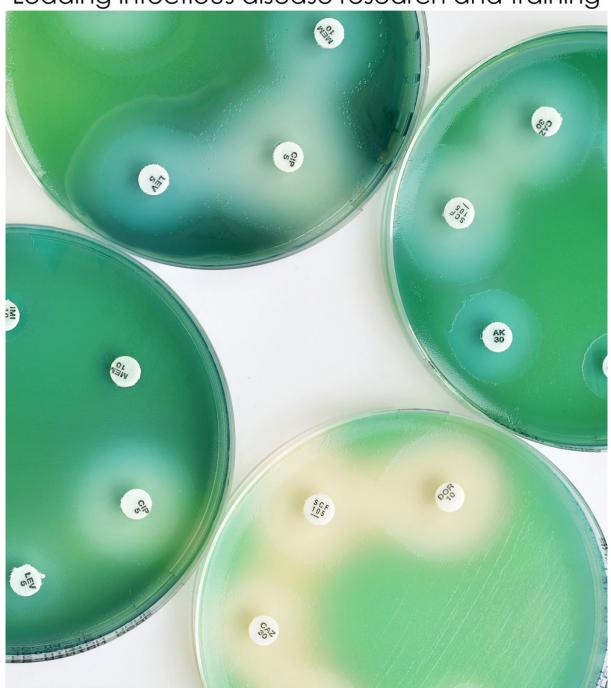
# Edinburgh Infectious Diseases

Leading infectious disease research and training



Annual Report 2016/17

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## **Executive summary**

## What is Edinburgh Infectious Diseases

We are the Network of researchers across the University of Edinburgh and associated organisations with an interest in infectious diseases.

## The aims of Edinburgh Infectious Diseases

- 1. Represent the strengths of infectious disease science in Edinburgh through our symposia, workshops, outreach activity and internet profile;
- 2. Maintain a strategic overview of infectious disease research in Edinburgh, to maximise synergy between established activities and promote new avenues for investigation;
- 3. Foster infectious disease teaching and training at all levels within the University, including the development of new postgraduate initiatives.

### Our major activities and achievements in 2016/17

- We have seen continued growth in size and spread of membership of *Edinburgh Infectious Diseases* to over 180 principal investigators.
- Our members were awarded over £34.5M for infectious disease research in FY 2015/16 and published over 470 research papers in 2016.
- Edinburgh Infectious Diseases has contributed to the development and coordination of major successful international funding applications.
- We launched our strategy for Antimicrobial Resistance Research in Edinburgh in March 2017.
- We continue to facilitate increased communication and collaboration between social scientists and basic and clinical researchers.
- We have supported the launch of the Wellcome Trust Four Year PhD Programme in Hosts, Pathogens and Global Health doctoral training programme.
- We have added value to the experience of undergraduate students in infectious diseases through support of the Society for Immunology and Infectious Diseases and organisation of seminars for the Infectious Diseases Honours class.
- We are connected to our members through compilation and circulation of a weekly newsletter of relevant seminars and news, and organisation of effective research networking events, which have contributed to building the vibrant community of researchers in Edinburgh working on infectious diseases.

## Director's Statement

The membership of *Edinburgh Infectious Diseases* continues to increase in size with a total of 850 scientists and clinicians including 180 group leaders. In the last year, we have continued to serve the community through our network activities, symposia, strategic workshops etc. We have also established key strategic priority areas for particular focus by *Edinburgh Infectious Diseases*. A key goal was the development of a strategy for antimicrobial resistance research (AMR) in Edinburgh built around 4 key AMR-related research themes led by theme leaders who are international leaders within their fields. The theme leaders will co-ordinate activities within and between themes underpinned by One Health and Social Science cross-thematic approaches. The AMR strategy was launched recently at a symposium held the Playfair library with over 200 registered attendants from Edinburgh along with guests from some of the major national funding bodies. Recent funding successes by *Edinburgh Infectious Diseases* members in AMR research are encouraging and we must now build on the new structures to establish Edinburgh as a leading centre of excellence for AMR research.

In other priority areas, our activities to promote interactions between basic and social scientists will culminate on 24 April in a one-day symposium 'Global Challenges in Infectious Disease: Showcasing social science research in Edinburgh'. We anticipate that these activities will lead to stronger research links between our basic and social scientists and funded collaborative projects. With regard to Global Challenges, we have recently surveyed the *Edinburgh Infectious Diseases* membership to identify current partnerships with low and middle income countries and we continue to promote Global Challenge Research fund (GCRF) bids by *Edinburgh Infectious Diseases* members, such as the recent successful TIBA NIHR grant application. Moving forward, in the light of the new Industrial Strategy Challenge Fund we will pro-actively support the development of partnerships with industry in readiness for ICSF funding calls.

