Alistair Bailey

I currently work as a learning technologist supporting Librarians at the University of Southampton. 7

An engineer by training, I have since worked primarily as an informatician and research scientist investigating antigen processing and presentation by major histocompatibility molecules.

My most recent project was CRUK Accelerator: Improving immunotherapy treatment for cancer patients.² Other projects I have worked on include research into influenza, COVID19, skin sensitization to chemical allergens, asthma and contagious cancer in the Tasmanian Devil. I am also a Data and Software Carpentry³ instructor.



EDUCATION

2017 | 2017

CARPENTRIES INSTRUCTOR

Worldwide

The Carpentries

 I trained as a Carpentries⁴ instructor as part of their volunteer led mission to increase global capacity in essential data and computational skills for conducting efficient, open, and reproducible research.

2016 | 2016

MACHINE LEARNING

Stanford University

Coursera

• 10 week online introduction to machine learning.

2015 | 2015

DATA SCIENCE SPECIALIZATION

John Hopkins University

Coursera

 \bullet 12 month online set of courses on data science using R, git and command line tools.

2013 | 2008

PHD, IMMUNOLOGY

Cancer Sciences, University of Southampton

Southampton, UK

 Thesis: Relating the structure, function and dynamics of the MHC Class I antigen presenting molecule.

2008 | 2005

BENG, CIVIL ENGINEERING

University of Southampton

Southampton, UK

• First Class Honours in Civil Engineering.

2005 | 2004

ENGINEERING, SCIENCE & MATHEMATICS FOUNDATION YEAR University of Southampton ♥ Southampton, UK

Maths and physics foundation year preparation for undergraduate study.



View this CV online with links at ab604.uk/cv/cv.html

CONTACT

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- ¥ alistair604
- github.com/ab604

@ ab604.uk

1 0000-0003-0023-8679

LANGUAGE SKILLS

R
Python
Bash
SQL
Markdown
Git
Latex

Made with the R package pagedown.

The source code is available on github.com/ab604/abailey-cv.

The font is Atkinson Hyperlegible

Last updated on 2024-01-14.

1994 | 1992

BTEC ND AUDIO-VISUAL PRODUCTION

Bournemouth & Poole College of Art & Design

Page Bournemouth, UK

• Foundation course in film, photography, TV and radio production.

Τ

RESEARCH EXPERIENCE

2023 | 2018

RESEARCH FELLOW

Centre for Proteomic Research/Cancer Sciences, University of Southampton

♥ Southampton, UK

 Cancer Research UK Accelerator: this project aims to identify potential treatment targets for hard to treat cancers such as lung cancer using peptidomics methods.

In my role, I process, analyse and manage data from various Omics technologies, primarily whole exome sequencing, RNAseq and proteomics. Proteomics data I receive as Thermo raw data and process with Peaks Studio⁵, and post-process in R and RStudio. Whole exome and transcriptomics data I receive as fastq files and I use a mixture of command line tools using bash scripts and R and RStudio. I tend to follow the Broad Institute Best Practices for genomic data analysis⁶ and Cornell Bioinformatics Core⁷ for transcriptomic data processing. Scripts and processed data are managed using git version control. Raw data is backed up remotely and deposited along with processed outputs public repositories such as EBI PRIDE⁸ and the European Phenome-Genome Archive⁹ following FAIR protocols¹⁰. My primary computer is a Linux Ubuntu machine, but I also use Windows.

- We have also developed our method to identify treatment targets for infectious diseases such as influenza.
- In 2020 I also worked to develop a COVID19 test using proteomics methods.

2018 | 2015

RESEARCH FELLOW

Centre for Proteomic Research/Cancer Sciences, University of Southampton

Southampton, UK

 Developed peptidomics methodology at the UoS for research into the role of MHC molecules in skin sensitisation to chemical allergy.

