

# Dr. Mark Norman Read

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## Education

<b>Ph.D. in Computer Science</b> Computer Science, University of York, UK	Oct 2007 - Feb 2012
<b>MEng in Computer Science &amp; Systems Engineering</b> <i>1<sup>st</sup> Class Honours</i> Computer Science, University of York, UK	Oct 2002 - Oct 2007
<b>Graduate Certificate in Educational Studies (Higher Education)</b> <i>High Distinction.</i> The University of Sydney, Australia	2020

## Employment History

<b>Lecturer in Digital Health (Health Informatics)</b> School of Computer Science, University of Sydney	Sept 2019 - current
<b>Research Fellow</b> School of Chemical and Biomolecular Engineering, University of Sydney	Sept 2018 - Aug 2019
<b>Research Fellow</b> School of Life and Environmental Sciences and the Charles Perkins Centre, University of Sydney	Sept 2013 - Sept 2018
<b>Research Associate</b> Electronics Engineering, University of York	April 2011 - August 2013

## Research and Scholarship

I conduct research in Health Informatics, devising novel data science and computational modelling methods that advance biomedical discovery and healthcare delivery. I establish and lead collaborative projects with clinicians and experimentalists to maximise the translational benefit of my work.

I have three areas of principle focus:

- Using computational modelling and data science to understand the impact of gut microbial systems on our health
- Developing new computational algorithms and models to study the immune system through high-throughput molecular profiling
- Innovative computational modelling and analytical methodologies to investigate collective motion: how mobile agents achieve higher-level goals through coordination and communication. E.g. analysing how immune cells hunt for cancers and pathogens.

I have published 36 full-length peer-reviewed manuscripts, and have attracted over \$850,000 in competitive funding to support my research.

## Publications

Authorship conventions differ across the disciplines I publish in, and I adhere to those of the given journal/-conference. In Computer Science relative contribution is indicated by authorship order. In Medicine and Biology, last authorship positions and corresponding author status are highly esteemed: they denote seniority in conceiving, convening and managing the project.

### Refereed Journal Articles (\* corresponding author)

- J31 *JP Molina Ortiz, MN Read, DD McClure, A Holmes, F Deghani, ER Shanahan.* (2022). High throughput genome scale modeling predicts microbial vitamin requirements contribute to gut microbiome community structure. **Gut Microbes** 14(1):2118831.
- J30 *Rebecca C Simpson, Erin R Shanahan, Marcel Batten, Irene LM Reijers, Mark Read, Ines P Silva, Judith M Versluis, Rosilene Ribeiro, Alexandra S Angelatos, Jian Tan, Chandra Adhikari, Alexander M Menzies, Robyn PM Saw, Maria Gonzalez, Kerwin F Shannon, Andrew J Spillane, Rebecca Velickovic, Alexander J Lazar, Ashish V Damania, Aditya K Mishra, Manoj Chelvanambi, Anik Banerjee, Nadim J Ajami, Jennifer A Wargo, Laurence Macia, Andrew J Holmes, James S Wilmott, Christian U Blank, Richard A Scolyer, Georgina V Long.* (2022). Diet-driven microbial ecology underpins associations between cancer immunotherapy outcomes and the gut microbiome. **Nature Medicine** 28(11):2344-2352.
- J29 *Jian Tan, Duan Ni, Jemma Taitz, Gabriela Veronica Pinget, Mark Read, Alistair Senior, Jibran Abdul Wali, Reem Elnour, Erin Shanahan, Huiling Wu, Steven J Chadban, Ralph Nanan, Nicholas Jonathan Cole King, Georges Emile Grau, Stephen J Simpson, Laurence Macia.* (2022). Dietary protein increases T-cell-independent sIgA production through changes in gut microbiota-derived extracellular vesicles. **Nature Communications**, 13(1):4336.
- J28 *Givanna H Putri, Jonathan Chung, Davis N Edwards, Felix Marsh-Wakefield, Irena Koprinska, Suat Dervish, Nicholas JC King, Thomas M Ashhurst, Mark N Read \*.* (2022). TrackSOM: Mapping immune response dynamics through clustering of time-course cytometry data **Cytometry Part A**, 103(1):54-70.
- J27 *Felix Marsh-Wakefield, Pierre Juillard, Thomas M Ashhurst, Annette Juillard, Diana Shinko, Givanna H Putri, Mark N Read, Helen M McGuire, Scott N Byrne, Simon Hawke, Georges E Grau.* (2022). Peripheral B-cell dysregulation is associated with relapse after long-term quiescence in patients with multiple sclerosis. **Immunology and Cell Biology**, 100(6):543-467.
- J26 *GH Putri, I Koprinska, TM Ashhurst, NJC King, MN Read \*.* (2021). Using single-cell cytometry to illustrate integrated multi-perspective evaluation of clustering algorithms using Pareto fronts. **Bioinformatics** :btab038.
- J25 *J Tan+, D Ni+, JA Wali, DA Cox, GV Pinget, J Taitz, CI Daien, A Senior, MN Read, SJ Simpson, NJC King+, and L Macia+.* (2021). Dietary carbohydrate, particularly glucose, drives B cell lymphopoiesis and function. **iScience**, accepted for publication. + equal first/senior contributions.
- J24 *T Ashhurst+, F Marsh-Wakefield+, G Putri+, A Spiteri, D Shinko, MN Read, A Smith, NJC King.* (2021). Integration, exploration, and analysis of high-dimensional single-cell cytometry data using Spectre. **Cytometry: Part A**, 101(3):237-253. + equal contribution.
- J23 *JD Hywood, G Rice, SV Paeon, MN Read, M Biro.* (2021). Detection and characterisation of chemotactic swarming without cell tracking. **Journal of the Royal Society Interface** 18(176), 20200879.
- J22 *M Koutsakos, LC Rowntree, L Hensen, BY Chua, CE van de Sandt, JR Habel1, W Zhang, X Jia, L Kedzierski, TM Ashhurst, GH Putri, F Marsh-Wakefield, MN Read, et al., K Kedzierska* (2021). Integrated immune dynamics define correlates of COVID-19 severity and antibody responses. **Cell Reports Medicine** 2(3):100208.

