Robert J. Noble

robjohnnoble.github.io
scholar.google.com/citations?user=lDDprHkAAAAJ

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

Academic employment

2020- Department of Mathematics, City St George's, University of London
Senior Lecturer in Applied Mathematics (2024-)
Lecturer in Applied Mathematics (2020-24)

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 and qualifications

2024 Advance HE Fellowship

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)

Year	Title / Authors	Source
2025	*Evolutionary tree balance predicts disease-free survival in the TRACERx non-small cell lung cancer cohort* Verity K, **Noble R**	medRxiv 2025.11.22.25340797
2025	*Preventing evolutionary rescue in cancer* Patil S, Viossat Y, **Noble R**	Genetics iyaf255
2025	*Expected and minimal values of a universal tree balance index* Manojlović V*, Ahmed A*, Viossat Y, **Noble R**	arXiv 2507.08615
2024	*A seven-step guide to spatial, agent-based modelling of tumour evolution* Colyer B, Bak M, Basanta D, **Noble R**	Evol. Appl. eva.13687
2023	*Selective sweep probabilities in spatially expanding populations* Stein A, Kizhuttil R, Bak M, **Noble RJ**	bioRxiv 2023.11.27.568915
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 IFNα 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*	Front. Ecol. Evol.
	Noble R, Walther V, Roumestand C, Hochberg ME, Hibner U, Lassus P	9:675638
2021	*A theoretical analysis of tumour containment*	Nature Ecol. Evol.
	Viossat Y, **Noble R**	5, 826-35
2020	*Identifying key questions in the ecology and evolution of cancer*	Evol. Appl.
	Dujon A, …, **Noble R**, …, Thomas F, Ujvari B	eva.13190
2020	*When, why and how tumour clonal diversity predicts survival*	Evol. Appl.
	Noble R*, Burley JT*, Le Sueur C, Hochberg ME	eva.13057
2017	*Spatial competition constrains resistance to targeted cancer therapy*	Nature Commun.
	Bacevic K*, **Noble R***, …, Hochberg ME, Krasinska L, Fisher D	8, 1995
2017	*Antibiotic stress selects against cooperation in the pathogenic bacterium* Pseudomonas aeruginosa	PNAS 114, 546-51
	Vasse M*, **Noble R***, …, Hochberg ME	
2017	*A framework for how environment contributes to cancer risk*	Ecol. Lett.
	Hochberg ME, **Noble R**	20, 117-34
2016	*Overestimating the role of environment in cancers*	Cancer Prev. Res.
	Noble R, Kaltz O, Nunney L, Hochberg ME	9, 773-6
2016	*A hypothesis to explain cancers in confined colonies of naked mole rats*	bioRxiv
	Hochberg ME, **Noble RJ**, Braude S	10.1101/079012
2015	*Peto's paradox and human cancers*	Phil. Trans. B
	Noble R, Kaltz O, Hochberg ME	370, 20150104
2013	*The antigenic switching network of* Plasmodium falciparum *and its implications for the immuno-epidemiology of malaria*	eLife 2013.2:e01074
	Noble R*, Christodoulou Z*, Pinches R, Kyes S, Recker M, Newbold CI	
2012	*Erasing the epigenetic memory and beginning to switch—the onset of antigenic switching of* var *genes in* Plasmodium falciparum	PLoS ONE 7, e34168
	Fastman Y, **Noble R**, Recker M, Dzikowski R	
2012	*A statistically rigorous method for determining antigenic switching networks*	PLoS ONE
	Noble R, Recker M	7, e39335

Software

| 2022 | *warlock: Automated computational workflow for simulating tumour evolution* | GitHub |
| 2017 | *ggmuller: Create Muller plots of evolutionary dynamics* | CRAN |

Teaching

2020- **Supervision (City St George's, University of London)**

PhD primary supervisor: Kate Bostock (2024-); Alexandru Chitiga (2024-); Kimberley Verity (2022-); Blair Colyer (2021-25); Veselin Manojlović (2020-24)

PhD second supervisor: Manjot Singh Bedi (2023-); Youssef Arafat (2021-25); Hasan Haq (2021-25)

