Robert J. Noble

<u>robjohnnoble.github.io</u> <u>scholar.google.com/citations?user=IDDprHkAAAAJ</u>

Research focus: Mathematical and computational modelling of cancer evolution and treatment.

Academic employment

2020-	Department of Mathematics, City, University of London Lecturer in Applied Mathematics
2018-2020	Department of Evolutionary Biology and Environmental Studies, University of Zurich Postdoctoral researcher (20% contract) advised by Hanna Kokko
2017-2020	Department of Biosystems Science and Engineering, ETH Zurich Postdoctoral researcher (80% contract from May 2018) advised by Niko Beerenwinkel
2014-2017	Institut des Sciences de l'Evolution de Montpellier (ISEM) Postdoctoral researcher advised by Michael Hochberg

Education

2009-2014	DPhil in Zoology (mathematical biology), University of Oxford
	Supervisors: Mario Recker and Sunetra Gupta
1999-2003	Master of Mathematics (First Class), University of York

Publications and preprints (* denotes equal contributions)

2023	A seven-step guide to spatial, agent-based modelling of tumour evolution Colyer B, Bak M, Basanta D, Noble R	arXiv 2311.03569
2023	A new universal system of tree shape indices Noble R, Verity K	bioRxiv 2023.07.17.549219
2023	Warlock: an automated computational workflow for simulating spatially structured tumour evolution Bak M, Colyer B, Manojlović V, Noble R	arXiv 2301.07808
2023	A survey of open questions in adaptive therapy: bridging mathematics and clinical translation West J, Adler F, Gallaher J,, Noble R , Viossat Y, Basanta D, Anderson ARA	eLife 12:e84263
2022	Robust, universal tree balance indices Lemant J, Le Sueur C, Manojlović V, Noble RJ	Syst. Biol. 71, 1210-24
2021	Spatial structure governs the mode of tumour evolution Noble R, Burri D, Le Sueur C, Lemant J, Viossat Y, Kather JN, Beerenwinkel N	Nature Ecol. Evol. 6, 207-17
2021	Drug-induced resistance evolution necessitates less aggressive treatment Kuosmanen T, Cairns J, Noble R , Beerenwinkel N, Mononen T, Mustonen V	PLoS Comput. Biol. 17: e1009418
2021	Inferring the dynamic of mutated hematopoietic stem and progenitor cells induced by IFNa in myeloproliferative neoplasms Mosca M*, Hermange G*, Tisserand A*, Noble R *,, Plo I	Blood 138, 2231-43
2021	Paracrine behaviors arbitrate parasite-like interactions between tumor subclones Noble R , Walther V, Roumestand C, Hochberg ME, Hibner U, Lassus P	Front. Ecol. Evol. 9:675638
2021	A theoretical analysis of tumour containment Viossat Y, Noble R	Nature Ecol. Evol. 5, 826-35
2020	Identifying key questions in the ecology and evolution of cancer Dujon A,, Noble R ,, Thomas F, Ujvari B	Evol. Appl. eva.13190
2020	When, why and how clonal diversity predicts survival Noble R*, Burley JT*, Le Sueur C, Hochberg ME	Evol. Appl. eva.13057
2017	Spatial competition constrains resistance to targeted cancer therapy Bacevic K*, Noble R *,, Hochberg ME, Krasinska L, Fisher D	Nature Commun. 8, 1995