- J21 *MN Read*\*, *K Alden*, *J Timmis*, *PS Andrews*. (2020). Strategies for Calibrating Models of Biology. **Briefings in Bioinformatics** 21(1):24-35
- J20 *N Lamm*, *MN Read*, *M Nobis*, *D Van Ly*, *SG Page*, *VP Masamsetti*, *P Timpson*, *M Biro* and *AJ Cesare*. (2020). Nuclear F-actin counteracts nuclear deformation and promotes fork repair during replication stress. **Nature Cell Biology** 22(12):1460-1470.
- J19 *JLG Nino*, *SV Pagoon*, *SS Tay*, *F Colakoglu*, *D Kempe*, *J Hywood*, *JK Mazalo*, *J Cremasco*, *M Govendir*, *LF Dagley*, *K Hsu*, *S Rizzetto*, *J Zieba*, *G Rice*, *V Prior*, *G O'Neill*, *RJ Williams*, *DR Nisbet*, *B Kramer*, *AI Webb*, *F Luciani*, *MN Read*, and *M Biro*. (2020). Cytotoxic T cells swarm by homotypic chemokine signalling. **eLife** 9:e56554.
- J18 *G Rangan*, *A Raghubanshi*, *A Chaitarvornkit*, *AN Chandra*, *R Gardos*, *A Munt*, *MN Read*, *S Saravanabavan*, *JQJ Zhang*, *ATY Wong* (2020). Current and emerging treatment options to prevent renal failure due to autosomal dominant polycystic kidney disease. **Expert Opinion on Orphan Drugs** 8(8):285-302.
- J17 *J Zoll* <sup>+</sup>, *MN Read* <sup>+</sup>, *S Heywood*, *E Estevez*, *J Marshall*, *H Kammoun*, *T Allen*, *A Holmes*, *M Febbraio*, and *D Henstridge* (2020). Fecal microbiota transplantation from high caloric-fed donors alters glucose metabolism in recipient mice, independently of adiposity or exercise status. **American Journal of Physiology-Endocrinology and Metabolism** 319(1):E203-2016. <sup>+</sup> co-first authors.
- J16 *CAM Fois*, *TYL Le*, *A Schindeler*, *S Naficy*, *DD McClure*, *MN Read*, *P Valtchev*, *A Khademhosseini* and *F Dehghani*. (2019). Models of the Gut for Analyzing the Impact of Food and Drugs. **Advanced Healthcare Materials** 8(21):1900968.
- J15 *GH Putri*, *MN Read*, *I Koprinska*, *D Singh*, *U Rohm*, *TM Ashhurst* and *NJC King*. (2019). ChronoClust: density-based clustering and cluster tracking in high-dimensional time-series data. **Knowledge Based Systems** 174:9-26.
- J14 *I Moran*, *A Nguyen*, *W Khoo*, *D Butt*, *K Bourne*, *C Young*, *J Hermes*, *M Biro*, *G Gracie*, *C Ma*, *C Munier*, *F Luciani*, *J Zaunders*, *A Parker*, *A Kelleher*, *S Tangye*, *P Croucher*, *R Brink*, *MN Read* and *T Phan*. (2018). Memory B Cells Are Reactivated in Subcapsular Proliferative Foci of Lymph Nodes. **Nature Communications** 9:3372.
- J13 *MN Read*\*, *AJ Holmes*. (2017). Towards an integrative understanding of diet-host-gut microbiome interactions. **Frontiers in Immunology** 8:538.  
This paper is in the top 5% of research outputs scored by Altmetric (as of 2017-11-06, score=39).
- J12 *AJ Holmes*, *YV Chew*, *F Colakoglu*, *JB Cliff*, *E Klaassens*, *MN Read*, *SM Solon-Biet*, *AC McMahon*, *VC Cogger*, *K Ruohonen*, *D Raubenheimer*, *DG Le Couteur*, *SJ Simpson*. (2017). Diet-microbiome interactions in health are controlled by intestinal nitrogen source constraints. **Cell Metabolism** 25:1-12.  
This paper is in the top 5% of research outputs scored by Altmetric (as of 2017-11-06, score=253).
- J11 *MN Read*\*, *K Alden*, *L Rose* and *J Timmis*. (2016). Automated multi-objective calibration of biological agent-based simulations. **Journal of the Royal Society Interface** 13(122):20160543.
- J10 *MN Read*\*, *J Bailey*, *J Timmis* and *T Chtanova*. (2016). Leukocyte motility models assessed through simulation and multi-objective optimization-based model selection. **PLoS Computational Biology** 12(9):e1005082.
- J9 *J Hywood*, *M Read*, *G Rice*. (2016). Statistical analysis of spatially homogeneous dynamic agent-based processes using functional time series analysis. **Spatial Statistics** 17:199-219.
- J8 *J Cosgrove*, *JA Butler*, *K Alden*, *M Read*, *V Kumar*, *J Timmis* and *M Coles*. (2015). Agent-based modelling in systems pharmacology. **CPT: Pharmacometrics & Systems Pharmacology** 4(11):615-629.
- J7 *K Alden*, *M Read*, *P Andrews*, *J Timmis* and *M Coles*. (2014). Applying spartan to understanding parameter uncertainty in simulations. **The R Journal** 6(2):1-10.
- J6 *M Read*\*, *PS Andrews*, *J Timmis* and *V Kumar*. (2014). Modelling biological behaviours using the unified modelling language: an immunological case study and critique. **Journal of the Royal Society Interface** 11(9):20140704.

