

Day 2: Population growth, species interactions and time series

Day 2: Population growth, species interactions and time series

Caveat:

You are pioneers... because this is version 1.0 of the course
We don't know what you know.

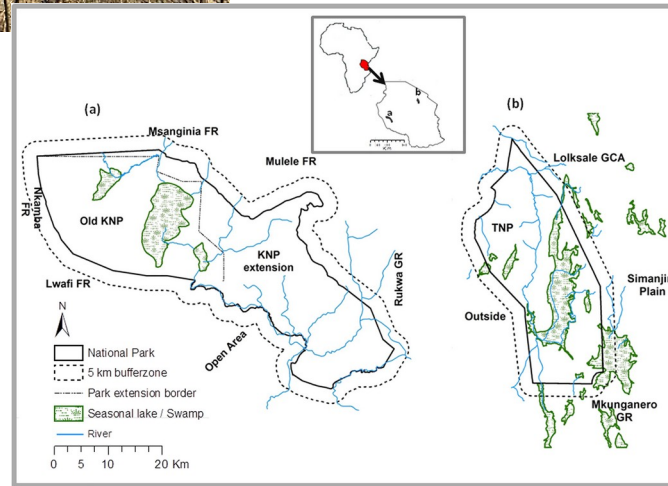
Please give us feedback during and after the course.

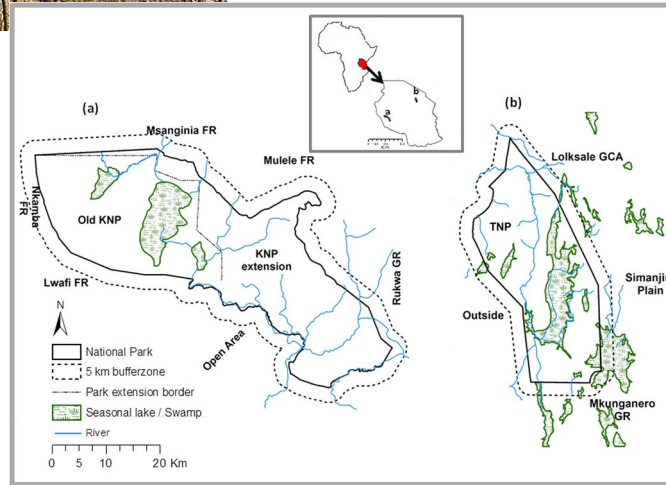
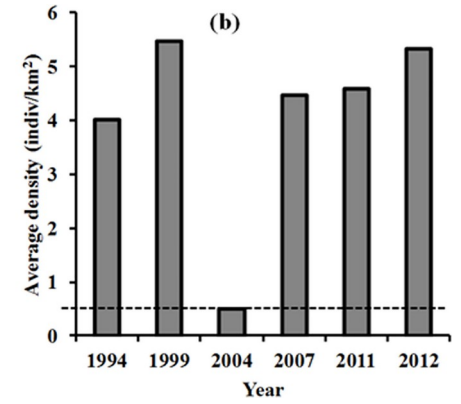
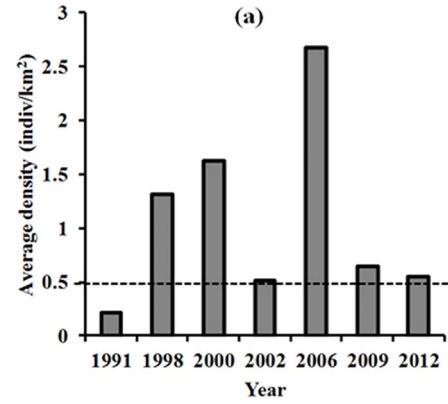
Day 2: Population growth, species interactions and time series

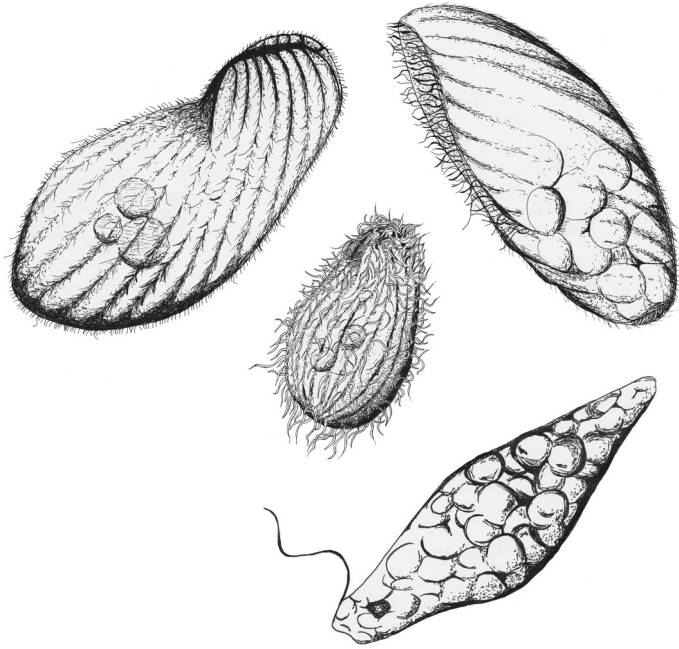
Schedule:

