ALISTAIR BAILEY

An engineer by training, I have since worked primarily as an informatician and research scientist. I am currently a Research Fellow in Microfluidic Hydrogen-Deuterium Exchange at the University of Southampton.¹

The topic of my research career has been the role of HLA antigen processing and presentation in disease recognition by T cells. In cancer² this has focused on HLApresented tumour antigens3, and in infectious disease the focus has been HLApresented viral⁴ and bacterial antigens. Exploiting these targets has the potential for enhancing personalised therapies, vaccine development and understanding allergy.

I have contributed to research into COVID19⁵, skin sensitization to chemical allergens⁶, asthma⁷ and contagious cancer in the Tasmanian Devil⁸.

My core skills are processing and analysing data from whole exome sequencing, RNAseq, scRNAseq and proteomics assays. My workflow combines command line tools with micromamba, the R programming language and git version control.

Proteomics data I have curated, deposited and I am the data controller for is deposited at the PRoteomics IDEntifications Archive9. Whole Exome and RNAseq data I have curated, deposited and I am the data controller for is deposited at the European Genome-phenome Archive¹⁰.

I am a Data and Software Carpentry¹¹ instructor and I have also created and delivered my own workshops to teach foundational R coding and data science skills¹² to bioscientists and web design¹³ to librarians.



A EDUCATION

2017

CARPENTRIES INSTRUCTOR

Worldwide

The Carpentries

• I trained as a Carpentries¹⁴ 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

MACHINE LEARNING

Stanford University

Coursera

• 10 week online introduction to machine learning.

2015

DATA SCIENCE SPECIALIZATION

John Hopkins University

Coursera

• 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.



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

CONTACT

- ab604@soton.ac.uk
- @ ab604.uk
- **D** 0000-0003-0023-8679
- github.com/ab604
- **a** @ab604.uk

LANGUAGE SKILLS

R
Bash
Markdown
Latex
Git
Python
SQL

Made with the R package pagedown.

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

The fonts are Inter and Permanent Marker

Last updated on 2025-05-13.

2005 2004

ENGINEERING, SCIENCE + MATHEMATICS FOUNDATION YEAR

University of Southampton

Southampton, UK

• Maths and physics foundation year preparation for undergraduate study.

1994 1992

BTEC ND AUDIO-VISUAL PRODUCTION

Bournemouth & Poole College of Art & Design

Pournemouth, UK

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

RESEARCH EXPERIENCE

2025

RESEARCH FELLOW

Cancer Sciences, University of Southampton

Southampton, UK

· Research Fellow in Microfluidic HDX

2023

RESEARCH FELLOW

School of Biological Sciences, University of Southampton

Southampton, UK

• scRNAseq analysis of T-cell response to neutrophil exposure. Bioinformatician maternity leave cover for Medical Research Council funded project.

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 using multi-omics methods. In this project our focus was on oesophageal, lung and neuroendocrine cancers.

As an informatician I processed, analysed and managed data from whole exome sequencing, RNAseq, scRNAseq and proteomics.

For sequencing fastq data, my workflow comprised of a mixture of command line tools using bash scripts and R/RStudio. I followed the Broad Institute Best Practices for genomic data analysis¹⁵ and Cornell Bioinformatics Core¹⁶. For proteomics data, my worklfow used Peaks Studio¹⁷, and post-process in R and RStudio.

Scripts and processed data were managed using git version control. Raw data was deposited along with processed outputs in PRoteomics IDEntifications Archive¹⁸ and the European Phenome-Genome Archive¹⁹.

We also developed our method to identify treatment targets for infectious diseases from influenza and bacterial proteins. In 2020 I also worked to develop a COVID19 test using proteomics methods.

RESEARCH FELLOW 2018 Southampton, UK Centre for Proteomic Research/Cancer Sciences, 2015 University of Southampton • Developed peptidomics methodology at the UoS for research into the role of MHC molecules in skin sensitisation to chemical allergy. 2015 RESEARCH FELLOW Southampton, UK Cancer Sciences, University of Southampton 2013 MRC Centenary Fellow TEACHING EXPERIENCE 2024 WEBPAGE DESIGNED Southampton, UK University of Southampton • I created a webpage design workshop and materials for Librarians at the University of Southampton 2022 BIOL 2013: INTRODUCTION TO BIGINFORMATICS Southampton, UK University of Southampton 2021 • I taught the undegraduate introduction to bioinformatics module on variant discovery using the University Galaxy Server. 2020 CODING TOGETHER Southampton, UK University of Southampton 2019 • I created and taught an eight week series of collaborative workshops to teach foundational R coding and data science skills based on Carpentries materials. ACADEMIC SUPPORT TUTOR 2019 Southampton, UK IntoUniversity Millbrook 2018 • IntoUniversity²² 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. 2018 SOFTWARE CARPENTRY Umeå, Sweden Umeå University • Taught R for Reproducible Research and assisted in Command Line Basics. BRITISH SOCIETY FOR PROTEOMICS ZOIR DATA SCIENCE WORKSHOP 25 2018 Paradford, UK University of Bradford • I created and taught a proteomics data science workshop including

introduction to R, Volcano plots, heatmaps and peptide logos.