- J5 M Read\*, P Andrews, J Timmis, R Williams, R Greaves, H Sheng, M Coles, V Kumar. (2013). Determining disease intervention strategies using spatially resolved simulations. **PLoS ONE** 8(11):e80506.
- J4 K Alden, M Read, J Timmis, P Andrews, H Veiga-Fernandes and M Coles. (2013). Spartan: a comprehensive tool for understanding uncertainty in simulations of biological systems. **PLoS Computational Biology** 9(2):e1002916.
- J3 R Greaves, M Read, J Timmis, P Andrews, J Butler, B Gerckens and V Kumar. (2013). In silico investigation of novel biological pathways: the role of CD200 in regulation of T cell priming in experimental autoimmune encephalomyelitis. **Biosystems** 112(2):107-121.
- J2 R Williams, R Greaves, M Read, J Timmis, P Andrews and V Kumar. (2013). In silico investigation into dendritic cell regulation of CD8Treg mediated killing of Th1 cells within murine experimental autoimmune encephalomyelitis. **BMC Bioinformatics** 14(Suppl 6):S9.
- J1 M Read\*, P Andrews, J Timmis and V Kumar. (2011). Techniques for grounding agent-based simulations in the real domain: a case study in experimental autoimmune encephalomyelitis. **Mathematical and Computer Modelling of Dynamical Systems** 17(4):296-302.

#### Refereed Conference Papers (\* corresponding author)

Typical in computer science and engineering is the publication of full-length papers in conference proceedings. All the conferences in which I publish employ a peer review process, typically using at least 3 reviewers, and a round of revisions prior to decisions regarding acceptance for publication.

- C9 GH Putri, MN Read\*, I Koprinska, TM Ashhurst and NHC King. (2019). Dimensionality reduction for clustering and cluster tracking of cytometry data. **International Conference on Artificial Neural Networks** pp 624-640.
- C8 B Naylor, M Read, A Turner, J Timmis and A Tyrrell. (2014). The Relay Chain: A scalable dynamic communication link between an exploratory underwater shoal and a surface vehicle. **Artificial Life (ALife)**, pages 290-297.
- C7 M Read\*, C Möeslinger, T Dipper, D Kengyel, J Hilder, R Thenius, A Tyrrell, J Timmis and T Schmickl. (2013). Profiling underwater swarm robotic shoaling performance using simulation. **Towards Autonomous Robotic Systems (TAROS)**, pages 404-416.
- C6 D Sutantyo, P Levi, C Möslinger and M Read (2013). Collective-adaptive Levy flight for underwater multi-robot exploration. **IEEE Conference on Mechatronics and Automation**, pages 456-462.
- C5 RB Greaves, M Read, J Timmis, PS Andrews, V Kumar. (2012). Extending an established simulation: exploration of the possible effects using a case study in experimental autoimmune encephalomyelitis. **International Conference on Information Processing in Cells and Tissues (IPCAT)**, LNCS 7223, pages 150-161.
- C4 T Schmickl, R Thenius, C Möslinger, J Timmis, A Tyrrell, M Read, J Hilder, J Halloy, C Stefanini, L Manfredi, A Campo, T. Dipper, D Sutantyo, S Kernbach. (2011). CoCoRo – The self-aware underwater swarm. **IEEE International Conferences on Self-Adaptive and Self-Organizing Systems (SASO)**, pages 120-126.
- C3 F Polack, P Andrews, T Ghetiu, M Read, S Stepney, J Timmis and A Sampson. (2010). Reflections on the simulation of complex systems for science. **IEEE International Conference on Engineering Complex Computer Systems (ICECCS)**, pages 276-285.
- C2 M Read\*, J Timmis, P Andrews and V Kumar. (2009). A domain model of Experimental Autoimmune Encephalomyelitis. **2nd Workshop on Complex Systems Modelling and Simulation**, pages 9-44.
- C1 M Read\*, J Timmis and P Andrews. (2008). Empirical investigation of an artificial cytokine network. **International Conference on Artificial Immune Systems (ICARIS)**, LNCS 5132, pages 340-351.