2017	Antibiotic stress selects against cooperation in the pathogenic bacterium Pseudomonas aeruginosa Vasse M*, Noble R *,, Hochberg ME	PNAS 114, 546-51
2017	A framework for how environment contributes to cancer risk Hochberg ME, Noble R	Ecol. Lett. 20, 117-34
2016	Overestimating the role of environment in cancers Noble R, Kaltz O, Nunney L, Hochberg ME	Cancer Prev. Res. 9, 773-6
2016	A hypothesis to explain cancers in confined colonies of naked mole rats Hochberg ME, Noble RJ , Braude S	bioRxiv 10.1101/079012
2015	Peto's paradox and human cancers Noble R, Kaltz O, Hochberg ME	Phil. Trans. B 370, 20150104
2013	The antigenic switching network of Plasmodium falciparum and its implications for the immuno-epidemiology of malaria Noble R*, Christodoulou Z*, Pinches R, Kyes S, Recker M, Newbold CI	eLife 2013.2:e01074
2012	Erasing the epigenetic memory and beginning to switch—the onset of antigenic switching of var genes in Plasmodium falciparum Fastman Y, Noble R , Recker M, Dzikowski R	PLoS ONE 7, e34168
2012	A statistically rigorous method for determining antigenic switching networks Noble R , Recker M	PLoS ONE 7, e39335
Softwa	re	
2022	warlock: Automated computational workflow for simulating tumour evolution	GitHub
2019	demon: Deme-based oncology model	GitHub
2017	ggmuller: Create Muller plots of evolutionary dynamics	CRAN
Teachir	ng	
2020-	Supervision (City, University of London) PhD primary supervisor, Kimberley Verity (from 2022) PhD primary supervisor, Blair Colyer (from 2021) PhD primary supervisor, Veselin Manojlović (from 2020) PhD secondary supervisor, Youssef Arafat (from 2021) PhD secondary supervisor, Hasan Haq (from 2021) Final year BS-MS thesis, Srishti Patil Postdoctoral research supervisor, Maciej Bak Lecturing and tutoring as module leader (City, University of London) Mathematical processes for finance (BSc) Mathematics for economists post A Level (BSc) Tutoring (City, University of London) Algebra (BSc); Functions, vectors and calculus (BSc); Final-year group projects	(BSc)
2017- 2020	Supervision (ETH Zurich) Second year MSc thesis, Alexander Stein (next step: PhD at QMUL) Second year MSc thesis, Jeanne Lemant (next step: PhD at University of Basel) Second year MSc thesis, Dominik Burri (next step: PhD at University of Basel) Research internship (eight months), Cécile Le Sueur (next step: PhD at EMBL) Lecturing and tutoring (ETH Zurich) Evolutionary dynamics (MSc three terms)	
2016	Supervision (ISEM) First year MEME MSc project, John Burley (next step: PhD at Brown University)	
2010- 2013	Tutoring (University of Oxford) Quantitative methods (BSc two terms) Demonstrating (University of Oxford) Quantitative methods (BSc; three terms); Epidemiology (BSc; two terms); Epidemiology (MSc one term)	niological models

Funding and awards

Co-awardee of US NSF grant *Quantifying and modeling the transmission dynamics of bivalve transmissible neoplasia* (postdoc and travel support in 2023-27)

Sub-awardee of US NCI Arizona Cancer Evolution Center U54 grant (\$150K in 2020-2023)

Awardee of LMS Research in Pairs grant (£850 in 2024-2026)

Awardee of ACE Pilot Project funding (\$8K in 2022)

Awardee of City University Pump Priming funding (£10K in 2022)

Co-awardee of FMJH grant Optimization of a new type of cancer therapy (€5K in 2019-2020)

Biotechnology and Biological Sciences Research Council PhD fellowship 2009-2013

Travel grants: Lorentz Center 2017; Moffitt Cancer Center 2015; ECMTB 2011

City University Images of Research Competition 2020-21: First prize (£250)

External professional activities

Secretary (elected) of the International Society for Evolution, Ecology and Cancer 2022-2025; Advisory Committee member (elected) 2018-2021

Supervisory Board member of the EvoGamesPlus Innovative Training Network 2021-2025

Associate Editor of Journal of Molecular Evolution; Guest Associate Editor of PLoS Computational Biology

Reviewer: American Naturalist, Cancer Research, Computational and Systems Oncology, Evolutionary Applications, F1000Research, Frontiers Ecology and Evolution, Journal of Evolutionary Biology, Journal of Theoretical Biology, Nature Communications, Nature Ecology & Evolution, Nature Genetics Nature Reviews Bioengineering, npj Genomic Medicine, Peer Community in Evolutionary Biology, PLoS Computational Biology, PNAS, Proceedings of the Royal Society B, Royal Society Open Science, Scientific Reports

Co-organizer: "Evolutionary approaches to understand cancer across scales" symposium (SMBE 2023); "Cancer Adaptive Therapy Models" workshop (2020); "Aging & cancer through the lens of evolution" symposium (ESEB 2019); "How does spatial structure affect tumour evolution?" symposium (MBE 2017)

Other employment

2008-2009 International HIV/AIDS Alliance: Communications

TU Dresden (hosted by Andreas Deutsch)

2004-2008 AVERT (HIV/AIDS charity): Science/heath communication and web development

Invited departmental seminars

Apr 2023	Quantifying and explaining modes of evolution University of Leeds (hosted by Tyler Cassidy)
Feb 2022	Explaining the modes of tumour evolution University of Warwick (virtual, hosted by Simon Graham)
Oct 2021	Explaining the modes of tumour evolution University of Basel (virtual, hosted by Dominik Burri)
Sept 2020	Characterizing and forecasting tumour evolution Cancer Research UK Cambridge Institute (virtual, hosted by Florian Markowetz)
Jun 2020	Characterizing and forecasting tumour evolution Virtual Seminar on Modeling Biocomplexity (hosted by Andreas Deutsch)
May 2020	Characterizing and forecasting tumour evolution Moffitt Cancer Center (virtual, hosted by David Basanta)
Jan 2020	Cancer: evolution, ecology and bad luck University of Bath (hosted by Ben Ashby)
Dec 2019	The logic of containing tumours University of Oxford (hosted by Eamonn Gaffney)
Sep 2019	Cancer: evolution, ecology and bad luck University of Southampton (hosted by Lindy Holden-Dye)
Feb 2019	Characterising the evolutionary modes of cancer and normal tissue