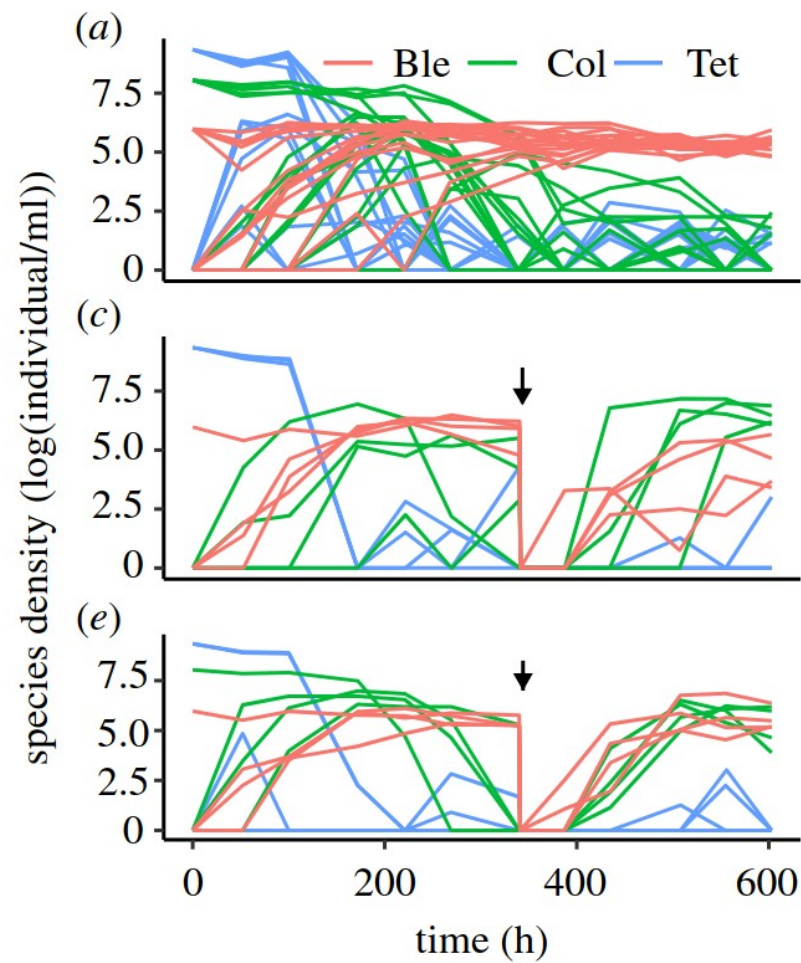
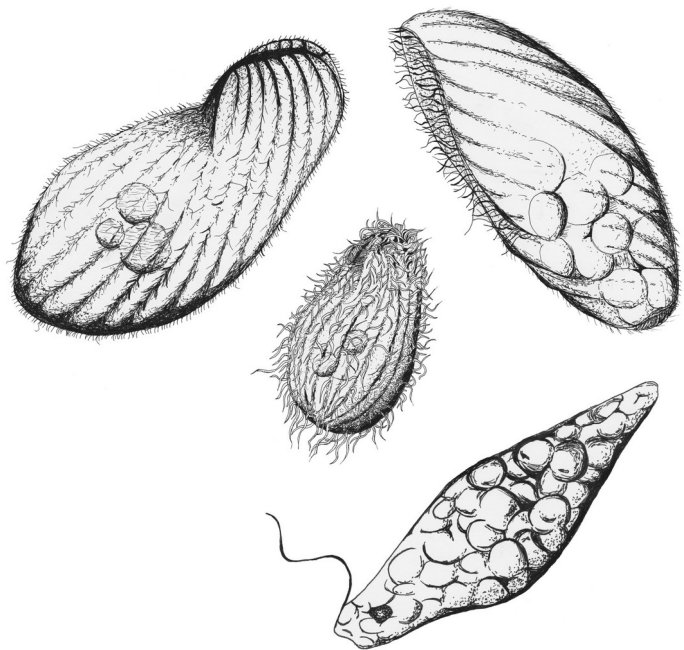
09:00 – 10:00	Introduction (Emanuel Fronhofer)
10:15 – 11:15	Fitting population dynamics in stan/rstan (Camille Saade)
11:30 – 12:30	Introducing process error and state-space models (Benjamin Rosenbaum)
14:00 – 17:00	It's your turn!
17:00 – 17:30	Fitting multiple time series at once (Camille Saade)
18:00 – 19:00	Evening lecture: Frédéric Barraquand

