

Robert J. Noble

robjohnnoble.github.io

scholar.google.com/citations?user=IDDprHkAAAAJ

Academic employment

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

Education

- Oct 2009- Department of Zoology, **University of Oxford**
Jan 2014 DPhil: *Antigenic variation and its evolution in P. falciparum malaria*
Supervisors: Sunetra Gupta and Mario Recker
- Oct 1999- **University of York**
Jul 2003 Master of Mathematics (First Class)

Publications

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|------|---|--|
| 2021 | <i>A theoretical analysis of tumour containment</i>
Viossat Y, Noble R | Nature Ecol. Evol. |
| 2020 | <i>Identifying key questions in the ecology and evolution of cancer</i>
Dujon A, ..., Noble R , ..., Thomas F, Ujvari B | Evol. Appl.
eva.13190 |
| 2020 | <i>When, why and how clonal diversity predicts survival</i>
Noble R *, Burley JT*, Le Sueur C, Hochberg ME
(* equal contributions) | Evol. Appl.
eva.13057 |
| 2017 | <i>Spatial competition constrains resistance to targeted cancer therapy</i>
Bacevic K*, Noble R *, Soffar A, Ammar OW, Boszonyik B, Prieto S, Vincent C, Hochberg ME, Krasinska L, Fisher D
(* equal contributions) | Nature Commun.
8, 1995 |
| 2017 | <i>Antibiotic stress selects against cooperation in the pathogenic bacterium Pseudomonas aeruginosa</i>
Vasse M*, Noble R *, Akhmetzhanov AR, Torres-Barceló C, Gurney J, Simon Benateau, Gougat-Barbera C, Kaltz O, Hochberg ME
(* equal contributions) | PNAS
114, 546-51 |
| 2017 | <i>A framework for how environment contributes to cancer risk</i>
Hochberg ME, Noble R | Ecology Letters
20, 117-34 |
| 2016 | <i>Overestimating the role of environment in cancers</i>
Noble R , Kaltz O, Nunnery L, Hochberg ME | Cancer Prev. Res.
9, 773-6 |
| 2015 | <i>Peto's paradox and human cancers</i>
Noble R , Kaltz O, Hochberg ME | Phil. Trans. B
370, 20150104 |

2013	<i>The antigenic switching network of Plasmodium falciparum and its implications for the immuno-epidemiology of malaria</i> Noble R* , Christodoulou Z*, Pinches R, Kyes S, Recker M, Newbold CI (* equal contributions)	eLife 2013.2:e01074
2012	<i>Erasing the Epigenetic Memory and Beginning to Switch—The Onset of Antigenic Switching of var Genes in Plasmodium falciparum</i> Fastman Y, Noble R , Recker M, Dzikowski R	PLoS ONE 7, e34168
2012	<i>A statistically rigorous method for determining antigenic switching networks</i> Noble R , Recker M	PLoS ONE 7, e39335

Submitted for publication

Submitted	<i>Paracrine behaviors arbitrate parasite-like interactions between tumor subclones</i> Noble R , Walther V, Roumestand C, Hibner U, Hochberg ME, Lassus P.	bioRxiv 10.1101/2020.12.14.422649v2
Submitted	<i>Determinants of successful IFNα therapy in myeloproliferative neoplasms</i> Mosca M*, Hermange G*, Tisserand A*, Noble R* , ..., Plo I (* equal contributions)	
Submitted	<i>Drug-induced resistance evolution necessitates less aggressive treatment</i> Kuosmanen T, Cairns J, Noble R , Beerenwinkel N, Mononen T, Mustonen V	bioRxiv 10.1101/2020.10.07.330134
Submitted	<i>Spatial structure governs the mode of tumour evolution</i> Noble R , Burri D, Le Sueur C, Lemant J, Viossat Y, Kather JN, Beerenwinkel N	bioRxiv 10.1101/586735

Software

2017	<i>ggmuller: Create Muller Plots of Evolutionary Dynamics</i>	CRAN
2019	<i>demon: Deme-based oncology model</i>	GitHub

Teaching

2020-	Supervision (City, University of London) PhD, Veselin Manojlovic	
	Lecturing and tutoring as module leader (City, University of London) Mathematical processes for finance (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: research scientist at Swiss TPH) 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 assistance (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	Supervision (University of Oxford) Second year BSc project, Charlotte Ward	
	Tutoring (University of Oxford) Quantitative Methods (BSc two terms)	
	Demonstrating (University of Oxford)	

Quantitative Methods (BSc; three terms); Epidemiology (BSc; two terms); Epidemiological Models (MSc one term)

Funding

\$150K personal funding for 2020-2023 from the NCI, via the Arizona Cancer Evolution Center

Co-awardee of Fondation Mathématique Jacques Hadamard grant *Optimization of a new type of cancer therapy* (€5K to support international collaboration in 2019-2020)

Biotechnology and Biological Sciences Research Council PhD fellowship 2009-2013

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

Professional activities

Elected Advisory Committee member of the International Society for Evolution, Ecology and Cancer 2018-21

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

Co-organizer: “Cancer Adaptive Therapy Models” workshop (2020); “Aging & cancer through the lens of evolution” symposium (ESEB conference 2019); “How does spatial structure affect tumour evolution?” symposium (MBE conference 2017)

Other employment

Dec 2008- **International HIV/AIDS Alliance**
Sep 2009 Communications

Aug 2004- **AVERT (HIV/AIDS charity)**
Dec 2008 Science/health communication and web development

Invited departmental seminars

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 Bartłomiej 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 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 virtual conference
- 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