An important remit for *Edinburgh Infectious Diseases* is supporting teaching and training and we have acted as a vehicle towards improving coordination and collaboration between existing undergraduate teaching programmes of relevance to infectious diseases in Edinburgh. With regard to post-graduate training and capacity building, we wish to build on existing research and infrastructure strengths in Edinburgh to train the next generation of scientists by establishing new doctoral training programmes. Current concepts under discussion include 'One Health models of Infectious Diseases' and Physical sciences and AMR'.

Finally, we have just launched our new website which along with our outreach activities, aims to extend the reach of *Edinburgh Infectious Diseases*, engaging the outside world and reporting our key activities, initiatives and the successes of our membership.

Overall, we look ahead to the next year with enthusiasm, aiming to build on recent successes, while continuing to identify new opportunities to promote excellent science and infectious disease training in Edinburgh.

Professor Ross Fitzgerald, Chair of Molecular Bacteriology, The Roslin Institute

Director of Edinburgh Infectious Diseases

April 2017

## Organisation and membership of Edinburgh Infectious Diseases

Edinburgh Infectious Diseases is coordinated by Director (Professor Ross Fitzgerald) and Executive Manager (Dr. Hilary Snaith), and supported by an administrative assistant (Jennifer Hurst). The Network has regular input from an Executive Committee, which meets once a month.

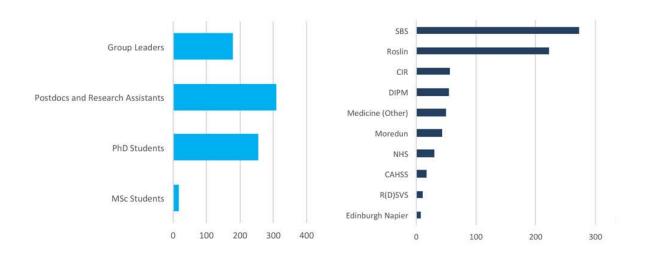
Table 1: Current members of the Edinburgh Infectious Diseases executive committee				
Member	Affiliation			
Dr. Till Bachmann	Division of Infection and Pathway Medicine, Royal Infirmary of Edinburgh, Little			
Dr. Till Bacilliaini	France			
Dr. Amy Buck	Institute of Immunology & Infection Research, Ashworth Laboratories, King's			
Dr. Arry Buck	Buildings			
Prof Harry Campbell	Usher Institute of Population Health Sciences and Informatics, Teviot Place			
Prof David Dockrell	Centre for Inflammation Research, Queen's Medical Research Institute, Little			
FIOI David Dockieli	France			
Dr. Bernadette Dutia	The Roslin Institute, Easter Bush			
Prof Gary Entrican	The Moredun Research Institute			
Prof Ross Fitzgerald	The Roslin Institute, Easter Bush			
Prof Clifford Leen	Department of Infectious Diseases, Western General Hospital			
Prof Keith Matthews	Institute of Immunology & Infection Research, Ashworth Laboratories, King's			
FIOI Reitii Matthews	Buildings			
Prof Harish Nair	Usher Institute of Population Health Sciences and Informatics, Teviot Place			
Prof Jürgen Schwarze	Centre for Inflammation Research, Queen's Medical Research Institute, Little			
rioi juigeii sciiwaize	France			
Dr. Hilary Snaith	Edinburgh Infectious Diseases, Ashworth Laboratories, King's Buildings			
Prof Mark Stevens	The Roslin Institute, Easter Bush			
Dr. Alice Street	School of Social and Political Science, George Square			
Dr. Kate Templeton	Royal Infirmary of Edinburgh, Little France			
Prof Sue Welburn	Division of Infection and Pathway Medicine, Royal Infirmary of Edinburgh, Little			
FIOI SUE WEIDUITI	France			

Over the past year membership of *Edinburgh Infectious Diseases* has grown to over 180 academics, drawn from the University of Edinburgh, Heriot Watt and Edinburgh Napier Universities, NHS Lothian and associated Institutes, including the Moredun Research Institute and Scotland's Rural College. A full listing of our members is on our website: http://www.eid.ed.ac.uk/members.

As of April 2017 Edinburgh Infectious Diseases has over 850 members, with 180 group leaders, over 300 postdocs and research assistants, and over 250 PhD students (see graph below left). The Roslin Institute and the School of Biological Studies continue to have the highest numbers of infectious disease researchers, but there are significant concentrations located at the Little France site, both in the Queen's Medical Research Institute and in the Chancellor's Building (see graph below right). We are also pleased to be engaging with more members of the College of Arts, Humanities and Social Sciences as part of our aim to foster interdisciplinary approaches to addressing questions in infectious disease research.