2015 | 2013

RESEARCH FELLOW

Cancer Sciences, University of Southampton

Southampton, UK

• MRC Centenary Fellow



INDUSTRY EXPERIENCE

2012 | 2012

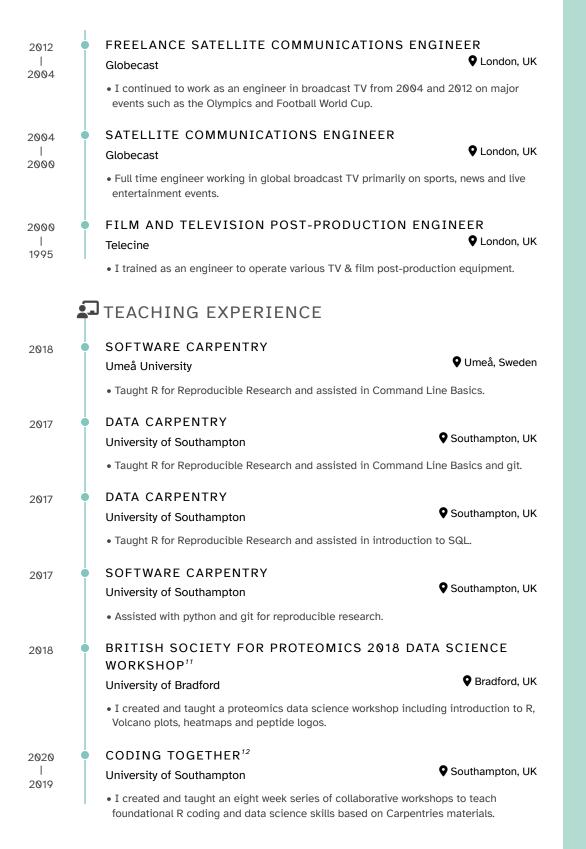
INTERNSHIP

Microsoft Research

Cambridge, UK

• Helped develop computational model of MHC I peptide selection.

I have worked in a variety of roles ranging from engineering to research scientist. I like collaborative environments where I can learn from my peers.



I am passionate about teaching foundational coding and data science skills to researchers and developing evidence-based best practices. I am especially interested in helping novices and making coding more accessible to all.

2019 | 2018

ACADEMIC SUPPORT TUTOR

IntoUniversity Millbrook

Southampton, UK

• IntoUniversity¹³ supports young people from disadvantaged backgrounds to attain either a university place or another chosen aspiration. I volunteered as an academic support tutor for secondary school learners.



PUBLICATIONS

2022

IDENTIFICATION OF NEOANTIGENS IN ESOPHAGEAL ADENOCARCINOMA14

Immunology

• Ben Nicholas, Alistair Bailey, Katy J. McCann, Oliver Wood, Robert C. Walker, Robert Parker, Nicola Ternette, Tim Elliott, Tim J. Underwood, Peter Johnson, Paul Skipp

2022

ANALYSIS OF CELL-SPECIFIC PERIPHERAL BLOOD BIOMARKERS IN SEVERE ALLERGIC ASTHMA IDENTIFIES INNATE IMMUNE DYSFUNCTION15

Clinical & Experimental Allergy

• Ben Nicholas, Jane Guo, Hyun-Hee Lee, Alistair Bailey, Rene de Waal Malefyt, Milenko Cicmil, Ratko Djukanovic

2021

THE DIFFERENTIATION STATE OF THE SCHWANN CELL PROGENITOR DRIVES PHENOTYPIC VARIATION BETWEEN TWO CONTAGIOUS CANCERS¹⁶

PLOS Pathogens

• Rachel S. Owen, Sri H. Ramarathinam, Alistair Bailey, Annalisa Gastaldello, Kathryn Hussey, Paul J. Skipp, Anthony W. Purcell, Hannah V. Siddle

2021

CHARACTERIZATION OF THE CLASS I MHC PEPTIDOME RESULTING FROM DNCB EXPOSURE OF HACAT CELLS¹⁷

Toxicological Sciences

• Alistair Bailey, Ben Nicholas, Rachel Darley, Erika Parkinson, Ying Teo, Maja Aleksic, Gavin Maxwell, Tim Elliott, Michael Ardern-Jones, Paul Skipp.

2022

IMMUNOPEPTIDOMIC ANALYSIS OF INFLUENZA A VIRUS INFECTED HUMAN TISSUES IDENTIFIES INTERNAL PROTEINS AS A RICH SOURCE OF HLA LIGANDS18

PLoS Pathogens

• Ben Nicholas, Alistair Bailey, Karl J. Staples, Tom Wilkinson, Tim Elliott, Paul Skipp.