I enjoy 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. 2017

DATA CARPENTRY

University of Southampton

Southampton, UK

 Taught R for Reproducible Research and assisted in Command Line Basics and git.

2017

DATA CARPENTRY

University of Southampton

Southampton, UK

• Taught R for Reproducible Research and assisted in introduction to SQL.

2017

SOFTWARE CARPENTRY

University of Southampton

Southampton, UK

• Assisted with python and git for reproducible research.



RESEARCH DATA

Immunopeptidomic analysis of influenza A virus infected human tissues identifies internal proteins as a rich source of HLA ligands²⁴, Publicly released

Proteomics data: PRIDE Project PXD022884²⁵

Dentification of Negantigens in Esophageal Adenocarcinoma²²,

Publicly released

- Proteomics data: PRIDE Project ID PXD031108²⁷
- WES & RNAseq data EGA Study ID EGAS00001005957

CHARACTERIZATION OF THE CLASS I MHC PEPTIDOME RESULTING FROM DNCB EXPOSURE OF HACAT CELLS²⁷, Publicly released

• Proteomics data: PRIDE Project PXD021373²⁹

NEGANTIGEN IDENTIFICATION IN PANCREATIC NEUROENDOCRINE TUMOURS, UNRELEASED PENDING PUBLICATION

- Proteomics data: PRIDE Project ID PXD037449
- WES & RNAseq data EGA Study ID EGAS00001006722

Immunopertidomics guided identification of Neoantigens in Non-Small cell lung cancer, Unreleased Pending Publication

- Proteomics data: PRIDE Project ID PXD028990
- WES & RNAseg data EGA Study ID EGAS00001005499

Immunopertidomics of a brain tumour cell line to identify HLA presented Zika, Unreleased Pending publication

• Proteomics data: PRIDE Project ID PXD037627

Non-small cell lung cancer global proteomics, Unreleased pending publication

• Proteomics data: PRIDE Project ID PXD054390

GESOPHAGEAL ADENOCARCINOMA GLOBAL PROTEOMES, UNRELEASED PENDING PUBLICATION

• Proteomics data: PRIDE Project ID PXD054428

INDUSTRY EXPERIENCE

2012 • INTERNSHIP

Microsoft Research

♥ Cambridge, UK

• Helped develop computational model of MHC I peptide selection.

Freelance Satellite Communications Engineer

Globecast

♥ London, UK

• I continued to work as an engineer in broadcast TV from 2004 and 2012 on major events such as the Olympics and Football World Cup.

SATELLITE COMMUNICATIONS ENGINEER

Globecast

◆ London, UK

 Full time engineer working in global broadcast TV primarily on sports, news and live entertainment events.

FILM AND TELEVISION POST-PRODUCTION ENGINEER

Telecine

◆ London, UK

• I trained as an engineer to operate various TV & film post-production equipment.

E PUBLICATIONS

EVIDENCE OF FOCUSING THE MHC CLASS I IMMUNOPEPTIDOME BY

Frontiers in Immunology

• Rachel Darley, Patricia T. Illing, Patrick Duriez, Alistair Bailey, Anthony W. Purcell, Andy van Hateren, Tim Elliott.

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.

l

2012

2004

2004

2000

2000 | 1995

2025

COMPARATIVE ANALYSIS OF PROTEIN EXPRESSION BETWEEN DESOPHAGEAL 2025 ADENOCARCINOMA AND NORMAL ADJACENT TISSUESI PLOS One • Ben Nicholas, Alistair Bailey, Katy J. McCann, Robert C. Walker, Peter Johnson, Tim Elliott, Tim J. Underwood, Paul Skipp 2025 COMPARATIVE ANALYSIS OF TRANSCRIPTOMIC AND PROTEOMIC EXPRESSION BETWEEN TWO NON-SMALL CELL LUNG CANCER SUBTYPES 52 Journal of Proteome Research • Ben Nicholas, Alistair Bailey, Katy J McCann, Peter Johnson, Tim Elliott, Christian Ottensmeier and Paul Skipp PROTEOGENOMICS GUIDED IDENTIFICATION OF FUNCTIONAL NEGANTIGENS 2024 IN NON-SMALL CELL LUNG CANCERSS bioRxiv • Ben Nicholas, Alistair Bailey, Katy J McCann, Oliver Wood, Eve Currall, Peter Johnson, Tim Elliott, Christian Ottensmeier, Paul Skipp OPERATION MOONSHOT: RAPID TRANSLATION OF A SARS-COV-Z 2022 TARGETED PEPTIDE IMMUNOAFFINITY LIQUID CHROMATOGRAPHY-TANDEM MASS SPECTROMETRY TEST FROM RESEARCH INTO ROUTINE CLINICAL USESY Clinical Chemistry and Laboratory Medicine • Jenny Hällqvist, Benjamin I. Nicholas, Alistair Bailey et al. Identification of negantigens in esophageal adenocarcinomass 2022 **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 DYSFUNCTIONS Clinical & Experimental Allergy • Ben Nicholas, Jane Guo, Hyun-Hee Lee, Alistair Bailey, Rene de Waal Malefyt, Milenko Cicmil, Ratko Djukanovic

Immunopeptidomic analysis of influenza A virus infected human tissues identifies internal proteins as a rich source of HLA ligands⁵⁷

PLoS Pathogens

2022

 Ben Nicholas, Alistair Bailey, Karl J. Staples, Tom Wilkinson, Tim Elliott, Paul Skipp. 2021 THE DIFFERENTIATION STATE OF THE SCHWANN CELL PROGENITOR DRIVES PHENOTYPIC VARIATION BETWEEN TWO CONTAGIOUS CANCERSS?