## Books

- B1 *S Stepney, FAC Polack, K Alden, P Andrews, J Bown, A Droop, RB Greaves, MN Read, A Sampson, J Timmis, A Winfield.* (2018). **Engineering Simulations as Scientific Instruments: A Pattern Language.** Springer International Publishing.

## Book Chapters (\* corresponding author)

- CH1 *M Read\*, P Andrews and J Timmis.* (2011). An introduction to artificial immune systems. **The Handbook of Natural Computing**, edited by G Rozenberg, T Back and J Kok. Springer, pages 1575-1597.

## Refereed Conference Abstracts (\* corresponding author)

The following represent peer-reviewed abstracts (1-2 pages) published in conference proceedings. In-person presentations to conference attendees facilitate rapid dissemination before pursuing publication in a journal; this is common in engineering and computer science.

- A8 *M Batten, E Shanahan, R Simpson, M Read, et al., and GV Long.* (2020). Gut microbiota predicts response and toxicity with neoadjuvant immunotherapy. **In: Proceedings of the Annual Meeting of the American Association for Cancer Research 2020.**
- A7 *M Read\*, J Timmis and T Chtanova.* (2015). Simulation-based analysis of in situ cellular motility. **European Conference on Artificial Life (ECAL)**, page 637.
- A6 *M Read\*, AJ Holmes, M Hartill-Law, S Solon-Biet, D Raubenheimer and SJ Simpson.* (2015). Simulating the influence of diet on the intestinal microbiome composition. **European Conference on Artificial Life (ECAL)**, page 638.
- A5 *M Read\*, M Tripp, H Leonova, L Rose and J Timmis.* (2013). Automated calibration of agent-based immunological simulation. **International Conference on Artificial Immune Systems track at European Conference on Artificial Life (ECAL)**, MIT Press, pages 874-875.
- A4 *E Hart, M Read, C McEwan, J Greensmith and U Aickelin.* (2013). On the role of the AIS practitioner. **Artificial Immune Systems track at European Conference on Artificial Life (ECAL)**, MIT Press, pages 891-892.
- A3 *M Read, J Butler, B Ole-Gerckens, J Timmis and V Kumar.* (2012). CD200 regulation can promote recovery from autoimmunity in Experimental Autoimmune Encephalomyelitis. Presented at **International Conference on Artificial Immune Systems (ICARIS)**.
- A2 *R Williams, M Read, J Timmis, P Andrews and V Kumar.* (2011). In silico investigation into CD8Treg mediated recovery in murine Experimental Autoimmune Encephalomyelitis. **International Conference on Artificial Immune Systems (ICARIS)**, LNCS 6825, pages 52-54.
- A1 *M Read\*, J Timmis, P Andrews and V Kumar.* (2009). Using UML to model EAE and its regulatory network (extended abstract). **International Conference on Artificial Immune Systems (ICARIS)**, LNCS 5666, pages 4-6.

## Other

- D3 *K Alden and M Read* (2013). Scientific software needs quality control. **Nature** 502:448. This is an editor-reviewed correspondence.
- D2 *M Read.* (2012). Statistical and Modelling Techniques to Build Confidence in the Investigation of Immunology through Agent-Based Simulation. **PhD thesis**, Department of Computer Science, University of York.

D1 M Read. (2007). Explicable Boolean Functions. **Masters degree dissertation**, Department of Computer Science, University of York.

## Research Funding

### Australian Research Council

- *T Chtanova, M Read*. Why do neutrophils swarm? **Australian Research Council Discovery Project**: \$560K, 2022-2024. DP220102278.
- *M Biro, M Read*. Search strategy optimisation by theory, functional analysis and simulation. **Australian Research Council Discovery Project**: \$388K, 2018-2021. DP180102458.

