Thomas Guillerme

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ORCID: 0000-0003-4325-1275 — ResearcherID: G-9833-2014

RESEARCH INTERESTS

I'm interested in evolutionary biology at a large scale: the stories behind the history of life and the methodologies with which it is investigated. By combining classical methods such as descriptive palaeontology with modern bioinformatics tools and mathematical modeling, I work on understanding how to use morphology in evolutionary biology: from phylogenetic theory to macroevolution and macroecology.

ACADEMIC CAREER

2017 - Present: Postdoctoral Research Fellow in Vera Weisbecker's group

School of Biological Sciences

University of Queensland, St Lucia, Australia Research Associate in Martin Brazeau's group

Faculty of Natural Sciences, Department of Life Sciences Silwood Park Campus, Imperial College London, UK

EDUCATION

2016 - 2017:

2012 - 2015: PhD in Zoology, Trinity College Dublin

Macroevolution with living and fossil species

Awarded as it stands.

http://dx.doi.org/10.6084/m9.figshare.1583337

2010 - 2012: MSc in Palaeontology (honours), Université de Montpellier

2nd year project: Molecular clocks and evolutionary rates among primates: a palaeontological and genomic approach.

1st year project: Southern France Upper Cretaceous Theropod

teeth variability.

2007 - 2010: BSc in Biology, Université de Montpellier

PUBLICATIONS

Peer reviewed - Google Scholar profile: Thomas Guillerme

2018: **Guillerme** T. dispRity: a modular R package for measuring disparity. *Methods in Ecology and Evolution*, DOI: 10.1111/2041-210X.13022

2018: **Guillerme T** and Cooper N. Time for a rethink: sub-sampling methods in disparity-through-time analysis. *Palaeontology*, DOI: 10.1111/pala.12364

2017: Healy K, **Guillerme T**, Kelly SBA, Inger R, Bearhop S, Jackson AL. SIDER: an R package for predicting trophic discrimination factors of consumers based on their ecology and phylogenetic relatedness. *Ecography*, DOI: 10.1111/ecog.03371

2017: Turvey SA, Crees JJ, Hansford J, Jeffree TE, Crumpton N, Kurniawan I, Setiyabudi E, **Guillerme** T, Paranggarimu U, Dosseto A, van den Bergh GD. Quaternary vertebrate faunas from Sumba, Indonesia: implications for Wallacean biogeography and evolution *Proceedings of the Royal Society B*, 284 20171278, DOI: 10.1098/rspb.2017.1278

2016: Kane A, Healy K, **Guillerme T**, Ruxton G, and Jackson AL. A Recipe for Scavenging - the natural history of a behaviour. *Ecography*, 39:001011, DOI: 10.1111/ecog.02817

2016: **Guillerme T** and Cooper N. Assessment of available anatomical characters for linking living mammals to fossil taxa in phylogenetic analyses. *Biology Letters*, 12 20151003 DOI: 10.1098/rsbl.2015.1003

2016: **Guillerme T** and Cooper N. Effects of missing data on topological inference using a Total Evidence approach. *Molecular Phylogenetics and Evolution* 94:146158, DOI: 10.1016/j.ympev.2015.08.023

2014: Healy K, **Guillerme T**, Finlay S, Kane A, Kelly SBA, McClean D, Kelly DJ, Donohue I, Jackson AL and Cooper N. Ecology and mode-of-life explain lifespan variation in birds and mammals. *Proceedings of the Royal Society B*, 281(1784). DOI: 10.1098/rspb.2014.0298

In review/in press - see my publications page.

AWARDS AND GRANTS

2017: Royal Society International Scientific Seminar Award (£5000)

co-awarded with Prof. Phillip Donoghue FRS and Dr Natalie Cooper

2017: Marie Skłodowska-Curie Action Seal of Excellence

2014: BES Photo Competition Student Prize (Ecology and Society)

2012: Trinity Postgraduate Research Studentship (€46800)

COMPUTER SKILLS

Coding: \mathbf{R}^* , \mathbf{C}^* , \mathbf{C}^* , \mathbf{E}^* , \mathbf{Bash}^* , \mathbf{HTML} , \mathbf{E}^* , \mathbf{git}^*

Phylogenetics: PAUP, MrBayes*, BEAST*, RAxML, PhyloBayes, CoEvol

MACSE, MAFFT, SEAview, Tracer, Mesquite, R{ape} R{phytools}, R{diversitree}, R{MCMCglmm}, R{caper}, ...