PLOS Pathogens

2021

2021

2019

2017

2017

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

CHARACTERIZATION OF THE CLASS I MHC PEPTIDOME RESULTING FROM DNCB EXPOSURE OF HACAT CELLS⁵⁷

Toxicological Sciences

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

THE IMMUNOPEPTIDOMES OF TWO TRANSMISSIBLE CANCERS AND THEIR HOST HAVE A COMMON, DOMINANT PEPTIDE MOTIF¹⁰

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 FLEXIBILITY"

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 $\mathbf I$ molecules⁴²

JBC

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

RECENT ADVANCES IN MAJOR HISTOCOMPATIBILITY COMPLEX CLASS I ANTIGEN PRESENTATION: PLASTIC MHC MOLECULES AND TAPBPR MEDIATED QUALITY CONTROL*5

F1000 Research

• Andy van Hateren, Alistair Bailey, Tim Elliott.

2015 SELECTOR FUNCTION OF MHC I MOLECULES IS DETERMINED BY PROTEIN PLASTICITY"

Scientific Reports

 Alistair Bailey, Neil Dalchau, Rachel Carter, Stephen Emmott, Andrew Phillips, Jörn M. Werner, Tim Elliott 2014 TWO POLYMORPHISMS FACILITATE DIFFERENCES IN PLASTICITY BETWEEN
TWO CHICKEN MAJOR HISTOCOMPATIBILITY COMPLEX CLASS I
PROTEINS"S

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 PROTEINS"

JBC

2013

2010

• 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 understanding" $^{\circ}$

Tissue Antigens

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



- 1. https://www.soton.ac.uk
- 2. https://www.cancerresearchuk.org/funding-for-researchers/accelerator-award /portfolio-funded-projects-outputs
- 3. https://doi.org/10.1111/imm.13578
- 4. https://doi.org/10.1371/journal.ppat.1009894
- 5. https://doi.org/10.1515/cclm-2022-1000
- 6. https://doi.org/10.1093/toxsci/kfaa184
- 7. https://doi.org/10.1111/cea.14197
- 8. https://doi.org/10.1111/imm.13307
- 9. https://www.ebi.ac.uk/pride/
- 10. https://ega-archive.org/
- 11. https://carpentries.org/
- 12. https://ab604.github.io/docs/coding-together-2019/
- 13. https://ab604.github.io/webpage-design/
- 14. https://carpentries.org/
- 15. https://gatk.broadinstitute.org/hc/en-us
- 16. https://abc.med.cornell.edu/
- 17. https://www.bioinfor.com/peaks-studio/
- 18. https://www.ebi.ac.uk/pride/
- 19. https://ega-archive.org/
- 20. https://ab604.github.io/webpage-design/
- 21. https://ab604.github.io/docs/coding-together-2019/
- 22. https://intouniversity.org/
- 23. https://ab604.github.io/docs/bspr_workshop_2018/index.html
- 24. https://doi.org/10.1371/journal.ppat.1009894
- 25. https://www.ebi.ac.uk/pride/archive/projects/PXD022884
- 26. https://doi.org/10.1111/imm.13578
- 27. https://www.ebi.ac.uk/pride/archive/projects/PXD031108

- 28. https://doi.org/10.1093/toxsci/kfaa184
- 29. https://www.ebi.ac.uk/pride/archive/projects/PXD021373
- 30. 10.3389/fimmu.2025.1563789
- 31. 10.1371/journal.pone.0318572
- 32. https://doi.org/10.1021/acs.jproteome.4c00773
- 33. https://doi.org/10.1101/2024.05.30.596609
- 34. https://doi.org/10.1515/cclm-2022-1000
- 35. https://doi.org/10.1111/imm.13578
- 36. https://doi.org/10.1111/cea.14197
- 37. https://doi.org/10.1371/journal.ppat.1009894
- 38. https://journals.plos.org/plospathogens/article?id=10.1371/journal.ppat.1010033
- 39. https://doi.org/10.1093/toxsci/kfaa184
- 40. https://doi.org/10.1111/imm.13307
- 41. https://doi.org/10.3389/fimmu.2019.00966
- 42. https://doi.org/10.1074/jbc.M117.809624
- 43. https://doi.org/10.12688/f1000research.10474.1
- 44. https://doi.org/10.1038/srep14928
- 45. https://doi.org/10.1371/journal.pone.0089657
- 46. https://doi.org/10.1074/jbc.M113.474031
- 47. https://doi.org/10.1111/j.1399-0039.2010.01550.x