We were delighted in October of 2016 when Professor David Dockrell moved to the University of Edinburgh from Sheffield to take up the new Chair of Infection Medicine. He brings with him a vibrant research group that is based in the MRC Centre for Inflammation Research at Little France, and is focused on understanding the innate immune system and how it can be harnessed to control

infection. David has already taken an active role in the *Edinburgh Infectious Diseases* community and is a new member of our Executive Committee.



Left: Total numbers of group leaders, postdocs and research assistants, and students working in infectious diseases across Edinburgh; right: distribution of infectious diseases researchers.

Another significant change has been the appointment of Professor Eleanor Riley as David Hume's successor as Director of the Roslin Institute. Eleanor has a long-standing interest in anti-malarial immunity, and will arrive at the Institute later in September 2017.

Other new appointments include Professor Debby Bogaert who has recently joined the MRC Centre for Inflammation Research from the University Medical Center in Utrecht. Debby is now building her research group to study the physiology and pathophysiology of respiratory infections and inflammation from an ecological perspective. Dr. Gavin Paterson has joined the Roslin Institute as a senior lecturer in Applied Diagnostics with a research programme including sequence-based diagnostic and genomic epidemiology of bacterial pathogens. The expansion of the Edinburgh-China teaching programme with Zhejiang University, has also allowed the recruitment of Dr. Richard Sloan from Barts and London Hospital to join the Division of Infection and Pathway Medicine. Richard's research focuses on the antiviral factors that inhibit HIV and other retroviruses. Over the next year it is anticipated that there will also be new appointments in the School of Biological Studies to further strengthen infectious disease research.

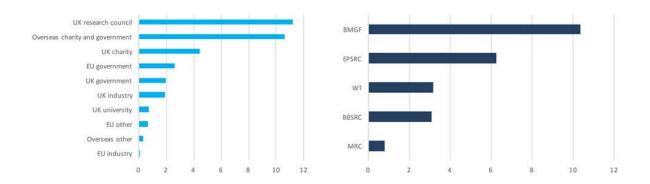
## Funding for our research

Members of *Edinburgh Infectious Diseases* have continued to be successful in attracting considerable external research funding over the past 18 months. Analysis of the Key Performance Indicator statistics produced by Edinburgh Research and Innovation demonstrates the success of our members in winning competitive research funding from a wide variety of sources.

From the data available for the 2015-2016 financial year, *Edinburgh Infectious Diseases* members were awarded £34.5M, in the context of total University of Edinburgh funding of £268M for FY 2015-2016.

These awards include a large grant of £10M from the Bill and Melinda Gates Foundation to fund the Center for Tropical Livestock Genetics and Health at the Roslin Institute. Other highlights included over £5M awarded to Mark Bradley and his team by the Engineering and Physical Sciences Research Council for molecular sensing and imaging, ~£1M awards by the European Commission and the Wellcome Trust to Rosalind Allen and Ross Fitzgerald respectively, to continue their work addressing antimicrobial resistance. Ivan Morrison and Tim Connelley received over £600K from the BBSRC to develop vaccines against bovine TB, and Finn Grey was awarded £510K by the MRC to continue his work on human Cytomegalovirus. Direct industry funding of £1.8M was also obtained, and further support was leveraged via RCUK LINK and IPA funding schemes (figures not currently available).

The graphs below quantify the funding sources including major UK and international charities, the UK and EU government research councils, and from industry collaborations.



Left: Total funding awarded to *Edinburgh Infectious Diseases* members in FY 2015-16; right: total funding from selected funders (BMGF – Bill and Melinda Gates Foundation; EPSRC – Engineering and Physical Sciences Research Council; WT – Wellcome Trust; BBSRC – Biotechnology and Biological Sciences Research Council; MRC – Medical Research Council.)

The first half of FY 2016-2017 has also seen considerable funding awards made for infectious disease research, with £17.5M awarded in total. Antimicrobial resistance research has over £3.5M with awards to David Dockrell and Meriem El Karoui from the MRC and the Wellcome Trust. Details of all the awards over £500K in FY 2015 - 2016, and in the first half of FY 2016 - 2017 are given in tables 2a and 2b below.

