FINAL CALL for "Applied Bayesian modelling for ecologists and epidemiologists" (ABME03)

Delivered by Dr. Matt Denwood and Emma Howard

www.prstatistics.com/course/applied-bayesian-modelling-ecologists-epidemiologists-abme03/

This 6 day course will run from 20th - 25th November 2017 at SCENE field station, Loch Lomond national park, Scotland.

This application-driven course will provide a founding in the basic theory & practice of Bayesian statistics, with a focus on MCMC modelling for ecological & epidemiological problems.

Starting from a refresher on probability & likelihood, the course will take Starting from a refresher on probability & likelihood, the course will take students all the way to cutting-edge applications such as state-space population modelling & spatial poin-process modelling. By the end of the week, you should have a basic understanding of how common MCMC samplers work and how to program them, and have practical experience with the BUGS language for common ecological and epidemiological models. The experience gained will be astificient foundation enabling you to understand current papers using Bayesian methods, carry out simple Bayesian analyses on your own data and springboard into more elaborate applications such as dynamical, spatial and hierarchical modelling.

Course content is as follows

- Day 1

 Revision of likelihoods using full likelihood profiles and an
- Probability and likelihood
- 0

Probability and incentiood Conditional, joint and total probability, independence, Baye's law Probability distributions Uniform, Bernoulli, Binomial, Poisson, Gamma, Beta and Normal distributions - their range, parameters and common usesoLikelihood and

- distributions their range, parameters and common ussoLikelihood and parameter estimation by maximum likelihood o Numerical likelihood profiles and maximum likelihood Introduction to Bayesian statistics Relationship between prior, likelihood & posterior distributions o Summarising a posterior distribution; The philosophical differences

between frequentist & Bayesian statistics, & the practical implications of

- o Applying Bayes' theorem to discrete & continuous data for common data types given different priors o Building a posterior profile for a given dataset, & compare the effect of different priors for the same data

Day 2 An introduction to the workings of mcmc, and the potential dangers of mcmc inference. Participants will program their own (basic) mcmc sampler to illustrate the concepts and fully understand the strengths and weaknesses of the general approach. The day will end with an introduction

- weaknesses of the general approach. The day will end with an introduction to the bugs language. Introduction to MCMC. The curse of dimensionality & the advantages of MCMC sampling to determine a posterior distribution.
- Monte Carlo integration, standard error, & summarising posteriori distributions in R.
 Writing a Metropolis algorithm & generating a posterior distribution for a simple problem using MCMC.
 Markov chains, autocorrelation & convergence.
 Definition of a Markov chain.

- Definition of a Markov chain. Autocorrelation, effective sample size and Monte Carlo error. The concept of a stationary distribution and burning. Requirement for convergence diagnostics, and common statistics for ssing convergence. Adapting an existing Metropolis algorithm to use two chains, &

- Adapting an existing metropoins algorithm to use two enamis, &
 assessing the effect of the sampling distribution on the autocorrelation.
 Introduction to BUGS & running simple models in JAGS.
 Introduction to the BUGS language & how a BUGS model is translated
 to an MCMC sampler during compilation.
 The difference between deterministic & stochastic nodes, & the
- contribution of priors & the likelihood.

Running, extending & interpreting the output of simple JAGS models from within R using the runjags interface.

Day 3 • This day will focus on the common models for which jags/bugs would it was a provided a size of the formation of the size of the to this day with costs of the Common model in which age order you be be used in practice, with examples given for different types of model code. All aspects of writing, running, assessing and interpreting these models will be extensively discussed so that participants are able and confident to run similar models on their own. There will be a particularly heavy focus on practical sessions during this day. The day will finish with a discussion of how to assess the fit of mcmc models using the
 with a discussion of how to assess the ht of meme models using the deviance information criterion (dic) and other methods.

 o
 Using JAGS for common problems in biology.

 o
 Understanding and generating code for basic generalised linear mixed models in JAGS.

 o
 Syntax for quadratic terms and interaction terms in JAGS.

