Robert J. Noble

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

Academic employment

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

Education

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

Publications

2020	When, why and how clonal diversity predicts survival Noble R*, Burley JT*, Le Sueur C, Hochberg ME (* equal contributions)	Evol. Appl. eva.13057
2017	Spatial competition constrains resistance to targeted cancer therapy 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	Antibiotic stress selects against cooperation in the pathogenic bacterium Pseudomonas aeruginosa 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	A framework for how environment contributes to cancer risk Hochberg ME, Noble R	Ecology Letters 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
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 (* equal contributions)	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 PLoS ONE Noble R, Recker M 7, e39335

Submitted for publication

Submitted Drug-induced resistance evolution necessitates less aggressive treatment bioRxiv 10.1101/ Kuosmanen T, Cairns J, Noble R, Beerenwinkel N, Mononen T, Mustonen V 2020.10.07.330134

In revision The logic of containing tumors **bioRxiv** 10.1101/

> Viossat Y, Noble R 2020.01.22.915355

Spatial structure governs the mode of tumour evolution bioRxiv 10.1101/

> Noble R, Burri D, Le Sueur C, Lemant J, Viossat Y, Kather JN, 586735

Beerenwinkel N

Software

2017 **CRAN** ggmuller: Create Muller Plots of Evolutionary Dynamics 2019 demon: Deme-based oncology model **GitHub**

Teaching

2020-Supervision (City, University of London)

PhD, Veselin Manoilovic

2017-**Supervision (ETH Zurich)**

2020 Second year MSc thesis, Alexander Stein (next step: in progress)

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 (ETH Zurich)

Evolutionary dynamics (MSc three terms; group tutorials, setting and marking exercises, self-

authored two-hour lecture)

2016 Supervision (ISEM)

First year MEME MSc project, John Burley (next step: PhD at Brown University)

2010-Supervision (University of Oxford)

2013 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

€140K 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, 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

Symposia: Co-organizer of "How does spatial structure affect tumour evolution?" (MBE conference 2017); co-organizer of "Aging & cancer through the lens of evolution" (ESEB conference 2019)

Dec 2008- International HIV/AIDS Alliance, Preece House, Hove, BN3 1RE

Other employment

	Communications			
	AVERT, 4 Brighton Road, Horsham, West Sussex, RH13 5BA Science/heath 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 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 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
Mar 2015	Modelling the evolution of tumour heterogeneity Invited talk at the Evolution and Cancer Conference, Montpellier
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