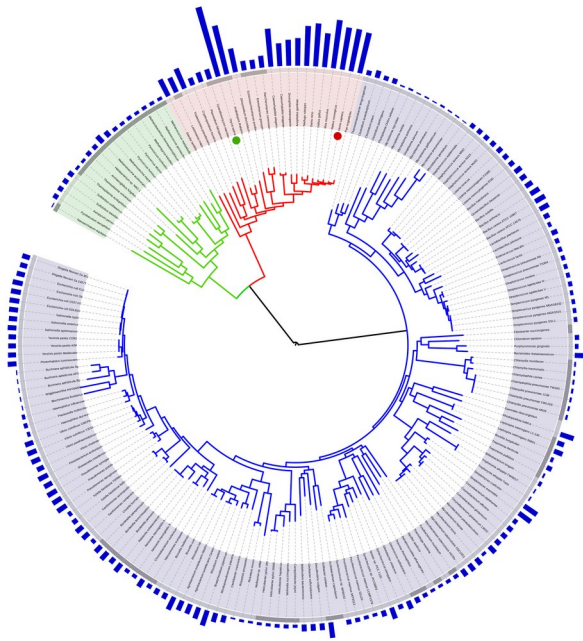


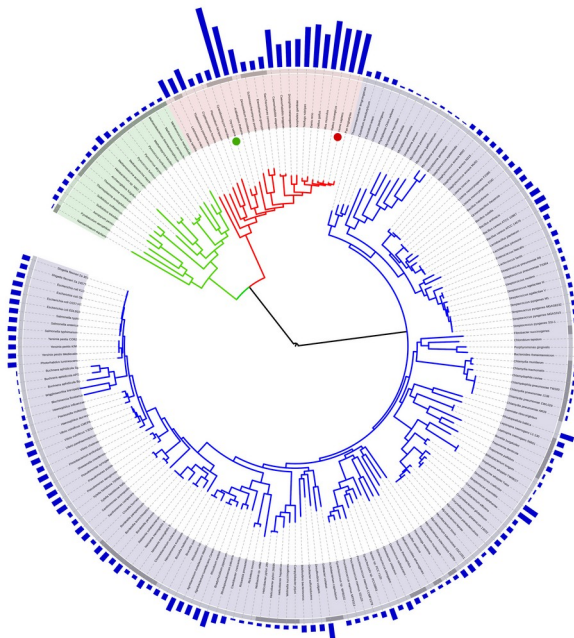




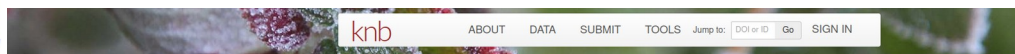








Global Population Dynamics Database



White, Owen Smith, John Lawton, Pablo Inchausti, et al. 2010. The Global Population Dynamics Database. Knowledge Network for Biocomplexity. doi:10.5063/F1B26328



Citations 0 Views 2.8K

Copy Citation Assessment report

Files in this dataset Package urn:uuid:bced9d4-1398-4788-a9f3-62007446a04

	File type	Size		Download All
Global Population Dynamics Database	EML v2.1.1	48 KB	2763 views	Download
	More info	text/csv	8 MB	2433 downloads
	More info	text/csv	794 KB	1710 downloads
	More info	text/csv	17 KB	1532 downloads

Show 5 more items in this data set

General

Identifier d35b:240.11

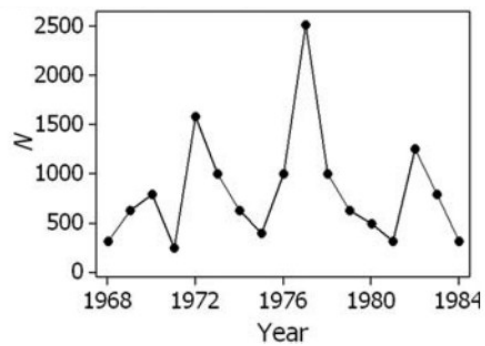
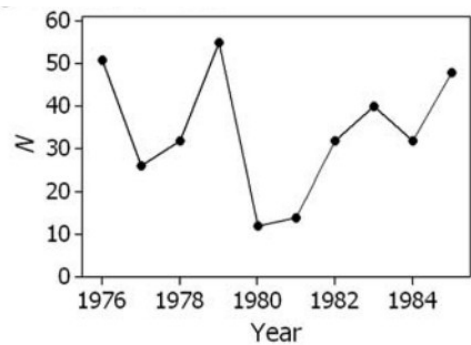
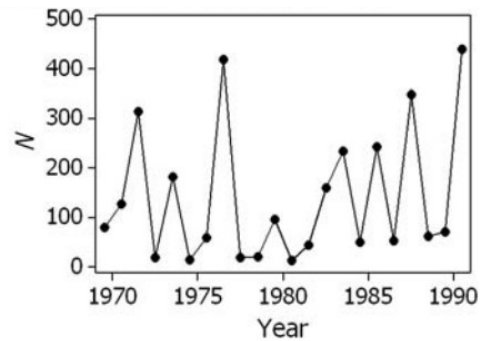
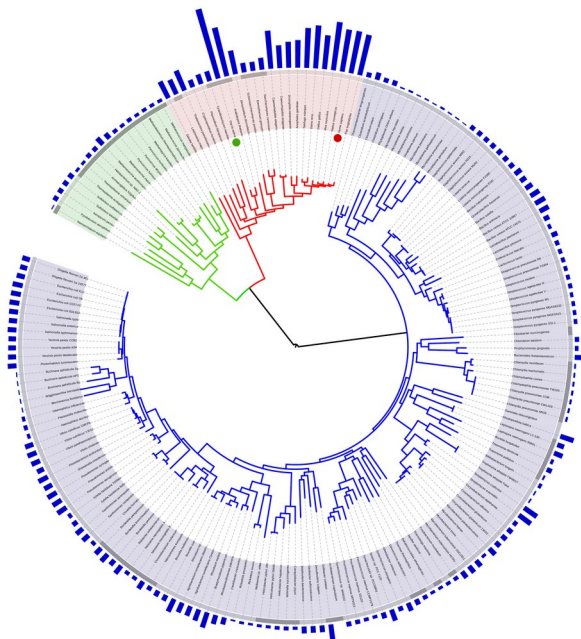
Abstract As a source of animal and plant population data, the Global Population Dynamics Database (GPDD) is unrivalled. Nearly five thousand separate time series are available here. In addition to all the population counts, there are taxonomic details of over 1400 species. The type of data contained in the GPDD varies enormously, from annual counts of mammals or birds at individual sampling sites, to weekly counts of zooplankton and other marine fauna. The project commenced in October 1994, following discussions on ways in which the collaborating partners could make a practical and enduring contribution to research into population dynamics. A small team was assembled and, with assistance and advice from numerous interested parties we decided to construct the database using the popular Microsoft Access platform. After an initial design phase, the major task has been that of locating, extracting, entering and validating the data in all the various tables. Now, nearly 5000 individual datasets have been entered onto the GPDD. The Global Population Dynamics Database comprises six Tables of data and information. The tables are linked to each other as shown in the diagram shown in figure 3 of the GPDD User Guide (GPDD-User-Guide.pdf). Referential integrity is maintained through record ID numbers which are held, along with other information in the Main Table. Its structure obeys all the rules of a standard relational database.