### External Competitive Grants

- *M Tracy, M Hinder, R Halliday, A McEwan, A Kumar, M Read, TA Goyen, M Cruz*. Virtual baby Project: (vbaby) predictive physiological modelling of critically ill preterm newborns. **Cerebral Palsy Alliance Research Foundation**: \$200K, 2020.
- *M Danta, S Ghaly, D Samocha-Bonet, C Bourke, M Read, L Macia, G Wark*. Metabolic monitoring of the microbiome in gastrointestinal disease study. **St Vincent's Clinic Foundation Research Grant**: \$40K, 2018.
- *C Clark, S Garcia, M Read*. Large dairy herds: Creating value from data. **Dairy Australia/Research and Development Grants**: \$158K, 2015-2017.
- *M Read*. Royal Academy of Engineering International Travel Grant: £500, 2009.

### Internal Competitive Grants

- *M Read, H McGuire, A Kumar, E Patrick, K Kott, T Ashhurst, B Di Bartolo*. Data science innovations to reveal the cellular foundations of atherosclerotic arterial disease. **Cardiovascular Bioengineering and Data Science Catalyst Award**: \$10K, 2020.
- *M Read, A Kumar, E Shanahan, et al*. Enabling personalised dietary recommendations to improve cardiometabolic health. **Cardiovascular Initiative Catalyst Award: Cardiovascular Precision Medicine**: \$10K, 2019.
- *M Read, G Rangan*. Understanding Polycystic Kidney Disease through Data-Driven Approaches. **Westmead Industrial Placement Scheme**: \$18.5K, 2019.
- *M Read, C Chow, A Kumar*. Advancing cardiovascular disease management through machine learning. **Westmead Industrial Placement Scheme**: \$18.5K, 2019.
- *T Ashurst, M Read, N King, U Roehm, I Koprinska, R Scalzo*. Mapping dynamic immunity: Next generation computational approaches to track the evolution of immune responses in West Nile virus and Zika virus encephalitis. **Marie Bashir Institute**: \$10,000, 2017.

## Prizes

- 2019 University of Sydney Faculty of Engineering, Research Conversazione award. Won by my PhD student, Juan Molina Ortiz.
- 2019 Dolby Australia Scientific Paper Competition, for the ChronoClust paper, output J15 above.

- 2017 Award for Reduction in the Use of Animals in Research, The University of Sydney. For simulations of the mouse gut, as related to output J12 above.
- Best computational immunology paper prize at the International Conference on Artificial Immune Systems (ICARIS) 2011, for output A2 (above). Voted for by conference attendees
- My PhD thesis was awarded 2<sup>nd</sup> prize in the University of York Computer Science Department’s “Best PhD Thesis Award 2011”
- My Masters degree dissertation was awarded the 2007 Science Engineering and Technology (SET) national prize in Information Technology
- My Masters degree dissertation was awarded “Best Project” prize by the University of York’s Computer Science department in 2007

## **Supervision of Research Students**

### **Current PhD**

- Jonathan Chung (2020). “Mapping disease state through machine learning and imaging mass cytometry.” Primary supervisor.
- Juan Ortiz (2019). “Modelling the gut: an in silico/in vitro approach to understanding the gut microbiome dynamics.” Primary supervisor.
- Givanna Haryono (2017). “Revealing the development of the immune response through dynamic clustering.” Primary supervisor.

### **Current MPhil**

- Jiadi Dong (2020). “Segmentation and classification of cells in imaging mass cytometry using deep learning methods.” Co-supervisor.

### **Current Honours-equivalent**

- Dijun Arthur Ng (2021). “Distinguishing junk- from informative-machine learning models in data-constrained biomedical contexts.” Primary supervisor.

### **Completed Masters and Honours-equivalent**

- Lina Gain (2020), Honours-equivalent research placement thesis. “Machine learning-based prediction of the blood metabolome from diet and gut microbial profiling.” Primary supervisor.
- Brendon Lam (2020), Honours thesis. “The immune system in Crohn’s disease as elucidated through computational modelling”, Primary supervisor.
- Aakanksha Jain (2020), Masters thesis. “Determining ecological roles in gut microbes through genome scale modelling and machine learning.” Primary supervisor.
- Yifan Shi (2019), Masters thesis. “Predicting post-prandial glucose responses from dietary, activity and gut microbiome quantifications through machine learning.” Primary supervisor.
- Alissa Chaitarvornkit (2020), Honours-equivalent research placement thesis. “Machine learning biomarkers for polycystic kidney disease.” Primary supervisor.
- Raymond Huang (2019), Honours. “Computational simulation of microbial monoculture growth dynamics; a prerequisite for rational microbiome-targeting interventions.” Primary supervisor.

