

## Robert J. Noble

D-BSSE, ETH Zurich, Basel, Switzerland; [robert.noble@bsse.ethz.ch](mailto:robert.noble@bsse.ethz.ch)  
IEU, University of Zurich, Zurich, Switzerland; [robert.noble@ieu.uzh.ch](mailto:robert.noble@ieu.uzh.ch)  
<https://scholar.google.com/citations?user=IDDprHkAAAAJ>

### 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)

### Academic employment

May 2018- Department of Evolutionary Biology and Environmental Studies, **University of Zurich**  
Postdoctoral researcher (20% contract) advised by **Hanna Kokko**  
Affiliated with the Arizona Cancer Evolution Center (led by Carlo Maley)  
  
Feb 2017- Department of Biosystems Science and Engineering, **ETH Zurich**  
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**

### Publications as first or co-first author

2017	<i>Spatial competition constrains resistance to targeted cancer therapy</i> Bacevic K*, <b>Noble R*</b> , Soffar A, Ammar OW, Boszonyik B, Prieto S, Vincent C, Hochberg ME, Krasinska L, Fisher D (* equal contributions)	<b>Nature Commun.</b> 8, 1995
2017	<i>Antibiotic stress selects against cooperation in the pathogenic bacterium Pseudomonas aeruginosa</i> Vasse M*, <b>Noble R*</b> , Akhmetzhanov AR, Torres-Barceló C, Gurney J, Simon Benateau, Gougat-Barbera C, Kaltz O, Hochberg ME (* equal contributions)	<b>PNAS</b> 114, 546-51
2016	<i>Overestimating the role of environment in cancers</i> <b>Noble R</b> , Kaltz O, Nunney L, Hochberg ME	<b>Cancer Prev. Res.</b> 9, 773-6
2015	<i>Peto's paradox and human cancers</i> <b>Noble R</b> , Kaltz O, Hochberg ME	<b>Phil. Trans. B</b> 370, 20150104
2013	<i>The antigenic switching network of Plasmodium falciparum and its implications for the immuno-epidemiology of malaria</i> <b>Noble R*</b> , Christodoulou Z*, Pinches R, Kyes S, Recker M, Newbold CI (* equal contributions)	<b>eLife</b> 2013.2:e01074
2012	<i>A statistically rigorous method for determining antigenic switching networks</i> <b>Noble R</b> , Recker M	<b>PLoS ONE</b> 7, e39335

### Other publications

2017	<i>A framework for how environment contributes to cancer risk</i> Hochberg ME, <b>Noble R</b>	<b>Ecology Letters</b> 20, 117-34
2012	<i>Erasing the Epigenetic Memory and Beginning to Switch—The Onset of Antigenic Switching of var Genes in Plasmodium falciparum</i> Fastman Y, <b>Noble R</b> , Recker M, Dzikowski R	<b>PLoS ONE</b> 7, e34168

**Submitted for publication**

In review	<i>The logic of containing tumors</i> Viossat Y, <b>Noble R</b>	<b>bioRxiv</b> 10.1101/ 2020.01.22.915355
In revision	<i>When, why and how clonal diversity predicts future tumour growth</i> <b>Noble R*</b> , Burley JT*, Le Sueur C, Hochberg ME (* equal contributions)	<b>bioRxiv</b> 10.1101/ 2019.12.17.879270
In revision	<i>Spatial structure governs the mode of tumour evolution</i> <b>Noble R</b> , Burri D, Kather JN, Beerenwinkel N	<b>bioRxiv</b> 10.1101/ 586735

**Software**

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

**Teaching**

2017- 2019	<b>Supervision (ETH Zurich)</b> Second year MSc thesis, Alexander Stein (six months; ongoing): <i>Modelling solid tumour growth and evolution with diffusion approximations</i> Second year MSc thesis, Jeanne Lemant (six months; ongoing): <i>Sample vs clone trees: A mathematical analysis of tumour phylogenies</i> Second year MSc thesis, Dominik Burri (six months): <i>Study of clonal selection in healthy epidermal tissue</i> Research internship Cécile Le Sueur (seven months, ongoing) in computational modelling of tumour evolution First year MSc lab rotation project, Dominik Burri: <i>The speed of propagation in a model of invasive cancer</i>  <b>Lecturing and tutoring (ETH Zurich)</b> Evolutionary dynamics (MSc; three terms; group tutorials, setting and marking exercises, and a self-authored two-hour lecture)
2016	<b>Supervision (ISEM)</b> First year MEME MSc project, John Burley: <i>Forecasting tumour growth</i>
2010- 2013	<b>Supervision (University of Oxford)</b> Second year BSc project, Charlotte Ward: <i>Calculating the <math>R_0</math> of malaria</i>  <b>Tutoring (University of Oxford)</b> Quantitative Methods (BSc; two terms)  <b>Demonstrating (University of Oxford)</b> Quantitative Methods (BSc; three terms); Epidemiology (BSc; two terms); Epidemiological Models (MSc; one term)

**Funding**

€17K personal funding per year until May 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: 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)

## Other employment

Dec 2008- **International HIV/AIDS Alliance, Preece House, Hove, BN3 1RE**  
Sep 2009 Communications

Aug 2004- **AVERT, 4 Brighton Road, Horsham, West Sussex, RH13 5BA**  
Dec 2008 Science/health communication and web development

## Invited departmental seminars

- Jan 2020 *Cancer: evolution, ecology and bad luck*  
University of Bath (hosted by Ben Ashby)
- 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)  
Part of two weeks as a visiting researcher in the Program for Evolutionary Dynamics
- 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)  
Part of two weeks as a visiting researcher in the Integrated Mathematical Oncology department

## Conference talks

- 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 European Society for Mathematical and Theoretical Biology & 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
- Jun 2015 *Peto's paradox and human cancers*  
Invited talk at the Institute of Cancer Research, London
- 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
- Nov 2014 *Data-based modelling of tumour evolution.*  
Joint Meeting: Institute of Cancer Research & French Consortium on Cancer Evolution, London
- 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

## Referees

**Professor Niko Beerenwinkel**  
D-BSSE, ETH Zurich  
Mattenstrasse 26  
4058 Basel, Switzerland  
niko.beerenwinkel@bsse.ethz.ch  
+41 (0)61 387 31 69

**Professor Michael Hochberg**  
ISEM, University of Montpellier  
Place Eugène Bataillon  
Montpellier 34095, France  
michael.hochberg@umontpellier.fr  
+33 (0)4 67 14 34 80

**Professor Hanna Kokko**  
IEU, University of Zurich  
Winterthurerstrasse 190  
8057 Zurich, Switzerland  
hanna.kokko@ieu.uzh.ch  
+41 (0)44 635 47 40

**Dr Mario Recker**  
CME, University of Exeter  
Penryn Campus  
Penryn TR10 9FE, UK  
m.recker@exeter.ac.uk  
+44 (0)1326 259329