Keywords

Keyword

Time series data
animal populations
plant populations
count data

Type

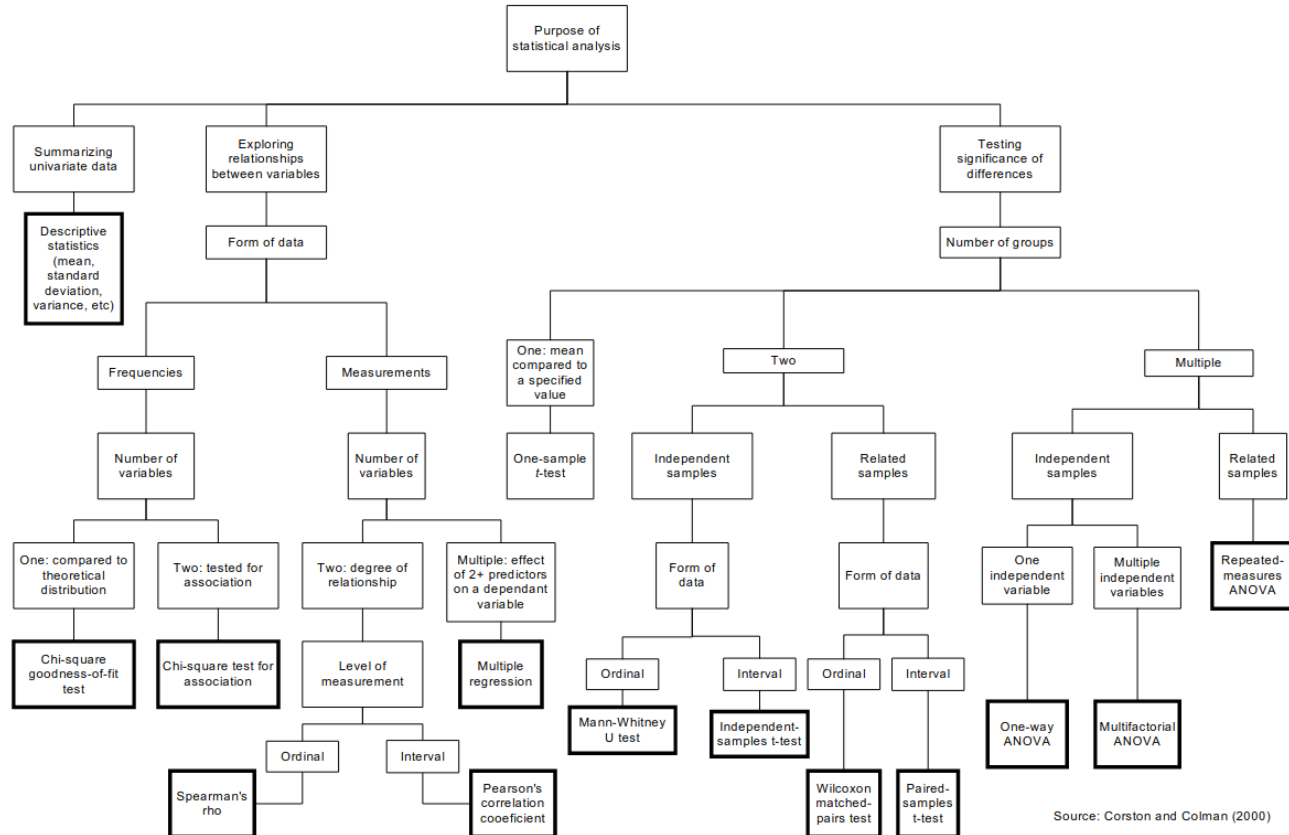


How should we analyse such data?