2021

THE IMMUNOPEPTIDOMES OF TWO TRANSMISSIBLE CANCERS AND THEIR HOST HAVE A COMMON, DOMINANT PEPTIDE MOTIF¹⁹

Immunology

• Annalisa Gastaldello, Sri H. Ramarathinam, Alistair Bailey, Rachel Owen, Steven Turner, N. Kontouli, Tim Elliott, Paul Skipp, Anthony W. Purcell, Hannah V. Siddle. DYNAMICALLY DRIVEN ALLOSTERY IN MHC PROTEINS:
PEPTIDE-DEPENDENT TUNING OF CLASS I MHC GLOBAL
FLEXIBILITY²⁰

Frontiers in Immunology

• Cory M. Ayres, Esam T. Abualrous, Alistair Bailey, Christian Abraham, Lance M. Hellman, Steven A. Corcelli, Frank Noé, Tim Elliott, Brian M. Baker.

DIRECT EVIDENCE FOR CONFORMATIONAL DYNAMICS IN MAJOR HISTOCOMPATIBILITY COMPLEX CLASS I MOLECULES²¹

JBC

• Andy van Hateren, Malcolm Anderson, Alistair Bailey, Jörn M. Werner, Paul Skipp, Tim Flliott

RECENT ADVANCES IN MAJOR HISTOCOMPATIBILITY
COMPLEX CLASS I ANTIGEN PRESENTATION: PLASTIC MHC
MOLECULES AND TAPBPR MEDIATED QUALITY CONTROL²²
F1000 Research

• Andy van Hateren, Alistair Bailey, Tim Elliott.

SELECTOR FUNCTION OF MHC I MOLECULES IS DETERMINED BY PROTEIN PLASTICITY²³

Scientific Reports

 Alistair Bailey, Neil Dalchau, Rachel Carter, Stephen Emmott, Andrew Phillips, Jörn M. Werner, Tim Elliott

TWO POLYMORPHISMS FACILITATE DIFFERENCES IN PLASTICITY BETWEEN TWO CHICKEN MAJOR HISTOCOMPATIBILITY COMPLEX CLASS I PROTEINS²⁴ PLoS One

• Alistair Bailey, Andy van Hateren, Tim Elliott, Jörn M. Werner.

A MECHANISTIC BASIS FOR THE CO-EVOLUTION OF CHICKEN TAPASIN AND MAJOR HISTOCOMPATIBILITY COMPLEX CLASS I PROTEINS²⁵

JBC

• Andy van Hateren, Rachel Carter, Alistair Bailey, Nasia Kontouli, Anthony P. Williams, Jim Kaufman, Tim Elliott.

THE CELL BIOLOGY OF MAJOR HISTOCOMPATIBILITY COMPLEX CLASS I ASSEMBLY: TOWARDS A MOLECULAR UNDERSTANDING²⁶

Tissue Antigens

• A. Van Hateren, E. James, A. Bailey, A. Phillips, N. Dalchau, T. Elliott

2015

2017

2017

2014

2013

2010

@ LINKS

- 1. https://www.soton.ac.uk
- 2. https://www.cancerresearchuk.org/funding-for-researchers/accelerator-award/portfolio-funded -projects-outputs
- 3. https://carpentries.org/
- 4. https://carpentries.org/
- 5. https://www.bioinfor.com/peaks-studio/
- 6. https://gatk.broadinstitute.org/hc/en-us
- 7. https://abc.med.cornell.edu/
- 8. https://www.ebi.ac.uk/pride/
- 9. https://ega-archive.org/
- 10. https://www.go-fair.org/fair-principles/
- 11. https://ab604.github.io/docs/bspr_workshop_2018/index.html
- 12. https://ab604.github.io/docs/coding-together-2019/
- 13. https://intouniversity.org/
- 14. https://doi.org/10.1111/imm.13578
- 15. https://doi.org/10.1111/cea.14197
- 16. https://journals.plos.org/plospathogens/article?id=10.1371/journal.ppat.1010033
- 17. https://doi.org/10.1093/toxsci/kfaa184
- 18. https://doi.org/10.1371/journal.ppat.1009894
- 19. https://doi.org/10.1111/imm.13307
- 20. https://doi.org/10.3389/fimmu.2019.00966
- 21. https://doi.org/10.1074/jbc.M117.809624
- 22. https://doi.org/10.12688/f1000research.10474.1
- 23. https://doi.org/10.1038/srep14928
- 24. https://doi.org/10.1371/journal.pone.0089657
- 25. https://doi.org/10.1074/jbc.M113.474031
- 26. https://doi.org/10.1111/j.1399-0039.2010.01550.x