Table 2a: Awards >£500,000 made in FY 2015 to 2016				
PI Name	PI Dept.	Sponsor Name	Project Title	Award Total (£)
David Hume	Roslin Institute	Bill and Melinda Gates Foundation	Center for Tropical Livestock Genetics and Health	10,000,000
Mark Bradley	Chemistry	EPSRC	Multiplexed 'Touch and Tell' Optical Molecular Sensing and Imaging	5,077,841
Rosalind Allen	Physics and Astronomy	European Commission	EVOSTRUC: The physics of antibiotic resistance evolution	1,304,989
Steven H Spoel	Institute of Molecular Plant Sciences	European Commission	IMMUNE-EXPRESS: Proteasome- Mediated Gene Expression in Plant Immunity	1,111,080

Ross Fitzgerald	Roslin Institute	Wellcome Trust	Understanding bacterial host adaptation to combat infectious diseases	981,065		
Bruce Whitelaw	Roslin Institute	BBSRC	Genome editing for quantitative traits in livestock	889,293		
Christopher Haslett	Centre for Inflammation Research	EPSRC	Multiplexed "touch and tell" optical molecular sensing and Imaging	823,260		
Dave Burt	Roslin Institute	BBSRC	Unravelling the networks that regulate seasonal rhythmicity in the epigenome	743,257		
Tim S Walsh	Division of Health Sciences	Selex Ltd	SELEX Pilot: LASER Assisted Staphylococcal ERadication prior to orthopaedic implantation surgery	716,511		
Ivan Morrison	R(D)SVS	BBSRC	NKp46+CD3+ T-Cells As A Novel Target For Vaccines Against Bovine TB	£605,367		
Finn Grey	Roslin Institute	MRC	Systematic identification of host proteins involved in Human Cytomegalovirus replication, assembly and egress	510,054		
Table 2b: Awards >£500,000 made in first half FY 2016 to 2017						
PI Name	PI Dept.	Sponsor Name	Project Title	Award Total (£)		
David Dockrell	Centre for Inflammation Research	MRC	Optimising Innate Host Defence to Combat Antimicrobial Resistance	2,631,980		
Harish Nair	Centre for Global Health Research	European Commission	Respiratory Syncytial Virus Consortium in Europe	1,770,250		
Sarah Reece	School of Biological Sciences	Wellcome Trust	Parasite offence or host defence? The roles of biological rhythms in malaria infection	1,527,986		
Meriem El Karoui	School of Biological Sciences	Wellcome Trust	DNA repair and genetic stability: elucidating the effects of cell physiology in Escherichia coli	955,487		
Jayne Hope	The Roslin Institute	BBSRC	VACCINE: Defining signature responses at the innate-adaptive interface to inform the design of vaccines inducing cellular immunity	870,609		
Jürgen Haas	Division of Infection & Pathway Medicine	MRC	Control of type III interferon expression and Herpes simplex virus type 1 replication by miR-200	690,239		
Emily Gwyer Findlay	Centre for Inflammation Research	Royal Society	Cathelicidin is Critical for Pathogenic T cell Development in Multiple Sclerosis	662,275		
Harish Nair	Centre for Global Health Research	Sanofi Pasteur MSD Limited	Nasopharyngeal pneumococcal carriage study in South Asian infants	647,015		
Debby Bogaert	Centre for Inflammation Research	Chief Scientist Office	Early-life effects on host-microbiome interactions	533,000		

## Facilitating collaborative funding applications

The environment for infectious disease research in Edinburgh is enhanced by the network of connected researchers generated by *Edinburgh Infectious Diseases*. Over the past year this has facilitated applications for several large multisite funding programmes.

## Respiratory Syncytial Virus Consortium in Europe



In January 2017 the Innovative Medicines' Initiative awarded £24M to the Respiratory Syncytial Virus (RSV) Consortium in Europe (RESCEU) project, of which £1.77M has come to the University of Edinburgh. The project is led by Harish Nair in the Usher Institute for Population Health Sciences and Informatics, and

involves several colleagues from the University of Edinburgh, alongside European partners. The RESCEU team will gather data to draw up best practice guidelines to improve the way RSV-associated disease is monitored across Europe and to inform future vaccination programmes.

#### Tackling Infection to Benefit Africa

Earlier on this year, Mark Woolhouse and Francisca Mutapi, alongside colleagues from across Edinburgh Infectious Diseases and collaborators in nine African countries, prepared a consortium

application for the call from the National Institute for Health Research for Global Health Research Units. At the time of writing we have just heard that the Programme has been awarded £7.4M to create a new multidisciplinary centre for Tackling Infections to Benefit Africa (the TIBA Centre; tiba means to cure an infection in Swahili).



TIBA is a partnership between the University of Edinburgh and leading African scientists in Botswana, Ghana, Kenya, Rwanda, South Africa, Sudan, Tanzania, Uganda and Zimbabwe. Uniquely, TIBA will work on all aspects of infectious diseases management – from basic research to uptake by local communities – in a single, unified framework. This meets an often articulated but rarely addressed need, to view infectious disease management not in isolation but in the context of local health systems and wide societal issues. TIBA will also contribute to the strategic goal of improving the health research ecosystem in Africa, providing a rare opportunity for different countries to work together in a common framework.