How should we analyse such data?

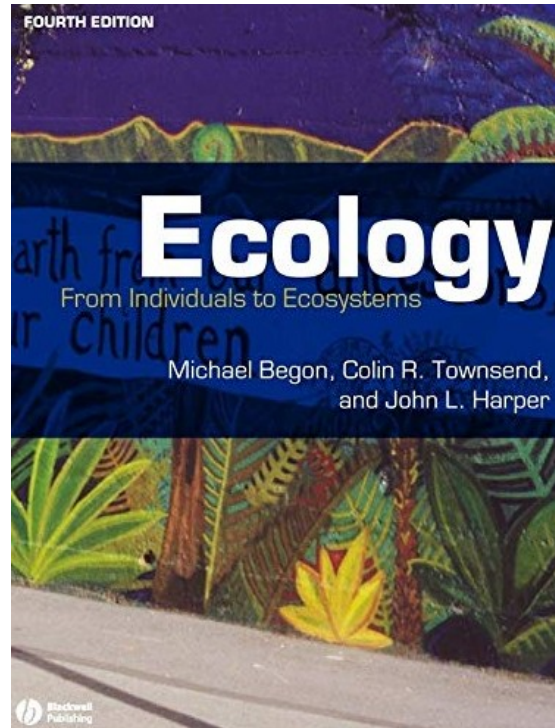
ANOVA
LM ? GAM
NLMM
(G)LMM

How should we analyse such data?

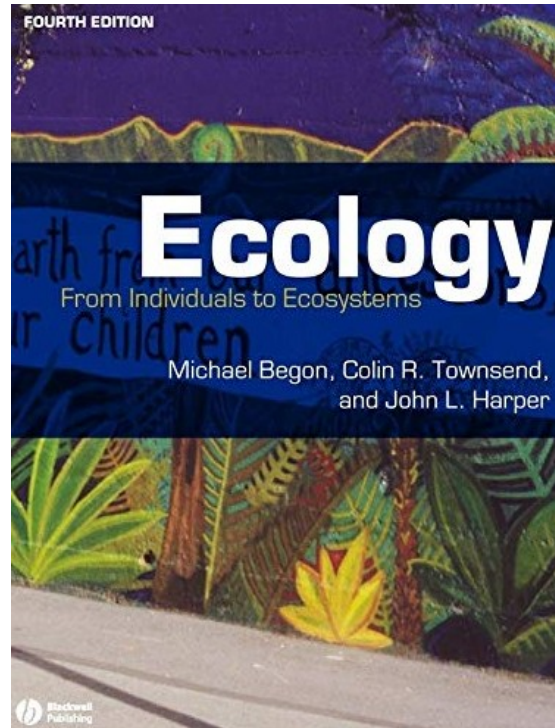


Source: Corston and Colman (2000)

How should we analyse such data?

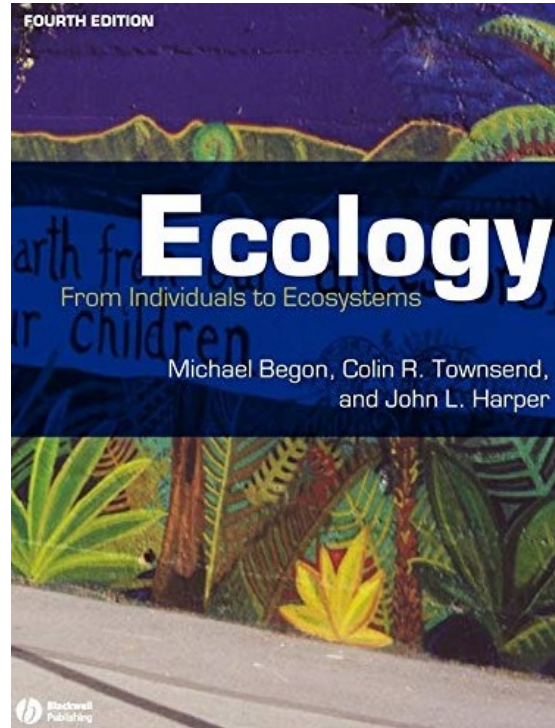


How should we analyse such data?



Why is this more interesting
than less informed models?

How should we analyse such data?



Why is this more interesting than less informed models?

- parameter estimation
- test of theory predictions
- maybe wrong inference?