Mar 2018	Characterising the evolutionary modes of cancer and normal tissue University of Basel (hosted by Richard Neher)
Feb 2018	The mode and predictability of intra-tumour evolution Wellcome Sanger Institute (hosted by Iñigo Martincorena)
Dec 2017	The mode and predictability of intra-tumour evolution Boston University (hosted by Kirill Korolev)
Nov 2017	Spatial constraints on intratumour evolution Harvard University (hosted by Martin Novak)
May 2017	Models for understanding tumour evolution and improving cancer therapy University of Edinburgh (hosted by Bartlomiej Waclaw)
Mar 2017	Evolution, ecology, and cancer risk: from naked mole rats to modern humans Chalmers University (hosted by Philip Gerlee)
Sep 2016	Cancer: evolution, ecology and bad luck Harvard University (hosted by Martin Novak)
Feb 2015	Data-based modelling of tumour evolution Moffitt Cancer Center (hosted by Robert Gatenby)
Conference	talks
May 2023	Tumour heterogeneity and survival of cancer patients Invited talk at French Cancer Society training session on tumour heterogeneity, Paris
Mar 2023	Evolutionary approaches to overcoming cancer cell plasticity Invited talk at the Systems Approaches Towards Cancer Cell Plasticity Symposium, London
Sep 2022	Robust, universal tree balance indices European Conference on Mathematical and Theoretical Biology, Heidelberg
Jul 2022	Parasite-like interactions between tumour subclones Mathematical Models in Ecology and Evolution, Reading
Jul 2021	The evolutionary logic of tumour containment International Society for Evolution, Medicine & Public Health conference (virtual)
Jun 2021	The evolutionary logic of tumour containment Evolution conference (virtual)
Jun 2021	Explaining modes of tumour evolution Society for Mathematical Biology conference (virtual)
Dec 2020	The logic of containing tumours Cancer Adaptive Therapy Models workshop (virtual)
Oct 2020	Characterizing and forecasting tumour evolution International Symposium on Mathematical and Computational Oncology (virtual)
Aug 2020	The logic of containing tumours Invited talk at the Society for Mathematical Biology conference (virtual)
Aug 2019	Spatial competition constrains resistance to targeted cancer therapy International Society for Evolution, Medicine & Public Health conference, Zurich
Jul 2019	Spatial structure governs the mode of tumour evolution Intelligent Systems for Molecular Biology / European Conference on Comp. Biology, Basel
Jun 2019	Spatial structure governs the mode of tumour evolution Modelling Ecology & Evolution Zurich seminar, Zurich
Sep 2018	Characterising the evolutionary modes of cancer and normal tissue Evolutionary Models of Structured Populations workshop, Plön
Dec 2017	Spatial competition constrains resistance to targeted cancer therapy International Society for Evolution, Ecology and Cancer Conference, Tempe
Oct 2017	Impact of tissue architecture on the nature and predictability of tumour evolution Satellite Symposium to the Louis-Jeantet Symposium, Geneva
Sep 2017	Impact of tissue architecture on the nature and predictability of tumour evolution Basel Computational Biology Conference, Basel
Jul 2017	Impact of tissue architecture on the nature and predictability of tumour evolution Intelligent Systems for Molecular Biology / European Conference on Comp. Biology, Prague

Apr 2017	Evolutionary ecology of senescence and cancer risk: from naked mole rats to modern humans Modelling Biological Evolution conference, Leicester
Nov 2016	Controlling drug resistance with adaptive therapy Invited talk at the second Modeling Tumour Evolution conference, Bielefeld
Sep 2016	Cancer: evolution, ecology and bad luck Invited talk at the first Modelling Tumour Evolution conference, Bielefeld
Jul 2016	Cancer risk: evolution, ecology and bad luck Joint Meeting of ESMTB & Society for Mathematical Biology, Nottingham
Dec 2015	Peto's paradox and human cancers Third International Biannual Evolution and Cancer Conference, San Francisco
Sep 2015	Modelling ecological interactions of cancer clones Cancer Evolution Through Space and Time workshop, Plön
Apr 2015	Eco-evolutionary models of tumour heterogeneity Invited talk at the Modelling Biological Evolution conference, Leicester
June 2011	Using iterative methods to determine an antigenic switching network in Plasmodium falciparum European Conference on Mathematical and Theoretical Biology, Krakow
May 2011	Determining the switch pathway of the var gene repertoire of Plasmodium falciparum Biology and Pathology of the Malaria Parasite, Heidelberg