#### Identifying links to connect researchers

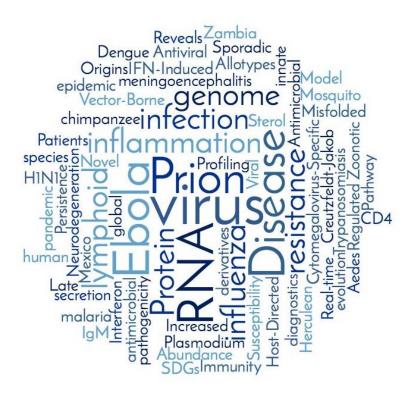
A key aspect to developing research proposals that are eligible for funding by the RCUK Grand Challenges Research Fund, is identifying the appropriate partners in low and middle income countries with whom to work. *Edinburgh Infectious Diseases* has contributed to building capacity in this area by compiling a data base of our members' collaborators in the developing world, which will be made available on our website. This has already been a valuable resource for the TIBA Centre application

(above), and we intend to continue augmenting the information available.

## Publication highlights

In 2016 Edinburgh Infectious Diseases members published over 470 papers (see Appendix 1). Selected highlights are three papers published in the Lancet, two in the Lancet Infectious Diseases, five in Nature, nine in Nature Communications, five in PLoS Pathogens, five in PLoS Neglected Tropical Diseases, three in PLoS Biology, three in Science, two in eLife and one in Cell. Details of some of the papers are below:

- Alderton, S., Macleod, E.T., Anderson, N.E., Schaten, K., Kuleszo, J., Simuunza, M., Welburn, S.C., and Atkinson, P.M. (2016). A Multi-Host Agent-Based Model for a Zoonotic, Vector-Borne Disease. A Case Study on Trypanosomiasis in Eastern Province, Zambia. PLoS Negl Trop Dis 10, e0005252.
- Alibhai, J., Blanco, R.A., Barria, M.A., Piccardo, P., Caughey, B., Perry, V.H., Freeman, T.C., and Manson, J.C. (2016). Distribution of Misfolded Prion Protein Seeding Activity Alone Does Not Predict Regions of Neurodegeneration. PLoS Biol *14*, e1002579.



 Clement, M., Marsden, M., Stacey, M.A., Abdul-Karim, J., Gimeno Brias, S., Costa Bento, D., Scurr, M.J., Ghazal, P., Weaver, C.T., Carlesso, G., et al. (2016). Cytomegalovirus-Specific IL-10-Producing CD4+ T Cells Are Governed by Type-I IFN-Induced IL-27 and Promote Virus Persistence. PLoS Pathog 12, e1006050.

