Alistair Bailey

I am a researcher at the Centre for Proteomic Research and Cancer Sciences Unit at the University of Southampton. My research interests are in antigen processing and presentation by class I MHC molecules, data science and Omics. My role is primarily as an informatician. The project I currently work on aims to improve immunotherapy treatment for cancer patients. I also contribute to research into the role of MHC molecules in skin sensitization to chemical allergens, and contagious cancer in the Tasmanian Devil. I am also a Data and Software Carpentry instructor.





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

CONTACT

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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 2022-09-25.

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.



🖵 RESEARCH EXPERIENCE

Current | 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 Studio⁴, 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 analysis⁵ and Cornell Bioinformatics Core⁶ 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 PRIDE⁷ and the European Phenome-Genome Archive⁸ following FAIR protocols⁹. 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

Ocambridge, 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.

2017 | 2017

SOFTWARE CARPENTRY

University of Southampton

• Assisted with python and git for reproducible research.

Southampton, UK

PUBLICATIONS

2022 | 2022 IDENTIFICATION OF NEOANTIGENS IN ESOPHAGEAL ADENOCARCINOMA 13

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 | 2022 ANALYSIS OF CELL-SPECIFIC PERIPHERAL BLOOD BIOMARKERS IN SEVERE ALLERGIC ASTHMA IDENTIFIES INNATE IMMUNE DYSFUNCTION¹⁴

Clinical & Experimental Allergy

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

2022 | 2022 IMMUNOPEPTIDOMIC ANALYSIS OF INFLUENZA A VIRUS INFECTED HUMAN TISSUES IDENTIFIES INTERNAL PROTEINS AS A RICH SOURCE OF HLA LIGANDS¹⁵

PLoS Pathogens

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

2021 | 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 | 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.

2021 | 2021 THE IMMUNOPEPTIDOMES OF TWO TRANSMISSIBLE CANCERS AND THEIR HOST HAVE A COMMON, DOMINANT PEPTIDE MOTIF 18

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: 2019 PEPTIDE-DEPENDENT TUNING OF CLASS I MHC GLOBAL 2019 FLEXIBILITY19 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 2017 MAJOR HISTOCOMPATIBILITY COMPLEX CLASS I 2017 MOLECULES²⁰ JBC • Andy van Hateren, Malcolm Anderson, Alistair Bailey, Jörn M. Werner, Paul Skipp, Tim RECENT ADVANCES IN MAJOR HISTOCOMPATIBILITY 2017 COMPLEX CLASS I ANTIGEN PRESENTATION: PLASTIC MHC 2017 MOLECULES AND TAPBPR MEDIATED QUALITY CONTROL²¹ F1000 Research • Andy van Hateren, Alistair Bailey, Tim Elliott. SELECTOR FUNCTION OF MHC I MOLECULES IS DETERMINED 2015 BY PROTEIN PLASTICITY²² 2015 Scientific Reports • Alistair Bailey, Neil Dalchau, Rachel Carter, Stephen Emmott, Andrew Phillips, Jörn M. Werner, Tim Elliott TWO POLYMORPHISMS FACILITATE DIFFERENCES IN 2014 PLASTICITY BETWEEN TWO CHICKEN MAJOR 2014 HISTOCOMPATIBILITY COMPLEX CLASS I PROTEINS²³ PLoS One • Alistair Bailey, Andy van Hateren, Tim Elliott, Jörn M. Werner. A MECHANISTIC BASIS FOR THE CO-EVOLUTION OF 2013 CHICKEN TAPASIN AND MAJOR HISTOCOMPATIBILITY 2013 COMPLEX CLASS I PROTEINS²⁴ JBC • Andy van Hateren, Rachel Carter, Alistair Bailey, Nasia Kontouli, Anthony P. Williams, Jim Kaufman, Tim Elliott. THE CELL BIOLOGY OF MAJOR HISTOCOMPATIBILITY 2010 COMPLEX CLASS I ASSEMBLY: TOWARDS A MOLECULAR 2010 UNDERSTANDING²⁵

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

Tissue Antigens



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