- Juan Ortiz (2018), Masters thesis. “The role of gut microbial symbiosis on community resilience and hysteresis.” Primary supervisor.
- Avneet Kaur (2018), Honours. “Understanding and engineering neutrophil swarming.” Primary supervisor.
- Joshua Won (2016-17), Honours. “Partial differential equation modelling of gut microbial dynamics.” Associate supervisor.
- Mika Herath (2017), Honours. “Developing a model for studying hematopoietic stem cell recirculation: A multidisciplinary approach.” Associate supervisor.
- Deeksha Singh (2016), Honours. “Incremental clustering of high dimensional, high throughput cytometry data to analyse immune response to West Nile Virus.” Primary Supervisor.
- Madison Hartill-Law (2015), Honours. “A computational approach to investigating the interaction between diet and the microbiome.” Associate supervisor, attained a mark of 84 (distinction).
- Magnus Tripp (2013), Master of Engineering thesis. “Automated calibration of agent-based simulation for an autoimmune disease.” Associate supervisor.
- Sophie Alexander (2013), Master of Engineering thesis. “Simulating underwater swarm robotic systems.” Associate supervisor.
- Hannah Leonova (2012), MRes in Computational Biology. “Automated calibration of agent-based simulation for an autoimmune disease.” Associate supervisor.
- Richard Greaves (2011), Master of Research in Computer Science. “Computational modelling of Treg networks in Experimental Autoimmune Encephalomyelitis” Associate supervisor.
- Richard Williams (2010), Master of Research in Computational Biology (Distinction). “In silico experimentation using simulation of experimental autoimmune encephalomyelitis (EAE).” Associate supervisor.

## Reviewing and Programme Committee Duties

In 2019 and 2020 I peer-reviewed for the Australian Research Council.

- Peer reviewed journal manuscripts for:
  - Applied Soft Computing
  - Artificial Intelligence
  - BMC Bioinformatics
  - Computers in Biology and Medicine
  - Engineering Applications of Artificial Intelligence
  - Food & Function (Royal Society of Chemistry)
  - Frontiers in Microbiology
  - IEEE Transactions on Evolutionary Computation
  - The International Society for Microbial Ecology (ISME) Journal
  - Natural Computing
  - PLoS Computational Biology
  - Robotica
  - Robotics and Autonomous Systems
  - Science of Computer Programming
  - Swarm Intelligence
  - Water Research



- Organising committee member of:
  - Workshop Chair for the 2013 Confidence in Computer Simulation workshop, part of the Summer Computer Simulation Conference (SCSC)
  - Workshop Chair for the 2012 CoSMoS project workshop
  - Publicity Chair of the 2011 ICST Conference on Bio-Inspired Models of Network, Information, and Computing Systems (BIONETICS)
- Programme committee member of:
  - Genetic and Evolutionary Computation Conference (GECCO) 2015, 2016, 2018, 2019
  - IEEE Congress on Evolutionary Computation (CEC): 2018, 2019, 2020
  - Complex Systems Modelling and Simulation Workshop (CoSMoS) 2014, 2015
  - Artificial Immune Systems (AIS) track at Genetic and Evolutionary Computation Conference (GECCO) 2014
  - Summer Computer Simulation Conference (SCSC) 2013
  - International Conference on Artificial Immune Systems (ICARIS) 2012
  - The Complex Systems Modelling and Simulation Infrastructure (CoSMoS) 2010 workshop
  - International Conference on Engineering Complex Computer Systems (ICECCS) 2010.
  - International Conference on Artificial Immune Systems (ICARIS) 2009

## Visiting Research Posts

My doctoral research was conducted in conjunction with Dr. Vipin Kumar, an immunologist who headed the Laboratory of Autoimmunity at the Torrey Pines Institute for Molecular Studies (TPIMS) San Diego. I undertook six week-long trips to TPIMS between 2008 and 2012.

## Invited Talks, Research Seminar, and Summer School Participation

“Using single-cell cytometry to illustrate the generalisable unbiased evaluation of clustering algorithms using Pareto fronts.” Australian Bioinformatics And Computational Biology Society Conference, (virtual) 2020.

“Plasma metabolomic and gut microbiome responses to prebiotic supplements.” Sanitarium Health Food Company, 2020.

“Characterising immune cell movement, and its consequences, through live imaging and simulations.” Artificial Intelligence in Medical Imaging, Sydney 2020.

“Computational approaches to advancing biomedical understanding.” Zreiqat Lab, USyd 2019.

“Delivering pre/probiotic health benefits by moving beyond the ‘One Size Fits All’ paradigm.” St George and Sutherland Clinical School Research in Progress Meeting, 2019.

“Microbes and the Mind.” Psychology School Seminar, USyd 2019.

“Delivering pre/probiotic health benefits and getting them to market by moving beyond the ‘One Size Fits All’ paradigm.” Centre for Advanced Food Enginomics industry engagement workshop, Sydney 2019.

“Untangling biological complexity through modelling. Leukocyte motility and search for targets, and gut microbial responses to diet.” The Garvan Institute of Medical Research, Sydney 2018.

“Host- vs diet-derived nutrient balance, and carbon vs nitrogen limitation, determines the effect of dietary intervention on the microbiome.” Society for Mathematical Biology, Sydney 2018.