- Donaldson, D.S., Sehgal, A., Rios, D., Williams, I.R., and Mabbott, N.A. (2016). Increased Abundance of M Cells in the Gut Epithelium Dramatically Enhances Oral Prion Disease Susceptibility. PLoS Pathog 12, e1006075.
- Duffin, R., O'Connor, R.A., Crittenden, S., Forster, T., Yu, C., Zheng, X., Smyth, D., Robb, C.T., Rossi, F., Skouras, C., et al. (2016). Prostaglandin E(2) constrains systemic inflammation through an innate lymphoid cell-IL-22 axis. Science *351*, 1333-1338.
- Gaunt, E., Wise, H.M., Zhang, H., Lee, L.N., Atkinson, N.J., Nicol, M.Q., Highton, A.J., Klenerman, P., Beard, P.M., Dutia, B.M., et al. (2016). Elevation of CpG frequencies in influenza A genome attenuates pathogenicity but enhances host response to infection. Elife *5*, e12735.
- Holmes, E.C., Dudas, G., Rambaut, A., and Andersen, K.G. (2016). The evolution of Ebola virus: Insights from the 2013-2016 epidemic. Nature *538*, 193-200.
- Jackson-Jones, L.H., Duncan, S.M., Magalhaes, M.S., Campbell, S.M., Maizels, R.M., McSorley, H.J., Allen, J.E., and Benezech, C. (2016). Fat-associated lymphoid clusters control local IgM secretion during pleural infection and lung inflammation. Nat Commun *7*, 12651.
- Laxminarayan, R., Sridhar, D., Blaser, M., Wang, M., and Woolhouse, M. (2016). Achieving global targets for antimicrobial resistance. Science *353*, 874-875.
- Mena, I., Nelson, M.I., Quezada-Monroy, F., Dutta, J., Cortes-Fernandez, R., Lara-Puente, J.H., Castro-Peralta, F., Cunha, L.F., Trovao, N.S., Lozano-Dubernard, B., et al. (2016). Origins of the 2009 H1N1 influenza pandemic in swine in Mexico. Elife *28*.
- Miesen, P., Ivens, A., Buck, A.H., and van Rij, R.P. (2016). Small RNA Profiling in Dengue Virus 2-Infected Aedes Mosquito Cells Reveals Viral piRNAs and Novel Host miRNAs. PLoS Negl Trop Dis 10, e0004452.
- Moore, R.A., Head, M.W., Ironside, J.W., Ritchie, D.L., Zanusso, G., Pyo Choi, Y., and Priola, S.A. (2016). The Distribution of Prion Protein Allotypes Differs Between Sporadic and latrogenic Creutzfeldt-Jakob Disease Patients. PLoS Pathog 12, e1005416.
- Quick, J., Loman, N.J., Duraffour, S., Simpson, J.T., Severi, E., Cowley, L., Bore, J.A., Koundouno, R., Dudas, G., Mikhail, A., et al. (2016). Real-time, portable genome sequencing for Ebola surveillance. Nature 530, 228-232.
- Robertson, K.A., Hsieh, W.Y., Forster, T., Blanc, M., Lu, H., Crick, P.J., Yutuc, E., Watterson, S., Martin, K., Griffiths, S.J., et al. (2016). An Interferon Regulated MicroRNA Provides Broad Cell-Intrinsic Antiviral Immunity through Multihit Host-Directed Targeting of the Sterol Pathway. PLoS Biol 14, e1002364.
- Sridhar, D. (2016). Making the SDGs useful: a Herculean task. Lancet 388, 1453-1454.
- Sundararaman, S.A., Plenderleith, L.J., Liu, W., Loy, D.E., Learn, G.H., Li, Y., Shaw, K.S., Ayouba, A., Peeters, M., Speede, S., et al. (2016). Genomes of cryptic chimpanzee Plasmodium species reveal key evolutionary events leading to human malaria. Nat Commun *7*, 11078.
- Swain, P.S., Stevenson, K., Leary, A., Montano-Gutierrez, L.F., Clark, I.B., Vogel, J., and Pilizota, T. (2016). Inferring time derivatives including cell growth rates using Gaussian processes. Nat Commun 7, 13766.
- Syed, S.N., Ducrotoy, M.J., and Bachmann, T.T. (2016). Antimicrobial resistance diagnostics: time to call in the young? The Lancet. Infectious diseases *16*, 519-521.

## A research strategy in Edinburgh to combat Antimicrobial Resistance

Antimicrobial resistance (AMR) is one of the most important challenges currently facing humankind. The government-commissioned O'Neill report (Review on Antimicrobial Resistance, 2014-2016) predicted a massive increase in deaths per year by 2050 as a direct result of the acquisition of resistance to antibiotics (from a baseline of 700,000 deaths per year currently). Furthermore, without effective policies to prevent the spread of AMR, it is estimated that the cost to the global economy would be 100 trillion USD. These potentially devastating predictions have led to the prioritisation of AMR research as a major focus for UK research funding by RCUK, and Wellcome Trust, and additional dedicated resources have been provided in the form of the Fleming fund, and Global Innovation fund, etc.



Edinburgh Infectious Diseases has recently assessed the strengths of AMR research in Edinburgh, highlighting potential areas for future focus, synergy and enhancement in the context of the current and ongoing national AMR research priority areas. It is clear that there will be considerable and sustained funding opportunities in the area of AMR research over the coming years and it is imperative that Edinburgh researchers are successful in competing for available funding and provide leadership in addressing this global challenge. In the new strategy we have identified four key AMR-related research areas with potential for excellence in Edinburgh; (i) Global and local epidemiology of AMR; (ii) Rapid diagnostics for more effective use of antibiotics; (iii) Biological, physical and evolutionary mechanisms underpinning AMR; and (iv) Alternatives to antimicrobials for controlling infections.

To lead advances across these themes Mark Woolhouse (Centre for Immunity, Infection and Evolution) will be the AMR Champion. Alongside three other theme leaders, Rosalind Allen (School of Physics and Astronomy), David Dockrell (Centre for Inflammation Research) and Till Bachmann (Division of Infection and Pathway Medicine), Mark will promote our AMR science, co-ordinate activities and spearhead funding applications. Future success will also depend on building capacity in excellent AMR research through support of early career scientists, and continued development of links with international (particularly in low, middle income countries) and industrial partners.

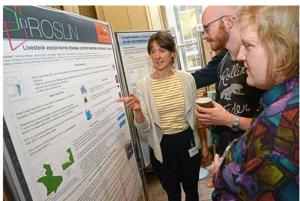
#### Launch of Antimicrobial Resistance Research Strategy

The new strategy was launched at the Edinburgh AMR Research Symposium on Wednesday 22 March 2017, to an audience of over 180. As AMR Champion, Mark Woolhouse set the stage for the day in his keynote lecture, explaining how AMR has arisen, what implications this has for human and animal

health and what is needed to ensure we are able to use antibiotics effectively into the future. The rest of day focused on the four research themes and highlighted some of the ways Edinburgh is already contributing solutions to counter AMR.