External master's project supervisor: Srishti Patil at IISER (2022-23); Carissa Mayo at PNRI (2024-)

Lecturing and tutoring as module leader (City St George's, University of London)

Mathematical processes for finance (BSc); Mathematics for economists post A Level (BSc)

Tutoring (City St George's, University of London)

Algebra (BSc); Functions, vectors and calculus (BSc); Final-year group projects (BSc)

2017- **Supervision (ETH Zurich)**
2020 Master's project supervisor: Alexander Stein; Jeanne Lemant; Dominik Burri

Research internship (eight months): Cécile Le Sueur

Lecturing and tutoring (ETH Zurich)
Evolutionary dynamics (MSc)

2016	**Supervision (ISEM)**
	First year MEME MSc project: John Burley

2010- 2013	**Tutoring & demonstrating (University of Oxford)**
	Quantitative methods (BSc); Epidemiology (BSc); Epidemiological models (MSc)

Funding and awards

Co-awardee of US NSF grant *Quantifying and modeling the transmission dynamics of bivalve transmissible neoplasia* (co-supervised PhD student in 2023-27)

Sub-awardee of US NCI Arizona Cancer Evolution Center U54 grant (£110K in 2020-25)

Awardee of LMS Research in Pairs grant (£850 in 2024-26)

Awardee of Arizona Cancer Evolution Center Pilot Project funding (£6K 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* (£4K in 2019-20)

Biotechnology and Biological Sciences Research Council PhD fellowship (2009-13)

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

City University Images of Research Competition 2021: First prize (£250)

External professional activities

Associate Editor of the Journal of Evolutionary Biology (2025-); Guest Associate Editor of PLoS Computational Biology (2021-); Editorial Board Member of Scientific Reports (2024-25); Associate Editor of the Journal of Molecular Evolution (2022-25)

Vice-President (elected 2025) of the International Society for Evolution, Ecology and Cancer; previously elected Secretary (2022-25) and Advisory Committee member (2018-21)

Supervisory Board member of the EvoGamesPlus Innovative Training Network (2021-25)

PhD external examiner: University of Oxford (2025); Queen Mary University of London (2022, 2025), Institute of Cancer Research (2021)

PhD advisory committee member: University of Montpellier, ETH Zurich, Université Grenoble Alpes

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)

Peer review for journals: American Naturalist, Bulletin of Mathematical Biology, Cancer Cell International, Cancer Research, Computational and Systems Oncology, eLife, Evolution, Evolutionary Applications, F1000Research, Frontiers Ecology and Evolution, Journal of Evolutionary Biology, Journal of Theoretical Biology, Mathematical Biosciences, 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

Peer review for funding agencies: French National Research Agency, Pancreatic Cancer UK

Other employment

2008-09	International HIV/AIDS Alliance (now known as Frontline AIDS): Communications
2004-08	AVERT (HIV/AIDS charity): Science and health communication

Invited seminar talks

Nov 2025	*Using mathematical modelling to support trials of evolutionary cancer treatment strategies* Institut de Recherche en Cancérologie de Montpellier (hosted by Daniel Fisher)
Sep 2025	*Understanding, predicting and controlling stochastic cancer evolution* University of Oxford (hosted by Bridget Penman)
Mar 2025	*Understanding, predicting and controlling stochastic cancer evolution* Mathematical and Computational Modeling of Cancer seminar (virtual, hosted by Ivana Bozic)

Dec 2024   *Understanding, predicting and controlling stochastic cancer evolution*
University of Sheffield (hosted by Alex Fletcher)

Feb 2024   *New directions in mathematical oncology*
Queen Mary University of London (hosted by Weini Huang)

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*
TU Dresden (hosted by Andreas Deutsch)

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

Oct 2025   *Using mathematical models to support clinical trials of evolutionary cancer therapies*
Plenary talk at the Mathematical Oncology meeting, St. Pete Beach

Aug 2025   *Selective sweep probabilities in tumours and other range expansions*
European Society for Evolutionary Biology conference, Barcelona

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