“Diagnostics and prognostics through machine learning; a tutorial and case study in gut microbiome-based weight-loss prediction.” Australian Society for Microbiology, Brisbane 2018.

“Host- vs diet-derived nutrient balance, and carbon vs nitrogen limitation, determines the effect of dietary intervention on the microbiome.” Cold Spring Harbour Asia conference on Microbiota, Metagenomics and Health, 2017.

“Determining immune cell search strategies: Studying motility through agent-based modelling and multi-objective optimization.” QUT School of Mathematical Sciences, 2017.

“Computational modelling and simulation techniques to investigate the gut bacterial ecosystems response to diet.” Invited talk at Theo Murphy Australian Frontiers of Science conference on the Microbiome, Adelaide, Australia, 2016.

“From an animal’s nutrient environment to the microbial environment.” Invited talk at Modelling the impact of the nutritional environment on host-microbiome interactions, the Cold Spring Harbour Asia Satellite Workshop, Suzhou, China, 2016.

“Simulating the influence of diet on the intestinal microbiome composition.” Invited talk at Australian Society for Microbiology conference, Canberra, Australia, 2015.

“Modelling the diet’s influence on gut bacterial communities.” Invited talk at the joint University of Sydney and Shanghai Jiao Tong University Research Alliance workshop, 2015.

Delivered an invited talk on the analysis and simulation of complex biological data to a hand-selected group of farmers from New South Wales, Australia, who are seeking data-driven ways to optimise their farming practices. Talk took place in December 2014.

“In Situ Imaging & Mechanistic Simulation of Cellular Swarming.” Invited talk at the Computer Graphics in Biomedical and Biological Imaging Data workshop, co-located with Computer Graphics International (CGI), Sydney, Australia, 2014.

“Modelling the gut microbiota response to host diet.” Invited talk at the Australia and New Zealand Obesity Society conference, Sydney, 2015.

“Capturing the immune system: from the wet-lab to the robot.” Invited guest lecture at the Awareness Summer School, Lucca, Italy, 2013. Further to this, I organized a case study for students to work on. The case study entailed deriving, in simulation, an algorithm run on underwater swarm robotic platforms that would organize the swarm into a dynamic chain that connected two (possibly moving) points in space.

“Determining disease intervention strategies using spatially resolved simulations.” Invited talk at the BSI Mathematical and Computational Modelling in Immunology conference, Cambridge, UK, 2013.

I was invited to participate in the AWARENESS (EU funded coordination action) “slides factory” in 2012, a meeting of representatives from projects examining self-awareness in autonomic systems who collectively wrote a popular science seminar and an academic course on the subject.

I have presented my research to a variety of research groups and departments within the University of York, to researchers at the Torrey Pines Institute for Molecular Studies in San Diego, and at the Cheltenham Science Festival 2010.

In 2011 I participated in an invitation-only Schloss Dagstuhl seminar series on Artificial Immune Systems (AIS), and lead a group examining how to derive AIS algorithms from established immunological simulations.

“Building confidence in immunological simulation.” Invited talk at the International Conference on Artificial Immune Systems, Cambridge, UK. 2011.

## Teaching and Education

I led the development of USyd's Master of Digital Health and Data Science, which will launch in 2022. The program will educate the next generation of students in data science principles and how they can solve data-driven health problems. This is a jointly offered, interdisciplinary course equally taught between the School of Computer Science and the Faculty of Medicine and Health's Discipline of Biomedical Informatics and Digital Health. My deep interdisciplinary knowledge was key to designing this degree; I understand the healthcare system context and the biological concepts that underpin a patient's health, and I have deep expertise in computational and data science technologies. As such, I expertly collaborated with partners across the Faculty of Engineering, the Faculty of Medicine and Health and the Office of the Deputy Vice Chancellor (Education) to position the program's curriculum and its strategic implementation across the faculties. My research program, which drives innovation in data science to advance healthcare, is ideally placed to support projects for the degree's Capstone experience.

I created, delivered and coordinated a **new unit of study, HTIN5005: Computational Approaches to Health Data** (2021). This is a core unit for USyd's Master of Digital Health and Data Science. It explores how tailored application of data science can enable personalised medicine, deliver remote care and clinical decision support systems, and optimise hospital operations using a wide range of health and biomedical data sources. From a technical perspective, the unit covers data types, descriptive and inferential statistical techniques, both supervised and unsupervised machine learning, computational simulations, and has a heavy focus on application in the R programming language. The cohort comprises students from both technical and health-related backgrounds. As such the material is designed to assume no prior knowledge, and yet students are able to apply data science techniques to model, visualise and interpret health data. The unit employs inquiry-based learning for assessments, wherein students have freedom to select healthcare problems and corresponding datasets on which to apply computational techniques to advance care. In 2021 the unit was delivered remotely over Zoom, given restrictions on in-person teaching due to the COVID-19 pandemic.

