Tim CD Lucas

PERSONAL INFORMATION

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PRESENT APPOINTMENT

2016-present **CBER Programmer**

Research Programmer I am currently the stuff programmer for the Centre of Biodiversity and Environment Research at UCL. I work on two main projects. I am working with the Madingley Model—an ecological model of all life, written in C#. My primary task is to enable this model to run on the *high performance cluster* at UCL. Secondly, I am translating code from *Mathematica to R* that performs statistical analyses for measurements of 3D objects used in paleontological research. I also provide technical support for the rest of the department.

EDUCATION

2012-present University College London, CoMPLEX

PhD Social structure and network epidemiology in bat zoonoses

> Description: I am using complex networks to study the epidemiology of bat-borne diseases. As bats carry a number of important zoonotic diseases, understanding the spread of these diseases within the bat population and how this affects spillover to humans and livestock is increasingly important. The unusually social nature of bat populations will strongly affect how diseases spread.

Supervisors: Prof. Kate Jones & Dr Hilde Wilkinson-Herbots

University College London, CoMPLEX 2011-2012

MRes Modelling Biological Complexity · Merit

> Description: Projects included adapting 'ideal gas' models to acoustic data, analysing moment closures for a pair-approximation model of plant ecology and

applying a novel *machine learning* method to a library of bat calls.

2006-2010 University of Sheffield, Animal & Plant Sciences

MBioSci Zoology · First

Description: For my final project I used wavelet analysis to study multi-annual

cycles in malaria incidence in Thailand.

PREVIOUS APPOINTMENTS

Autumn 2014 Zoön: An R package for reproducible SDMs

Internship I wrote the first version of an R package for reproducible species distribution modelling. The package uses an online repository of user submitted 'modules' to allow the software to keep up with this fast moving field and allow analyses

to be completely reproducible. [Github]

August 2011 Smithsonian Tropical Research Institute

Volunteer Two months fieldwork in Panamá on two projects: studying Anolis dewlap Fieldwork

evolution and studying gut length plasticity in Red-eyed tree frogs.

May 2011 Chiloé Silvestre, Chilé

Volunteer I spent two weeks trapping Darwin's foxes in Chilé to collect samples for **Fieldwork** geophylogenetics.

> August 2010 University of Sheffield

Summer Internship I studied the evolutionary response of plant communities to climate change with Dr Raj Whitlock. I collected, propagated and analysed plants collected

from the field.

August 2009 University of York, YCCSA

TRANSIT Internship I studied collective foraging behaviour by programming a complex 3D foraging model in Java and running simulations on a cluster at the York Centre for

Complex Systems Analysis.

OTHER APPOINTMENTS AND AFFILIATIONS

Peer Review Journals Reviewed for:

· Methods in Ecology and Evolution, National Academy Science Letters

Society Membership

2014-present · Royal Society of Tropical Medicine and Hygiene: Student member

CONFERENCES

The Zoön Project: Reproducible, Remixable and Shareable Species Distribution 2015 Modelling with R.

> Presentation at BES Annual Meeting by August T, Golding N, Lucas TCD, Gavaghan D, Isaac N, O'Hara B, van Loon E & McInerny G

Simple, Shareable and Reproducible Species Distribution Modelling with the Zoön R package.

Poster at BES Annual Meeting by N Golding, Lucas TCD, August T, Gavaghan D, Isaac N, O'Hara B, van Loon E & McInerny G

Comparative and computational studies of pathogen richness in bats.

Presentation at Research in Progress, RSTMH by Lucas TCD, Wilkinson-Herbots H & Jones KE.

A comparative and computational study of population structure and pathogen richness

Presentation at Epidemics5 conference by Lucas TCD, Wilkinson-Herbots H & Jones

An ideal gas model for estimating absolute abundances from bat detector data. Presentation at the National Bat Conference. [slides]

Pathogen diversity and bat population structure. Poster at British Parasitological Society Autumn Meeting.

Estimating abundance from camera traps and acoustic sensors.

Presentation at CEH, Wallingford seminar series.

Presentation at id2oxford conference. [slides] 2014

Poster at the CoMPLEX conference. [pdf]

Presentation at BritBats 2 [slides]. 2013

Invited attendance at ecoVIZ Tansley workshop.

Poster at the CoMPLEX conference and id2 conference. [pdf]

PUBLICATIONS

2016 Lucas TCD, Wilkinson-Herbots HM, & Jones KE. A comparative and

computational study of population structure and pathogen richness in bats. In prep.

Lucas TCD*, Moorcroft EA*, Freeman R, Rowcliffe MJ & Jones KE. (2015) A 2015 generalised random encounter model for estimating animal density with remote sensor data. Methods in Ecology and Evolution. doi: 10.1111/2041-210X.12346 [pdf]

Walters CL, Collen A, Lucas TCD, Mroz K, Sayer CA and Jones KE. (2013) 2013 Challenges of Using Bioacoustics to Globally Monitor Bats. in Bat Evolution, Ecology, and Conservation. Springer New York. 479-499.

* Co-first authors.

SOFTWARE

On CRAN

Lucas TCD, Golding N, August T, McInerny G, van Loon E (2015) Zoön: Reproducible, Accessible & Shareable Species Distribution Modelling. www.github.com/zoonproject/zoon

Lucas TCD (2015) palettetown: Use Pokemon Inspired Colour Palettes www.github.com/timcdlucas/palettetown

TEACHING

2015 · Demonstrator for reproducible species distribution modelling workshop run by Quantitative Ecology special interest group at BES.

2013–2014 · Online tutor for SysMIC, a course for teaching quantitative skills to biologists.

COMPUTATIONAL SKILLS

Languages R (eight years), Python, Matlab, Mathematica, Java, SQL.

OS Comfortable with Windows, Mac or Linux.

Other Experience in Git/Github, unit testing, LaTeX, web design, markdown, R package development, shell/ssh and high performance computing.

REFEREES

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