RICHARD J. ACTON

Professional Interests:

- · Ageing Biology
- Epigenetics
- · Adademic Software Development



EDUCATION

2019 2016-09

MRC PhD Studentship

Epigenomics of Human Ageing

University of Southampton

Including examination of changes in tRNA gene methylation with Age and performing EWAS for Bone health outcomes.

Supervisor: Dr. Chris Bell c.bell@gmul.ac.uk

2016-07 2015-09

MSc - Bioinformatics and Systems biology (Distinction)

Project: Differential Expression of circRNAs in daf-2 and daf-2/16 mutant C. elegans Neurons.

Birkbeck University of London

Supervisor: Irilenia Nobeli i.nobeli@mail.cryst.bbk.ac.uk

2015-07 2011-09

MSci - Biochemistry and Genetics (1st Hons)

Undergraduate Degree

♀ University of Nottingham

Research Projects During this period:

- 2015 CRISPR based KO and screens of B methylase in the N6methyladensine RNA modification system in Arabidopsis. (Supervisor: Dr. Rupert Fray rupert.fray@nottingham.ac.uk)
- 2014 BBSRC REP Summer project (See Grants).
- 2013 Interactions of PilZ proteins of the Cyclic-dimeric-3' → 5'-Guanosine Mono Phosphate (c-di-GMP) signalling system in Bdellovibrio bacteriovorus. (Supervisor: Prof. Liz Sockett liz.sockett@nottingham.ac.uk)
- 2012 Summer project Identifying possible chlamydial effector protein genes and expressing in Human cell lines (Supervisor: Dr. Alan Huett Alan.Huett@nottingham.ac.uk)

2011-07 2009-09

A-levels

Further education

The Sixth Form College Farnborough

Grade Subject Α Biology (A2) Chemistry (A2) Α Physics (A2) Α Maths (A2) В Economics (AS) General Studies (A2) B

CONTACT INFO

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D 0000-0002-2574-9611

前 DOB: 1992-10-30 **POB:** Scotland – UK

Programming & **Bioinformatic Skills***

R Perl5/6 •••00 Bash ••000 Python •0000 SQL •0000 HTML •0000 CSS •0000 javascript ●0000 PHP •0000 Linux Use ••••

Automated unit testing and continuous integration for R packages.

Basic Shiny app development for interactive data exploration and visualisation.

Use of High Performance Computing Clusters for 'Big Data' analysis, Experience with SLURM, PBS/TORQUE schedulers and the use of the future.batchtools R package for R processing on HPCs

2009-07 | 2005-09

GCSEs

Secondary School

♥ Frensham Heights School

Subject	Grade
Double Science	(A*)A*
Maths	A*
ICT	A*
French	A*
Photography	A*
Design Tech(RM)	A*
Art and Design 3D	A*
Geography	A*
English	Α
English Literature	В

(Photography taken as an extra-curricular in 2007, French taken 1 year early in 2008)

2005 | 1996

Early life Education

Primary Schooling

Q UK, US, UK, CH

- 2003 2005 École Internationale de Genève (La Grande Boissière campus), (Geneva, Switzerland)
- · 2000 2003 Cheddington Combined School (UK)
- 1999 2000 Osgood Elementary (Massachusetts, USA)
- 1997-1999 Cohasset Montessori (Massachusetts, USA)
- **1996-1997** Bagshot Montessori (*UK*)



PUBLICATIONS & PRESENTATIONS

Publications

- The Genomic Loci of Specific Human tRNA Genes Exhibit Ageing-Related DNA Hypermethylation (Pre-print 2019) @10.1101/870352
- Obligatory and facilitative allelic variation in the DNA methylome within common disease-associated loci, Nature Communications (2018) @10.1038/s41467-017-01586-1 (journal-article)
- Cancer detection and tissue of origin determination with novel annotation and scoring of cell-free methylated DNA, AME Medical Journal (2017) (2017) (2017)
 10.21037/amj.2017.08.02 (Comment)

Presentations

- (2019-01-16:17 Advances at the interface between metabolism and epigenetics) Poster Presentation, Specific Isoacceptor and Isodecoder tRNA Genes Exhibit Increased DNA Methylation with Age.
- **(2018-11-21 Institute of Developmental Sciences Review)** *Oral Presentation,* Ageing-Associated DNA Hypermethylation in tRNA genetic loci.
- (2018-01-31:02-02 Healthy Ageing: From Molecules to Organisms) Lightning talk & Poster Presentation, Specific Isoacceptor and Isodecoder tRNA Genes Exhibit Increased DNA Methylation with Age.
- **(2017-11-14:17 Epigenomics of Common Disease)** *Poster Presentation* 1st Poster Prize, Specific Isoacceptor and Isodecoder tRNA Genes Exhibit Increased DNA

丛 'Wet' Lab Skills

Molecular cloning (Multiple Techniques), DNA and RNA extraction and purification, PCR, Acrylamide and agarose gel electrophoresis, Basic chromatographic protein purification and hanging drop vapour diffusion crystallisation, Basic cell culture with transfection, staining and light microscopy.

Methylation with Age.

· (2017-11-08 - Institute of developmental sciences open day) Poster Presentation, Specific Isoacceptor and Isodecoder tRNA Genes Exhibit Increased DNA Methylation with Age.



CONFERENCES, MEETING & WORKSHOPS

- 2019-08-20 R Package development with Hadley Wickham (Southampton)
- 2019-08-21 Boosting your productivity using Tidyverse with Hadley Wickham (Southampton)
- · 2017-11-29 QMUL, London Epigenetic Club meeting
- 2017-06-05:07 Epigenetic Epidemiology Short Course (University of Bristol)
- 2017-03-08:10 Clinical Epigenetics Society (CLEPSO) meeting (Düsseldorf)
- 2017-03-03 London Epigenetics Club / London Chromatin Club joint meeting on Epigenetics and Metabolism
- · 2016-11-25 Research Data management Workshop (Southampton)
- · 2016-11-16 Galton institute conference (London)
- 2016-11-01:04 Epigenomics of Common Diseases (Cambridge)



■ GRANTS

2014-08 2014-06

BBSRC - Research Experience Placement grant

Project: Deletion, 2-hybrid vector creation and crystallography of the Bdellovibrio bacteriovorus Bd1483 gene.

Universities of Nottingham Birmingham Supervisors: Professor Liz Sockett liz.sockett@nottingham.ac.uk, Dr. Andy Lovering@bham.ac.uk



FOOTNOTES

Scoring system* - A subjective assessment of my own skill level. 5 is ~ daily use of the language, extensive knowledge of its ecosystem as well as deep knowledge of its structure and experience of developing production apps in that language. 1 is occasional use with heavy reliance of reference material but with at least one real world task such as creating a website or pipline accomplished using that tool.

> This resume was made with the R package pagedown.

Last updated on 2019-12-17.