Back to basics: exponential growth, logistic growth and inter-specific competition

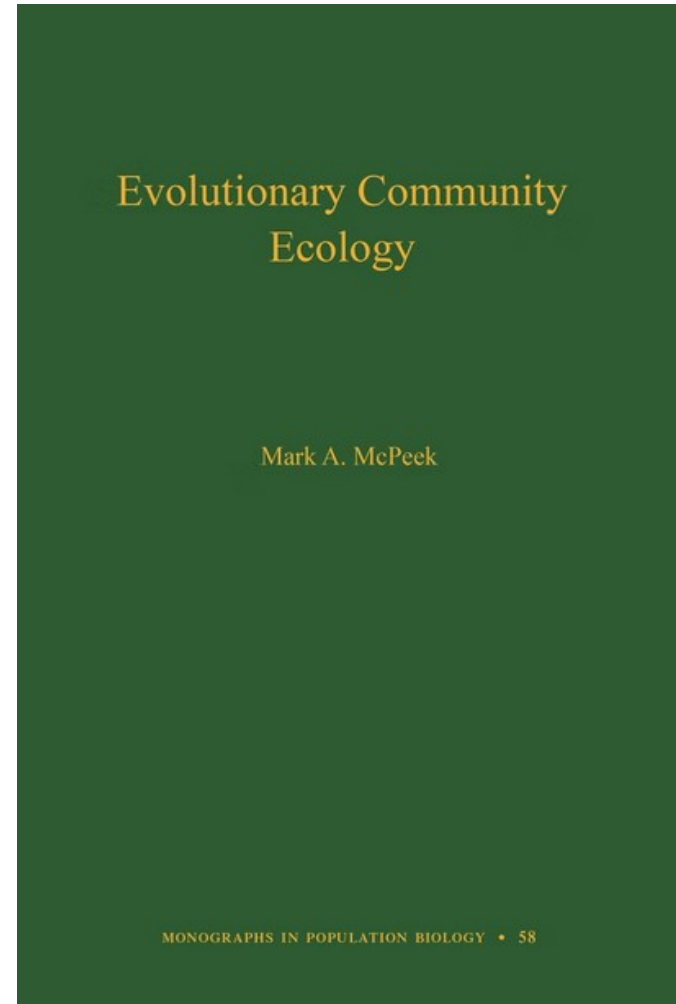
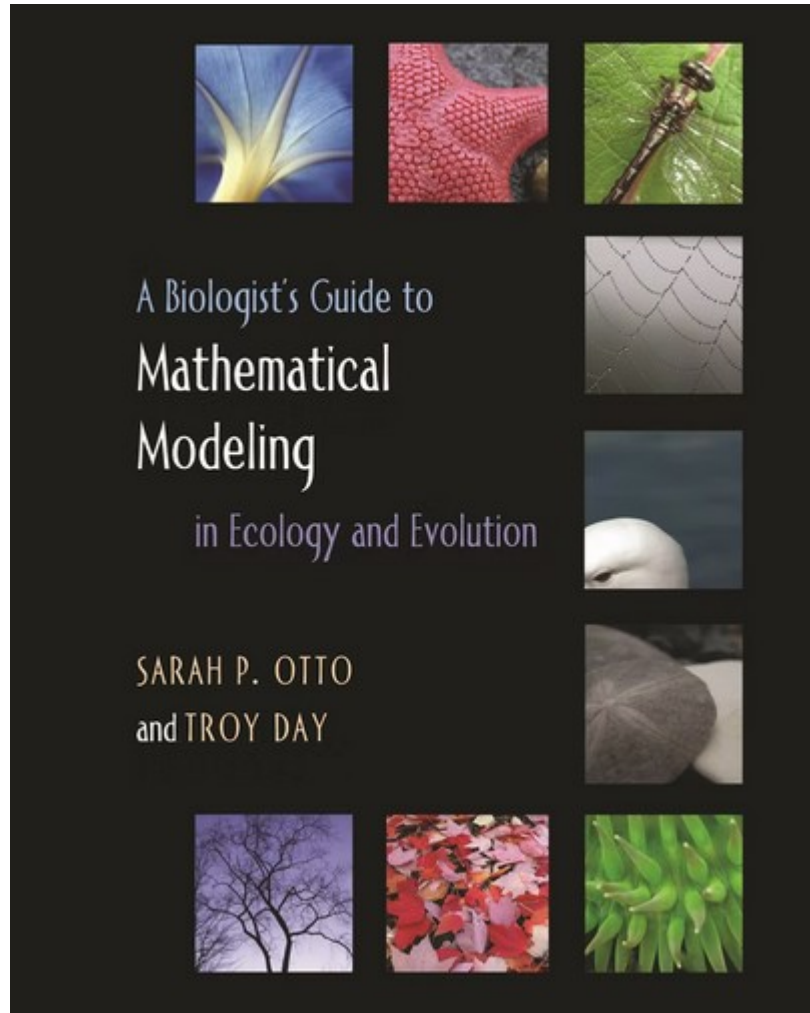
Evolutionary Ecology Research, 2012, **14**: 627–665

**The struggle for existence: how the notion of
carrying capacity, K , obscures the links
between demography, Darwinian evolution,
and speciation**

