

Robert J. Noble

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

Research interests

Using mathematical and computational models to investigate the evolution and ecology of cancer.

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, Mathematical biology, University of Oxford
Supervisors: Sunetra Gupta and Mario Recker
- 1999-2003 Master of Mathematics (First Class), University of York

Publications

* denotes equal contributions

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|------|--|---------------------------------|
| 2021 | <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 | Nature Ecol. Evol. |
| 2021 | <i>Drug-induced resistance evolution necessitates less aggressive treatment</i>
Kuosmanen T, Cairns J, Noble R , Beerenwinkel N, Mononen T, Mustonen V | PLoS Comput. Biol. 17: e1009418 |
| 2021 | <i>Inferring the dynamic of mutated hematopoietic stem and progenitor cells induced by IFNα in myeloproliferative neoplasms</i>
Mosca M*, Hermange G*, Tisserand A*, Noble R *, ..., Plo I | Blood
blood.2021010986 |
| 2021 | <i>Paracrine behaviors arbitrate parasite-like interactions between tumor subclones</i>
Noble R , Walther V, Roumestand C, Hochberg ME, Hibner U, Lassus P | Front. Ecol. Evol.
9:675638 |
| 2021 | <i>A theoretical analysis of tumour containment</i>
Viossat Y, Noble R | Nature Ecol. Evol.
5, 826-35 |
| 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 | 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 | 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, Benateau S, Gougat-Barbera C, Kaltz O, Hochberg ME | PNAS
114, 546-51 |
| 2017 | <i>A framework for how environment contributes to cancer risk</i>
Hochberg ME, Noble R | Ecol. Lett.
20, 117-34 |

2016	<i>Overestimating the role of environment in cancers</i> Noble R , Kaltz O, Nunney 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	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

In revision	<i>Robust, Universal Tree Balance Indices</i> Lemant J, Le Sueur C, Manojlović V, Noble RJ	bioRxiv 10.1101/ 2021.08.25.457695
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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 primary supervisor, Veselin Manojlović PhD primary supervisor, Blair Colyer PhD secondary supervisor, Youssef Arafat PhD secondary supervisor, Hasan Haq 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)	
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 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	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 and awards

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

Professional activities

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

Guest Associate Editor: PLoS Computational Biology

Reviewer: American Naturalist, Cancer Research, Evolutionary Applications, F1000Research, Frontiers Ecology and Evolution, Journal of Evolutionary Biology, Journal of Theoretical Biology, Nature Communications, Nature Ecology & Evolution, Nature Genetics, 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

2008-2009 International HIV/AIDS Alliance: Communications

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

Invited departmental seminars

- 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 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

- 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