Morphometrics: R{geomorph}, MorphoJ, TPS

Graphics: Adobe Illustrator, Adobe Photoshop, Gimp, Inkscape

Developed software:

R{mulTree}: Performs MCMCglmm on multiple phylogenetic trees. *Role: main author*.

GitHub page: github.com/TGuillerme/mulTree

R{dispRity}: Measuring disparity in R. Role: main author.

GitHub page: github.com/TGuillerme/dispRity

R{SIDER}: Stable Isotope Discrimination Estimation in R. *Role: co-author*.

GitHub page: github.com/healyke/SIDER

Morphy: A C library for morphological characters phylogenetic analysis. *Role: co-author*.

GitHub page: www.morphyproject.org/

^{*} Skills taught at workshops or courses (see below)

PRESENTATIONS

Invited seminars

2017: ISEM seminar series - Université de Montpellier, France

Talk: Disparate views on mammalian disparity at the K-Pg.

2017: Silwood Social Seminars Series - Imperial College London, UK

Talk: Multidimensionality in morphology.

2016: Palaeobiology Discussion Group - University of Bristol, UK

Talk: Do we still need morphological characters data?

International Conferences

2018: Reconciling disparate perspectives on the evolution of disparity - Royal Society

International Scientific Meeting - Chicheley Hall, UK - Organiser

2017: BES Macro SIG meeting 2017 - London, UK

conference co-organiser

2017: Evolution 2017 - Portland OR, USA

Talk: Blue tail, red tail: a solution to inapplicable data in phylogenetics

Available here: goo.gl/g68NTZ

2016: BES Macro SIG meeting 2016 - Oxford, UK

Talk: Moving away from the consensus in Phylogenetic Comparative Methods

Available here: goo.gl/1PsRfi

2016: Evolution 2016 - Austin TX, USA

Palaeobiology/macroevolution session chair

Talk: New approaches to disparity-through-time analysis

Available here: goo.gl/nnSpVS

2015: Systematics Association Biennial - Oxford, UK

Talk: Total evidence phylogenies, the data issue

Available here: goo.gl/MVwMTY

2014: SVP annual meeting - Berlin, Germany

Symposium presentation (invited talk): Putting fossils in trees

Talk: Total evidence phylogenies and missing data

Available here: goo.gl/CBqSVJ

2014: BES Macroecology SIG meeting - University of Nottingham, UK

Talk: Trees with fossil and living species: the data issue

Available here: goo.gl/Sdcxon

2014: Evolution 2014 - Raleigh NC, USA

Symposium presentation: Ernst Mayr Symposium

Talk: Total evidence phylogenies: the missing data issue

Available here: goo.gl/oPfHZE

2014: Challenges in Macroecology: Scaling the Time Barrier - Natural History Museum

London, UK

Talk: Trees with fossil and living species: the missing data issue

Available here: goo.gl/c1RBER

2013: European Society for Evolutionary Biology Congress - Lisbon, Portugal

Poster: Combining fossils and molecular phylogenies

Available here: goo.gl/3HcRGH

2013: 11th International Mammalogical Congress - Queen's University, Belfast, UK

Poster: Combining fossils and molecular phylogenies

Available here: goo.gl/3HcRGH

PROFESSIONAL SERVICE

Reviewing: Biology Letters (Top reviewer for 2016), International Journal of Primatology, Methods in Ecology and Evolution, Molecular Phylogenetics and Evolution, Palaeontology, PLoS ONE, The Royal Society:

Interface focus, Scientific Reports, Systematic Biology.