- Synax for quantum terms and mechanism terms in TAGS.
 Essential fitting tips and model selection
 The need for minimal cross-correlation and independence between
 parameters and how to design a model with these properties.
 The practical methods and implications of minimizing Monte Carlo

- The practical methods and implications of minimizing Monte Carl error and autocorrelation, including thiming.
 Interpreting the DIC for nested models, and understanding the limitations of how this is calculated.
 Other methods of model selection and where these might be more useful than DIC.
- Most commonly used methods Rationale and use for fixed threshold,
- ABGD, K/theta, PTP, GMYC with computer practicals. o Other methods, Haplowebs, bGMYC, etc. with computer practicals

Day 4 • 1

Day 4 will focus on the flexibility of mcmc, and precautions • Day 4 will focus on the flexibility of mcmc, and precatitons required for using mcmc to model commonly encountered datasets. An introduction to conjugate priors and the potential benefits of exploiting gibbs sampling will be given. More complex types of models such as hierarchical models, latent class models, mixture models and state space models will be introduced and discussed. The practical sessions will follow on from day 3.

General guidance for model specification. The flexibility of the BUGS language and MCMC methods.

- The difference between informative and diffuse priors
- 0
- Conjugate priors and how they can be used. Gibbs sampling. State space models. Hierarchical and state space models. 0
- Latent class and mixture models.
- Conceptual application to animal movement. Hands-on application to population biology. Conceptual application to epidemiology

- Day 5 Day 5 will give some additional practical guidance for the use of Bayesian methods in practice, and finish with a brief overview of more advanced Bayesian tools such as inla and stan. O Additional Bayesian methods. Understord the usefulness of conjugate priors for robust analysis

- of proportions (Binomial and Multinomial data)

- O Be aware of some methods noninal data):
 O Be aware of some methods noninal data):
 Advanced Bayesian tools.
 O Strengths and weaknesses of Integrated Nested Laplace Approximation
 (INLA) compared to BUGS.
- Strengths and weaknesses of Stan compared to BUGS

round table discussion and problem solving with final Q and A. o The final day will consist of round table discussions, the class

o The final day will consist or round table unscussions, the class will be split in to smaller groups to discuss set topic/synoblems. This will include participants own data where possible. After an early lunch there will be a general question and answer time until approx. 2pm as a whole group before transport to Balloch train station.

There will be a 15 minute morning coffee break, an hour for lunch, and a 15 minute afternoon coffee break. We keep the timing of these flexible depending how the course advances. Breakfast is from 08:00-08:45 and dinner is at 18:00 each day.

Please email any inquiries to oliverhooker@prstatistics.com or visit our website www.prstatistics.com

Please feel free to distribute this material anywhere you feel is suitable

Upcoming PR statistics courses

- 1. 1. ECOLOGICAL NICHE MODELLING USING R #ENMR 16th 20th October 2017, SCENE, Scotland, Dr. Nefali Sillero https://urldemes.proofpoint.com/v2url?u=http-3A__www.pratiatistics.com_course_ecological=2Dniche=2Dmodelling=2Dusing=2Dr=2D&d=DwlF=g&c=Ngd= ta5yR/sqeUsEDgshcqsYYY1Xs5ogLxWPA_2Wlc4&r=e2OJ1arRFn8ihlzb2HxzT0AqoiqLvxfeeaTyN59ZLo1&m=CirbnhlidSinlamxy09MqEsIL41_7Glxe77IPUWcRnA&s=P2z9RRqG87ZoaUMvQl9GPTx&PbTle1emc6C2kR5wJ&&= enmr01/

2. INTRODUCTION TO BIOINFORMATICS USING LINUX #IBUL IGth – 20th October, SCENE, Scotland, Dr. Martin Jones https://urldemess.gorofpoint.com/2vul?u=https://sceness.gorofpoint.com/2vul linux_ibul02/