The day concluded with a very informative panel discussion with representatives from the Medical Research Council (Caroline Harris), The Engineering and Physical Sciences Research Council (Stephanie Newland), the Chief Medical Officer of Scotland (Alistair Leanord), Health Protection Scotland (Eleanor Anderson) and The University of Edinburgh Research Support Office (Catherine Burns). Their main take home messages were to think big and boldly, to be willing to take risks and to be multi- and interdisciplinary. Good advice, which was well received by all in the audience, as *Edinburgh Infectious Diseases* looks to support and develop new funding initiatives in AMR.





Left: The Edinburgh AMR leaders: left to right – David Dockrell, Till Bachmann, Ross Fitzgerald, Mark Woolhouse and Rosalind Allen; right: Poster session at the Edinburgh AMR Research Symposium. (Photos by Norrie Russell).

## Funding highlights for Antimicrobial Resistance Research

There have been several significant finding successes for AMR research over the past year. Notably Ross Fitzgerald (Roslin Institute) was awarded £981K by the Wellcome Trust, Meriem El Karoui (School of Biological Sciences) was awarded £955K, also by the Wellcome Trust, and David Dockrell (Centre for Inflammation Research) and colleagues received £2.6M from the Medical Research Council. Furthermore, Till Bachmann (Division of Infection and Pathway Medicine) is the leader of a JPIAMR Horizon 2020 funded network which will bring further prominence to our own research in AMR. We have also established links with the new AMR Centre based in Manchester, with a view to contributing to their rapid diagnostics platforms.

## Social science in basic and clinical infectious disease research



Another area we have sought to develop during the course of the past year are connections between researchers in clinical and basic science, and those active in infectious diseases research in the social sciences. During 2016 we ran three workshops with the Institute for

Advanced Studies in the Humanities "Common Ground", which brought together colleagues to explore the interface between research interests. A major common thread in these discussions was antimicrobial resistance; firstly, identifying the barriers to development and implementation of rapid diagnostics, and the secondly understanding how the behaviours of vets and farmers can affect the acquisition and spread of resistance in animals and the environment.

We have been working with the Edinburgh Centre for Medical Anthropology to host a one-day symposium "Global Challenges in Infectious Diseases", which will highlight social science researchers in Edinburgh, and provide a spring board for the development of new collaborative projects. The

symposium will also feed into a discussion for postgraduate students from across the University with the keynote speaker, Professor Vinh Kim Nguyen from the Université de Montréal, to promote interdisciplinarity at all levels.



We have also fostered interactions between clinical and basic science researchers and colleagues at the Edinburgh College of Art and in the School of Health and Social Sciences. We held a productive meeting to discuss possible approaches to recent Research Council funding calls; these discussions are still at an early stage, but we are pleased to have been instrumental in supporting our members to reach out to new collaborators for the future.

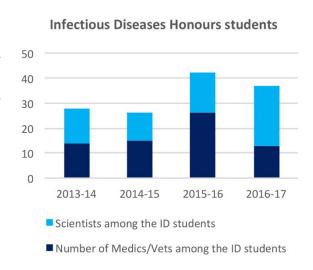
## Teaching and training

#### Undergraduate teaching in infectious diseases

The undergraduate honours programmes in Infectious Diseases and Immunology continue to attract talented students.

The Immunology programme has had 26 and 27 students for the past two years and is anticipating

similar numbers for the 2017/18 programme. The Infectious Diseases programme has also had steady numbers of students. It is likely that now all medical students at the University of Edinburgh will be required to undertake an intercalated degree between years 2 and 3 of their training, leading to increased demand for all Biomedical Honours programmes, including Infectious Diseases. This represents an opportunity for the programme, to promote the importance of infectious diseases to future doctors, and awareness of the possible research training options for clinicians in infectious disease in Edinburgh.



The Infectious Diseases and Immunology Honours programmes are run by the Biomedical Teaching Organisation (BMTO) and the Biology Teaching Organisation (BTO), respectively. As was discussed at the meeting of the Strategic Board in 2016, this separation of highly related subjects into two different School between in teaching has led both to some reduction in student course choice, and in duplication of some taught material.

Edinburgh Infectious Diseases, through Bernadette Dutia (Roslin Institute) has initiated discussion with the Heads of the BTO and the BMTO, and the Immunology and Infectious Diseases Honours Programme organisers, to explore possible improvements that can be made to programme delivery and integration. Although no major decisions have been made, these discussions have been encouraging and have led to some useful ideas, including, arranging timetabling to accommodate more sharing of elective courses and reassessing 3<sup>rd</sup> year requirements so that students are able to keep open their options for Honours programmes.