Workshops:

2018: Repeatability workshop: Organize your research from the get-go! — **organiser** Australian Mammals Society Annual meeting - Brisbane, Australia

2014: Tip-Dating workshop — *helper*

Nicholas Matzke & April Wright, SVP Annual meeting - Berlin, Germany

2014: Software Carpentry Bootcamp - R and UNIX — helper Devasena Inupakutika & Aur Saraf - Nottingham, UK

Conferences:

2018: Reconciling disparate perspectives on the evolution of disparity - Royal Society

International Scientific Meeting - Chicheley Hall, UK — organiser

2017: BES Macro SIG meeting 2017 - London, UK — co-organiser

Academic memberships:

2017-present: Member of the Society for the Study of Evolution

2015-present: Member of the Systematic Association

2014-present: Member of the Society of Vertebrate Palaeontology

2013-present: Member of Society of Systematic Biologists

2013-present: British Ecological Society (BES) including Macroecology Special Interest Group

Departmental services:

2018-2019: UQ Biology Postdocs monthly meetings co-organiser.
2016-2017: Silwood Ecology and Evolution Seminar Series organiser

Biweekly Silwood Park Campus computer workshops organiser (LATEX, version control,

webpages, R packages, etc.)

2013-2015: Trinity Zoology Postgraduate student social committee chair

TEACHING

Academia

All my teaching material is available on the following GitHub page.

2017: MSc students: C for biologists

Co-teacher, 30 hours, Imperial College - UK

2017: MSc students: Phylogenetic Comparative Methods Teaching assistant, 6 hours, Natural History Museum - UK

2017: MSc students: morphometrics

Co-teacher, 6 hours, Natural History Museum - UK

2016: MSc students: Rocks 'n' clocks (including a MrBayes practical)

Teacher, 6 hours, Imperial College London - UK

2014: MSc students: Timing Evolution

Teacher, 6 hours, Trinity College Dublin - IE

2014 : Software Carpentry Bootcamp - R and UNIX *Helper*, full day, University of Nottingham - UK

2013-14: 2nd year MSc students: Statistics with R

Teaching assistant, full semester, Trinity College Dublin - IE

2013-14: 4th year undergraduate students: Evolution

Teaching assistant, full semester, Trinity College Dublin - IE

2012-13: 4th year undergraduate students: Statistics with R

Teaching assistant, full semester, Trinity College Dublin - IE

2012-13: 1st year undergraduate students: Introduction to Biodiversity Evolution and the Environment

Teaching assistant, full semester, Trinity College Dublin - IE

Student supervision

2017: Joe Craig (Imperial College London - MSc project) How does adding data change node

age estimation in morphological phylogenies?

2017: Danielle Moraviec & Hayley Swalund (Imperial College London - MSc mini projects)
2016: Victoria Stanley Tsui (Imperial College London - MSc) *A preliminary study on the suture*

morphology of skull roofs in early gnathostomes and their ancestors. - Co-supervised with

Martin Brazeau.

2016: Asiem Sanyal (Imperial College London - MSc mini project)

2014-2015: Luke Hannon (Trinity College Dublin -4th year project) Orbit size and nocturnality in

living and fossil primates. - Co-supervised with Natalie Cooper.

2013-2014: Maura Judge (Trinity College Dublin -4th year project) *Bite force and diet in mammals.*

- Co-supervised with Natalie Cooper.

FIELDWORK AND DATA COLLECTION

2013-14: Museum data collection: Smithsonian Institute Natural History Museum (Washington

DC), American Museum of Natural History (New York), Museum of Comparative Zool-

ogy (Cambridge, MA), Muséum National d'Histoire Naturelle (Paris)

2011-12: Excavations in Cruzy, France

Southern France, Continental Cretaceous

2011: Excavations in Krasiejów, Poland

Southern Poland, Continental Trias

2009-10: Excavations in Espéraza, France - Site fore-person

Southern France, Continental Cretaceous

PUBLIC ENGAGEMENT AND OUTREACH

- Regular contributor to EcoEvo@TCD blog; rated best Irish Science and Technology blog (2014) and Bronze Award Irish Blog (2015).
- Complete list available here: goo.gl/MAq9oA
- Featured posts: How do lego cars evolve?; If you please draw me a dino...; The more the better?.
- Threesis video (early stage thesis project explained in three minutes winner of the NERD club threesis award 2013).

Outreach Events

2015 & 2014 & 2013: Trinity College Dublin Zoology Department, Researcher's night – participant

2014: 1st Soapbox Science Ireland – *helper*

2014: "What can we learn from skeletons", comparative anatomy practical – *Co-organiser*

2013: Trinity College Zoology Museum guided tours – tour guide

2010: Introduction to palaeontological excavations, Musée des Dinosaures (FR) – animator