REPRODUCIBLE DATA SCIENCE FOR POPULATION GENETICS #RDPG

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STRUCTURAL EQUATION MODELLING FOR ECOLOGISTS AND EVOLUTIONARY

BIOLOGISTS USING R #SEMR

DIVLOUID IS USING R NEMK 23rd - 2rth October 2017, Wales, Prof Jarrett Byrnes, Dr. Jon Lefcheck https://urldefense.proofpoint.com/v2/url?u=http=3A_www.prstatistics.com_course_structural-2Dequation-2Dmodelling-2Dfor-2D&d=DwIF-g&c=Ngd-ta5yRYsqU&EDgzhoqyYY1XS5ogLxWPA_2WIe4&r=e2OJ1azRFn8ihJzb2HxZT0AqoiqLvxfeeaTyN59ZLoI&m=CirbnhlkdStnIamxy09MqEsIL41_7Gixe77fPUWcRnA&s=8ZvJOWXEGCwjsXNFHEC_4-sD7u9pHhIrT7H110MOIH8&e= explorate active active intervisionize norm/U/ evolutionary-biologists-semr01

LANDSCAPE (POPULATION) GENETIC DATA ANALYSIS USING R #LNDG

 LANDSCAPE (POPULATION) OF NET DATA ANALYSIS USING K #LNDO 6th – 10th November 2017, Wales, Prof. Rodney Dyer https://url/defense.proofpoint.com/v2/url?u=http-3A_www.prstatistics.com_course_landscape-2Dgenetic-2Ddata-2Danalysis-2Dusing-2Dr-2D&d=DwIF-g&c=Ngd-ta5yRYsqeU&EDgxhcqsYYY1Xs5ogLxWPA_2W1c4&r=e2OJ1azRFn8ihJzb2HxZT0AqoiqLvxfeeaTyN59ZLol&m=CirbnhlkdSmlamxy09MqEsJL41_7Gixe77(PUWcRnA&s=c-vgcKomT2DSCyey51Ty2PQ8_q0p89MMs2lp7gyod1M&e= lndg02/

APPLIED BAYESIAN MODELLING FOR ECOLOGISTS AND EPIDEMIOLOGISTS #ABME

0. AFYLIED BATESIAN MODELLING FOR ECOLOGIES AND EFIDEMIOLOOIS S #ABME 20th - 25th November 2017, SCENE, Scotland, Dr. Matt Denwood https://rldefense.proofpoint.com/v2/urf?u=http-3A_www.prstatistics.com_course_applied-2Dbayesian-2Dmodelling-2Decologists-2D&d=DwIF-g&c=Ngd-ta5yRYsqeUsEDgxhcqsYYY1Xs5ogLxWPA_2WIe4&r=e2OJ1azRFn8ihJzb2HxZT0AqoiqLvxfeeaTyN59ZLoI&m=CirbnhlkdStnIamxy09MqEsJL41_7GIxe77fPUWcRnA&s=Vrg69GJ5auUtmI9hWMt4w6sCw8KYKZmML5oNGQGGmmE&e=

epidemiologists-abme03/

 INTRODUCTION TO PYTHON FOR BIOLOGISTS #IPYB
 27th Nov – 1 st Dec, Wales, Dr. Marini Jones
 https://urldefense.proofpoint.com/v2/url?u=http-3A__www.prinformatics.com_course_introduction-2Dto-2Dpython-2Dfor-2Dbiologists-2D&d=DwIF-g&c=Ngd utsyRV squUsEDgsthcqsYYY1KS50gLxWPA_2WIc4&r=e2OI larRFn8ihJzb2HxzT0AqoiqLxzfeeaTyNS9ZLoI&m=CirbnhlkdStnIamxy09MqE3LL41_7Gixe77/PUWcRnA&s=UhHMe7mdtmyusmy7wRiGhA9gadQaX-b98pHbbGi7E1U&c= ipyb04/

ADVANCING IN STATISTICAL MODELLING USING R #ADVR 4th - 8th December 2017, Wales, Dr. Luc Bussiere, Dr. Tom Houslay, Dr. Ane

Timenes Laugen.