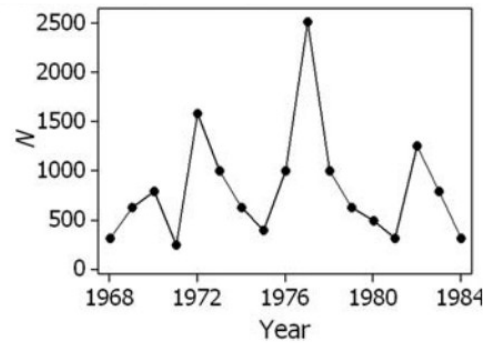
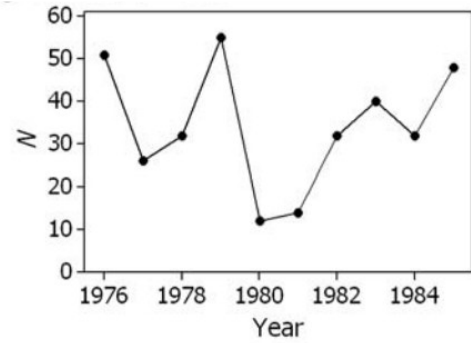
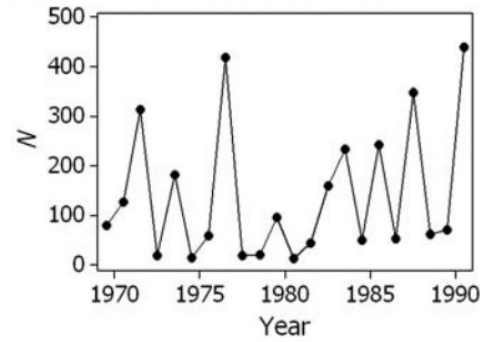
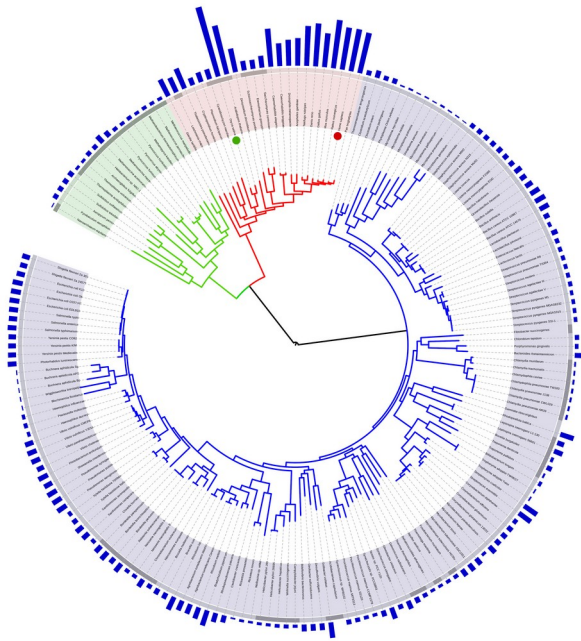
James Mallet

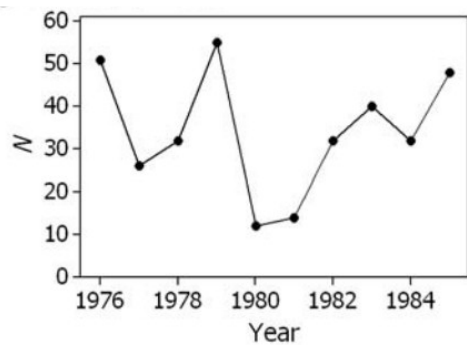
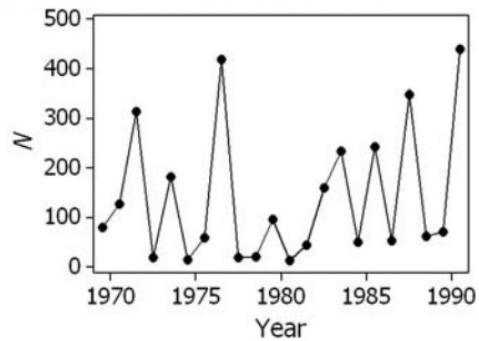
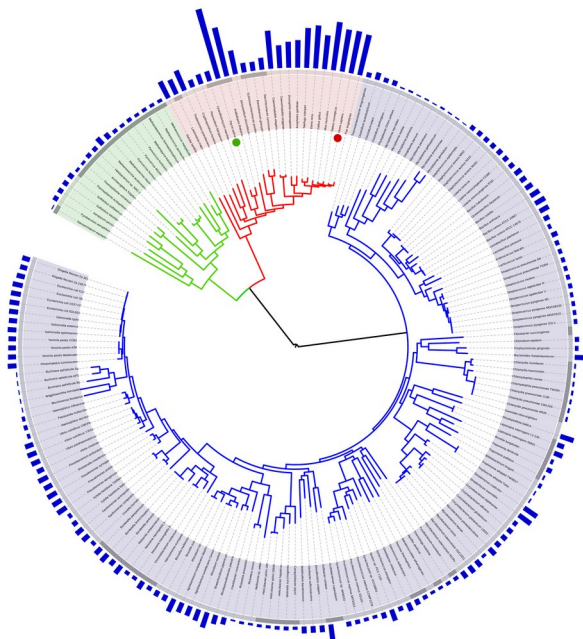
*Department of Organismic and Evolutionary Biology, Harvard University, Cambridge,
Massachusetts, USA and Department of Genetics, Evolution and Environment,
University College London, London, UK*

