

Readings

Alon, U. (2006) *An Introduction to Systems Biology: Design Principles of Biological Circuits* CRC Press.

Davies, Rafnar, Hellenthal and Hein (2009) *Integrative Genomics and Functional Explanation*

Gillespie (2007) *Stochastic Simulation of Chemical Kinetics*, Ann. Rev. Phys. Chem.. 58:35–55

3 Mathematics units and half units

The Mathematics units and half units that students may take are drawn from Part C of the Honour School of Mathematics. For full details of these units and half-units, see the Syllabus and Synopses for Part C of the Honour School of Mathematics, which are available on the web at

<http://www.maths.ox.ac.uk/current-students/undergraduates/handbooks-synopses>

The Mathematics units and half-units that are available are as follows:

- C1.1 (or C1.1a or C1.1b) Gödel's Incompleteness Theorems and Model Theory
- C1.2 (or C1.2a or C1.2b) Analytic Topology and Axiomatic Set Theory
- C2.1 (or C2.1a or C2.1b) Lie Algebras and Representation Theory of Symmetric Groups
- C3.1 (or C3.1a or C3.1b) Lie Groups and Differentiable Manifolds
- C3.2a Topology and Groups
- C4.1 (or C4.1a or C4.1b) Functional Analysis, Banach and C* Algebras
- C5.1: (or C5.1a or C5.1b or C5.2) Partial Differential Equations for Pure and Applied Mathematicians and Calculus of Variables
- C6.1 Solid Mechanics
- C6.2 Elasticity and Plasticity
- C6.3 (or C6.3a or C6.3b) Perturbation Methods and Applied Complex Variables
- C6.4 Special Topics in Fluids
- C7.1b Quantum Theory and Quantum Computers
- C7.2b General Relativity I
- C8.1 (or C8.1a or C8.1b) Mathematics and the Environment and Mathematical Physiology
- C9.1 (or C9.1a or C9.1b) Analytic Number Theory and Elliptic Curves
- C10.1 (or C10.1a or C10.1b) Stochastic Differential Equations and Brownian Motion in Complex Analysis
- C11.1a Graph Theory
- C12.1 (or C12.1a or C12.1b) Numerical Linear Algebra and Continuous Optimisation
- C12.2 (or C12.2a or C12.2b) Approximation Theory and Finite Element Methods

4 Registration

We ask that students register in advance for the classes they wish to take, by the end of week 9 Trinity Term 2009, using the form overleaf.

Because of the large number of options which are available in Part C, some lectures will clash. See the Syllabus and Synopses for Part C of the Honour School of Mathematics for information on which lectures may clash.

FHS MATHEMATICS AND STATISTICS
REGISTRATION FORM: PART C CLASSES 2009-2010

SURNAMEFIRST NAME

EMAIL ADDRESS

COLLEGE

Note: As described in Section 1, you need to do a total of 2 units in Part C (in addition to doing a dissertation on a statistics project). At least half a unit will be from the schedule of 'Statistics' units for Part C

Please give details of the subjects in which you wish to take classes.
I wish to take classes in the following subjects: [Please Tick]

- MS1a Graphical Models and Inference
- MS1b Statistical Data Mining
- MS2a Bioinformatics and Computational Biology
- MS2b Stochastic Models in Mathematical Genetics
- MS3b Lévy Processes and Finance
- MS4b Probabilistic Combinatorics
- MS5b High-throughput Data Analysis
- MS6a The Analysis of Biological Networks

For Mathematics units or half-units, please list the unit or half-unit code and name:
Unit code Unit name

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Please return this form to the Academic Administrator, Department of Statistics,
1South Parks Road, by the end of week 9 Trinity Term 2009.