https://url/defense.proofpoint.com/v2/url/u=http-3A_www.prstatistics.com_course_advancing-2Dstatistical-2Dmodelling-2Dusing-2Dr-2D&de=DwIF-g&c=Ngd-ta5yRYsqeUsEDgxhcqsYYY1Xs5ogLxWPA_2Wlc4&r=e2OJ1azRFn8ihJzb2HxZT0AqoiqLxxfeeaTyN59ZLoI&m=CirbnhlkdStnlamxy09MqEsIL41_7Glxe77fPUWcRnA&s=0XtuugMgd0IRKvWSqta81lijlK_tb8VnoJ5jonpnGRk&e=

INTRODUCTION TO BAYESIAN HIERARCHICAL MODELLING #IBHM

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10. PHYLOGENETIC DATA ANALYSIS USING R #PHYL

10. THEODEXCIPC DATA ANALISIS OSING KWITTE 208th Jan – Feb And 2018, SCENE, Scotland, Dr. Emmanuel Paradis https://urldefense.proofpoint.com/v2/url?u=https-3A_www.prstatistics.com_course_introduction-2Dto-2Dphylogenetic-2Danalysis-2D&d=DwIF-g&c=Ngd-ta5yRYsqeUsEDgxhcqsYYY1Xs5ogLxWPA_2WIc4&r=e2OJ1azRFn8ihJzb2HxZT0AqoiqLvxfeeaTyN59ZLo1&m=CirbnhlkdSmIamxy09MqEsJL41_7Glxe77fPUWcRnA&s=vOx90HQ2xSqxXM0cPU2QfFRPkIkaiSew513aTPXuZec&e=

with-r-phyg-phyl02/

11. MOVEMENT ECOLOGY #MOVE

John – 23rd February 2018, Wales, Dr Luca Borger, Dr Ronny Wilson, Dr Jonathan Potts

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GEOMETRIC MORPHOMETRICS USING R #GMMR
 19th – 23rd February 2018, Wales, Prof. Dean Adams, Prof. Michael Collyer, Dr. Antigoni Kaliontzopoulou

11 Junioran Automatic protocol https://url/demes.proof/point.com/v2/url/u=http-3A_www.prstatistics.com_course_geometric-2Dmorphometrics-2Dusing-2Dr-2Dgmmt01_&d=DwIF-g&c=Ngd-ta5yRYsqeUsEDgxhcqsYYY1Xs5ogLxWPA_2Wlc4&r=e2O11azRFn8ihJzb2HxZT0AqoiqLvxfeeaTyN59ZLol&m=CirbnhlkdStnlamxy09MqEsJL41_7Gixe77iPUWcRnA&s=hN6dFPEN5fcRWkcdscU-RDW_vXsEcEZakAL7hduBctU&e=

13. FUNCTIONAL ECOLOGY FROM ORGANISM TO ECOSYSTEM: THEORY AND COMPUTATION #FEER 5th – 9th March 2018, SCENE, Scotland, Dr. Francesco de Bello, Dr. Lars Götzenberger, Dr. Carlos Carmona Volzenerger, Jr. Canto Cantona https://urldenes.porofonit.com/2/url/u=http-3A_www.prstatistics.com_course_functional-2Decology-2Dfrom-2Dorganism-2Dto-2D&d=DwIF-g&c=Ngd-ta5yRYsqeUsEDgxhcqsYYY1Xs5ogLxWPA_2Wlc4&r=e2OJ1azRFn8ihJzb2HxZT0AqoiqLvxfeeaTyN59ZLoI&m=CirbnhlkdStnIamxy09MqEsJL41_7Gixe77fPUWcRnA&s=hPTVbmw_5kefEaQnedkgfQ3Y7TTO6jW7dEpqIwOfCjI&e= ecosystem-theory-and-computation-feer01/ 14. SPATIAL PRIORITIZATION USING MARXAN #MRXN 5th - 9th March 2018, Wales, Jennifer McGo

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15. ECOLOGICAL NICHE MODELLING USING R #ENMR