Back to basics: exponential growth, logistic growth and inter-specific competition



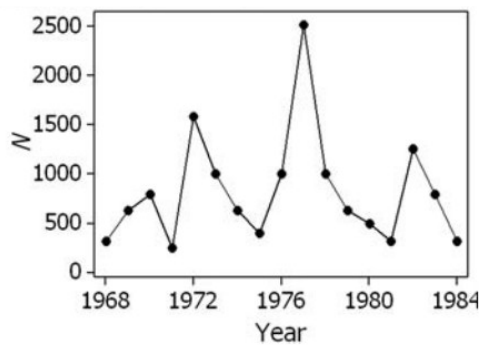
How should we fit such models?





$$\frac{dN}{dt} = r_0 \left[1 - \left(\frac{N}{K} \right)^\theta \right] N$$

least-squares



The theta-logistic is unreliable for modelling most census data

Francis Clark¹, Barry W. Brook¹, Steven Delean¹, H. Reşit Akçakaya² and
Corey J. A. Bradshaw^{1,3*}

¹*The Environment Institute and School of Earth & Environmental Sciences, University of Adelaide, Adelaide, SA 5005, Australia;* ²*Department of Ecology and Evolution, Stony Brook University, Stony Brook, NY 11794, USA; and*

³*South Australian Research and Development Institute, P.O. Box 120, Henley Beach, SA 5022, Australia*

The theta-logistic is unreliable for modelling most census data

Francis Clark¹, Barry W. Brook¹, Steven Delean¹, H. Reşit Akçakaya² and Corey J. A. Bradshaw^{1,3*}

¹The Environment Institute and School of Earth & Environmental Sciences, University of Adelaide, Adelaide, SA 5005, Australia; ²Department of Ecology and Evolution, Stony Brook University, Stony Brook, NY 11794, USA; and ³South Australian Research and Development Institute, P.O. Box 120, Henley Beach, SA 5022, Australia

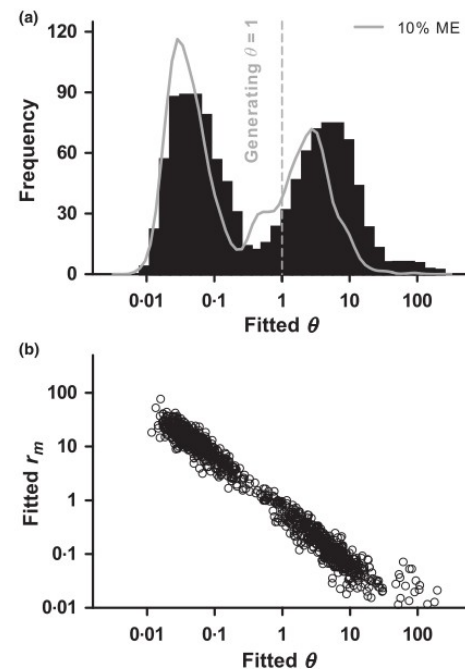


Fig. 2. Bimodality of fitted θ and relationship with r_m . (a) Fitted θ for 1000 simulations of 20 time-steps length, with generating model parameters: $r_m = 0.5$, $\theta = 1.0$, $\sigma = 0.1$ and $K = 100$ (vertical line at generating $\theta = 1.0$). Trace shows the effect of 10% measurement error (ME) applied to the abundance values; (b) \log_{10} – \log_{10} scatter plot of fitted $[r_m, \theta]$ pairs.

Richard McElreath

Anthropology, Evolutionary Ecology, Bayesian Data Analysis

HOME PUBLICATIONS SOFTWARE SR2 FOR INSTRUCTORS **STATISTICAL RETHINKING** TALKS TEACHING

Statistical Rethinking

FOLLOW ME ON TWITTER

[My Tweets](#)

A Bayesian Course with Examples in R and Stan (& PyMC3 & brms & Julia too, see links below)

Most recent set of free lectures: [Statistical Rethinking 2022](#)

Second Edition

The second edition is now out in print. Publisher information on the [CRC Press page](#). For more detail about what is new, [look here](#).

Materials

2nd Edition

- Book: [CRC Press](#)
- Book sample: [Chapters 1 and 2](#) (2MB PDF)
- Lectures and slides:
 - * Winter 2022 [materials](#) (ongoing)
 - * Winter 2019 [materials](#)
- Code and examples:
 - * R package: [rethinking](#) (github repository)
 - * R code examples from the book: [code.txt](#)
 - * Book examples in [Stan+tidyverse](#)
 - * brms + tidyverse conversion [here](#)
 - * PyMC3 code examples: [PyMC repository](#)
 - * [NumPyro!](#)
 - * More [NumPyro](#)
 - * TensorFlow Probability [notebooks](#)
 - * [Julia & Turing](#) examples (both 1st and 2nd edition)
 - * Another [Julia code translation](#) with clean outline in notebook format
 - * [R-INLA](#) examples

