Alistair Bailey

An engineer by training, I have since worked primarily as an informatician and research scientist investigating antigen processing and presentation by major histocompatibility molecules using multi-omics methods. I currently work as a learning technologist supporting Librarians at the University of Southampton.¹

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 and I have also created and delivered my own workshops to teach foundational R coding and data science skills to bioscientists.

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



EDUCATION

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

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.



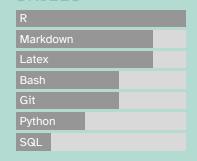
View this CV online with links at ab604.uk/cv/cv.html

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github.com/ab604



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 2024-02-10.

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

Bournemouth, UK

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



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



RESEARCH DATA

IMMUNOPEPTIDOMIC ANALYSIS OF INFLUENZA A VIRUS INFECTED HUMAN TISSUES IDENTIFIES INTERNAL PROTEINS AS A RICH SOURCE OF HLA LIGANDS11, PUBLICLY RELEASED

• Proteomics data: PRIDE Project PXD022884¹²

IDENTIFICATION OF NEOANTIGENS IN ESOPHAGEAL ADENOCARCINOMA13, PUBLICLY RELEASED

- Proteomics data: PRIDE Project ID PXD03110814
- WES & RNAseq data EGA Study ID EGAS00001005957

CHARACTERIZATION OF THE CLASS I MHC PEPTIDOME RESULTING FROM DNCB EXPOSURE OF HACAT CELLS¹⁵. PUBLICLY RELEASED

Proteomics data: PRIDE Project PXD021373¹⁶

NEOANTIGEN IDENTIFICATION IN PANCREATIC NEUROENDOCRINE TUMOURS, UNRELEASED PENDING **PUBLICATION**

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

IMMUNOPEPTIDOMICS GUIDED IDENTIFICATION OF NEOANTIGENS IN NON-SMALL CELL LUNG CANCER, UNRELEASED PENDING PUBLICATION

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

IMMUNOPEPTIDOMICS OF A BRAIN TUMOUR CELL LINE TO IDENTIFY HLA PRESENTED ZIKA, UNRELEASED PENDING **PUBLICATION**

• Proteomics data: PRIDE Project ID PXD037627



INDUSTRY EXPERIENCE

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 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 SOFTWARE CARPENTRY

University of Southampton

Southampton, UK

• Assisted with python and git for reproducible research.

PUBLICATIONS

2022

OPERATION MOONSHOT: RAPID TRANSLATION OF A SARS-COV-2 TARGETED PEPTIDE IMMUNOAFFINITY LIQUID CHROMATOGRAPHY-TANDEM MASS SPECTROMETRY TEST FROM RESEARCH INTO ROUTINE CLINICAL USE²⁰

Clinical Chemistry and Laboratory Medicine

• Jenny Hällqvist, Benjamin I. Nicholas, Alistair Bailey et al.

2022

IDENTIFICATION OF NEOANTIGENS IN ESOPHAGEAL ADENOCARCINOMA²¹

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 DYSFUNCTION²²

Clinical & Experimental Allergy

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

2022

IMMUNOPEPTIDOMIC ANALYSIS OF INFLUENZA A VIRUS INFECTED HUMAN TISSUES IDENTIFIES INTERNAL PROTEINS AS A RICH SOURCE OF HLA LIGANDS²³ **PLoS Pathogens**

• 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 CANCERS24

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 CELLS²⁵

Toxicological Sciences

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

2021

THE IMMUNOPEPTIDOMES OF TWO TRANSMISSIBLE CANCERS AND THEIR HOST HAVE A COMMON, DOMINANT PEPTIDE MOTIF²⁶

Immunology

 Annalisa Gastaldello, Sri H. Ramarathinam, Alistair Bailey, Rachel Owen, Steven Turner, N. Kontouli, Tim Elliott, Paul Skipp, Anthony W. Purcell, Hannah V. Siddle.

2019

DYNAMICALLY DRIVEN ALLOSTERY IN MHC PROTEINS: PEPTIDE-DEPENDENT TUNING OF CLASS I MHC GLOBAL FLEXIBILITY²⁷

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.

2017

DIRECT EVIDENCE FOR CONFORMATIONAL DYNAMICS IN MAJOR HISTOCOMPATIBILITY COMPLEX CLASS I MOLECULES²⁸

JBC

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

2017

RECENT ADVANCES IN MAJOR HISTOCOMPATIBILITY
COMPLEX CLASS I ANTIGEN PRESENTATION: PLASTIC MHC
MOLECULES AND TAPBPR MEDIATED QUALITY CONTROL²⁹

F1000 Research

• Andy van Hateren, Alistair Bailey, Tim Elliott.

2015

SELECTOR FUNCTION OF MHC I MOLECULES IS DETERMINED BY PROTEIN PLASTICITY³⁰

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 PROTEINS³¹

PLoS One

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

2013

A MECHANISTIC BASIS FOR THE CO-EVOLUTION OF CHICKEN TAPASIN AND MAJOR HISTOCOMPATIBILITY COMPLEX CLASS I PROTEINS³²

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 UNDERSTANDING³³

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://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://doi.org/10.1371/journal.ppat.1009894
- 12. https://www.ebi.ac.uk/pride/archive/projects/PXD022884
- 13. https://doi.org/10.1111/imm.13578
- 14. https://www.ebi.ac.uk/pride/archive/projects/PXD031108
- 15. https://doi.org/10.1093/toxsci/kfaa184
- 16. https://www.ebi.ac.uk/pride/archive/projects/PXD021373
- 17. https://ab604.github.io/docs/coding-together-2019/
- 18. https://intouniversity.org/
- 19. https://ab604.github.io/docs/bspr_workshop_2018/index.html
- 20. https://doi.org/10.1515/cclm-2022-1000
- 21. https://doi.org/10.1111/imm.13578
- 22• https://doi.org/10.1111/cea.14197
- 23. https://doi.org/10.1371/journal.ppat.1009894
- 24. https://journals.plos.org/plospathogens/article?id=10.1371/journal.ppat.1010033
- 25. https://doi.org/10.1093/toxsci/kfaa184
- 26. https://doi.org/10.1111/imm.13307
- 27. https://doi.org/10.3389/fimmu.2019.00966
- 28. https://doi.org/10.1074/jbc.M117.809624
- 29. https://doi.org/10.12688/f1000research.10474.1
- 30. https://doi.org/10.1038/srep14928
- 31. https://doi.org/10.1371/journal.pone.0089657
- 32. https://doi.org/10.1074/jbc.M113.474031
- 33• https://doi.org/10.1111/j.1399-0039.2010.01550.x