12. ECOLOGICAE INCLE arGUERATIO CONTROL MALENTE 12th - 16th March 2018, SCENE, Scotland, Dr. Neftali Sillero https://urldefense.proofpoint.com/v2/url?u=http-3A_www.prstatistics.com_course_ecological-2Dniche-2Dmodelling-2Dusing-2Dr-2D&d=DwIF-g&c=Ngd-ta5yRYsqcUsEDgxhcqsYYY1Xs5ogLxWPA_2Wlc4&r=e2OJ1azRFn8ihJzb2HxZT0AqoiqLvxfeeaTyN59ZLo1&m=CirbnhlkdStn1amxy09MqE3JL41_7Gixe77fPUWcRnA&s=P2z9RrqG87ZoaUMvQl9GPTx8-PbTle1emc6C2kR5wJ8&e=

16. BEHAVIOURAL DATA ANALYSIS USING MAXIMUM LIKLIHOOD IN R #BDML 19th – 23rd March 2018, Scotland, Dr William Hoppitt COMING SOON www.PSstatistics.com

NETWORK ANAYLSIS FOR ECOLOGISTS USING R #NTWA 17.

9th - 13th April 2018, SCENE, Scotland, Dr. Marco Scotti

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INTRODUCTION TO STATISTICAL MODELLING FOR PSYCHOLOGISTS USING R 18. #IPSY

16th – 20th April 2018, SCENE, Scotland, Dr. Dale Barr, Dr Luc Bussierre COMING SOON www.PSstatistics.com

MULTIVARIATE ANALYSIS OF ECOLOGICAL COMMUNITIES USING THE VEGAN

PACKAGE #VGNR 23rd - 27th April 2018, SCENE, Scotland, Dr. Peter Solymos, Dr. Guillaume

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20. QUANTITATIVE GEOGRAPHIC ECOLOGY: MODELING GENOMES, NICHES, AND

COMMUNITIES #QGER 30th April – 4th May 2018, SCENE, Scotland, Dr. Dan Warren, Dr. Matt Fitzpatrick

COMING SOON www.PRstatistics.com

21. INTRODUCTION TO MIXED MODELS FOR ECOLOGISTS #IMMR 14th - 18th May 2018, CANADA (QUEBEC) STILL to be confirmed, Prof Subhash Lele, Dr. Guillaume Blanchet

22 STABLE ISOTOPE MIXING MODELS USING SIAR, SIBER AND MIXSIAR #SIMM 28th May – 1st June 2018, CANADA (QUEBEC) STILL to be confirmed Dr. Andrew Parnell, Dr. Andrew Jackson

23. SOCIAL NETWORK ANALYSIS FOR BEHAVIOURAL SCIENTISTS USING R #SNAR 2nd - 5th12th July 2018, Prof James Curley

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24. MODEL BASE MULTIVARIATE ANALYSIS OF ABUNDANCE DATA USING K #MBM v 8th – 12th July 2018, Prof David Warton https://url/defense.proofpoint.com/v2/url?u=https=3A_www.prstatistics.com_course_model-2Dbase-2Dmultivariate-2Danalysis-2Dof-2D&d=DwIF-g&c=Ngd-tstyRvsgel26EDgxhcqsYYY1Ks5gLxWPA_2Wlc4&r=e2OJ1azRFn8ihJzb2HxZT0AqoiqLvxfeeaTyN59ZLoI&m=CirbnhlkdStnlamxy09MqEsJL41_7Glxe77fPUWcRnA&s=oNd86BgXm77uzLQfIZqAvgn_BTMUEwb2gBZbCSztZB4&e= abundance-data-using-r-mbmv02/

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26. ADVANCES IN MULTIVARIATE ANALYSIS OF SPATIAL ECOLOGICAL DATA USING R#MVSP

Prof. Pierre Legendre, Dr. Olivier Gauthier - Date

Oliver Hooker PhD. PR statistics

2017 publications -

Ecosystem size predicts eco-morphological variability in post-glacial diversification. Ecology and Evolution. In press.

The physiological costs of prey switching reinforce foraging specialization. Journal of animal ecology.

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