Other course coordination:

- HTIN5004: Integrated Approaches to Chronic Disease (2016, 2017, 2020; USYD). A core unit on the Master of Health Technology Innovation. I helped design this course.
- HTIN5001: Nature of Systems (2020; USYD). A core unit on the Master of Health Technology Innovation.

I regularly serve as an independent assessor for Capstone, Honours and Masters-level student projects.

Lecturing experience (lecture title and details of unit given):

- HTIN5005: Computational Approaches to Health Data (2021). This unit teaches biomedical and health-care data science.
- "Personalized healthcare with machine learning." (2020; USyd, postgraduate). Guest lecture for Sustainability and Intelligence in Project Management (PMGT5896).
- "Advancing medicine and healthcare through machine learning" (2020; USyd, postgraduate). Guest lecture for Machine Learning and Data Mining (COMP5318).
- "ICT-enabled diagnostics, prognostics and clinical decision support" (2019; USyd, post-graduate unit). 1h guest lecture on Enterprise Healthcare Information Systems (INFO5306).
- "Computational Modelling in Biological Research" (2016 - 2019; USyd, post-graduate unit). 1h lecture as part of Integrated Approaches to Chronic Disease (HTIN5004).
- "Bioinformatics for Cytometry" (2017; USyd, medical unit). Guest lecture on cutting edge technologies for analysing high-dimensional cytometry data. Given to students of USyd's Medical Program, Cardiovascular Sciences stream.
- "Simulating the Gut Microbial Response to Diet" (2016; USyd, post-graduate unit). A 3 hour session delivered as part of The Nature of Systems (HTIN5001).

- “Workshop on Writing Academic Papers” (2015; USyd). I co-organised and co-wrote this course, and delivered lectures on the presentation of data and selecting a journal to publish in.
- “Providing an Integrative Perspective of Biological Systems using Computer Simulations” (2014; USyd, undergraduate unit). A 1h guest lecture to 3rd year Veterinary Science students, studying Animal Biotechnology.
- I developed and delivered 4 hours of material for the Swarm Intelligence master’s unit (2012; UY, post-graduate). Titles:
  - “Group Behaviour”
  - “Simulation of Swarm Robotic Systems”
  - “Statistics and Sensitivity Analysis”
  - “Communication-less Boids” (a flocking algorithm)
- As part of UY’s Combating Infectious Disease: Computational Approaches in Translational Science Ph.D. training program, I developed and delivered the following, in both 2011 and 2012.
  - “The Experimental Autoimmune Encephalomyelitis Simulator”
  - “Performing *in silico* Experiments”
  - “Calibration & Sensitivity Analysis”
- “Simulation in Swarm Robotics” (2011; UY, post-graduate). A 1h lecture I prepared and delivered for the Swarm Intelligence unit.

## Service

I currently serve on a cross-disciplinary committee of academics that develop University of Sydney’s strategy for engaging the Westmead Health Precinct.

I was a founding member and inaugural Chair (2014-2016) of the Early Career Researcher Initiative committee, at the University of Sydney’s Charles Perkins Centre (CPC), an interdisciplinary research institute of 900 researchers. The committee organises activities supporting EMCR professional development, social integration and career progression. It also advocates on their behalf to the CPC’s executive committee.

In March 2015 I joined the NSW EMCR (Early and Mid Career Researcher) Network organising committee, and served as the Vice President. This network runs roughly 4 events a year, which aim to build professional networks between EMCRs and the private sector. Specifically, this is to make EMCRs aware of professional opportunities and careers where their training as researchers would be valued, and to provide the NSW private sector with exposure to a wide range highly skilled researchers. I stood down in 2017.

## Public Engagement

I ran a workshop on machine learning for teachers of science at a University of Sydney event in 2018.

In 2016 I worked with a journalist in the University of Sydney’s Alumni magazine who wrote an article on the gut microbiome that featured my research.

In March 2015 I delivered a public lecture on the interactions between host diet and gut microbial communities in influencing host health, at Ultimo library, Sydney.

I was interviewed and quoted in Can strategies used against computer viruses help us fight biological viruses? by Andy Favell, 2020.

## Academic References

### **Professor Jon Timmis**

Professor of Intelligent and Adaptive Systems.

Deputy Vice-Chancellor (Commercial), The University of Sutherland, United Kingdom

jon.timmis@sunderland.ac.uk

Prof. Timmis was my Ph.D. supervisor, and, thereafter, line manager during my first post-doctoral position at the University of York.

### **Professor Vipin Kumar (Chaturvedi)**

Professor of Medicine.

School of Health Sciences, UC San Diego.

vckumar@ucsd.edu

Prof. Kumar was my host for several international collaborative research visits that I conducted whilst at the University of York.

### **Associate Professor Irena Koprinska**

Faculty of Engineering, The University of Sydney, Australia

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A/Prof. Koprinska is a close collaborator on my machine learning-related projects.