Edinburgh Infectious Diseases has continued to support the current Infectious Diseases Honours students throughout the year. The Executive manager met with the students at the start of the academic year to introduce the network and the activities they can access. The students receive the weekly newsletter listing the Edinburgh-wide seminars and events they can attend. We also supported the class to host a seminar by new chair of Infection Medicine David Dockrell, which was well attended by members from across Edinburgh. Support has also been given to the student-led Society for Immunology and Infectious Diseases, which has hosted three highly successful seminars this year from Edinburgh academics Sue Welburn and Sarah Howie, and from Quentin Sattentau at the University of Oxford.

## Undergraduate summer placements at Leiden University Medical Center

We have continued our very fruitful collaboration with Leiden University Medical Center, and again sent a third year Infectious Diseases honours student to spend the July and August 2016 in the Netherlands. Andres Rodriguez Barrientos worked with Prof Annemieke Geluk and her lab on the development of a diagnostic assay for *Mycobacterium leprae* for use in Bangladesh and Leiden.



About his time in Leiden, Andres says: "Working in the labs of the LUMC has been very special and has educated me more about how 'real' investigation happens.... Engaging with different projects helped me understand that research is very much a collaborative activity and involves scientists within and between teams".

We have already recruited two new students, one each from the Immunology and Infectious Diseases Honours programmes, who will travel to Leiden again in summer 2017.

## Postgraduate training

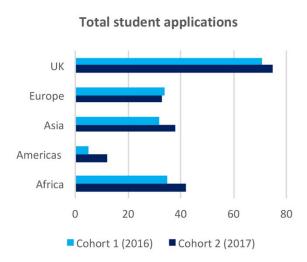
Doctoral training is a vital aspect of fostering young researchers. There are now over 250 PhD students at the University of Edinburgh currently embarked on infectious diseases-related projects.

#### Wellcome Trust Four Year PhD Programme Hosts, Pathogens and Global Health

The Wellcome Trust Four Year PhD Programme in Hosts, Pathogens and Global Health is led by co-Directors Keith Matthews (Institute of Immunology and Infection Research) and Mark Woolhouse (Centre for Immunity, Infection



and Evolution). It is centered in the Ashworth Labs at Kings Buildings with a total of 23 supervisors, including two based at Little France, and three at the Roslin Institute. The Programme launched in 2016 with the first cohort of students arriving in Edinburgh on 1 October. There are six students in total in this first year, who come from Ireland, Germany, South Africa and the UK. They have had a busy and productive year, carrying out three mini rotation projects, alongside tutorials in the three themes of the programme – Molecular Phylogeny and Epidemiology, Evolutionary Biology and Ecology, and Molecular Biology and Immunology. The programme has also included training in statistics and bioinformatics, presentation skills and relevant career development, to help the students make informed decisions as they navigate the course of their PhDs.



Recruitment for the second cohort was carried out in March 2017. We received over 200 applications for the available places, 105 from UK and EU candidates, and 95 from international candidates. Overall this represents an increase of 13% on total applications in comparison with 2016. The looming issues surrounding Brexit did not significantly deter candidates from EU countries, but we nevertheless saw a drop in applicants from 19.2 to 16.5% of the total. This year's recruits are from Hungary, Cyprus, Germany, The Gambia and the UK, and we look forward to welcoming them to Edinburgh in October.

### Edinburgh Infectious Diseases Ker Memorial PhD Studentship

Edinburgh Infectious Diseases is very fortunate to have the sponsorship of Miss Aileen Ker, who generously supports activities promoting infectious disease research at the University in memory of her father and grandfather who were both physicians in Edinburgh during the first half of the 20<sup>th</sup> century. Since 2009 some of her donation has been used to fund a PhD studentship, with the most recent recipient of the award, Mariya Goncheva (shown below left), completing her studies with Bernadette Dutia and Ross Fitzgerald at the Roslin Institute in September 2017.



Miss Ker is continuing to sponsor a new PhD student for a further four years, starting in October 2017. We have just appointed Greg Milne from the University of Bath to this studentship which will be held jointly in the School of Biological Sciences and the Deanery of Biomedical Sciences. The new studentship will be structured similarly to a University of Edinburgh Principal's Career Development Scholarship to provide additional training in teaching alongside research.

#### Ker Memorial Prize

In additional to the PhD studentship, Miss Ker also supports the Ker Memorial Prize, which is awarded annually to the student presenting the best PhD thesis in Infectious Diseases at the University of Edinburgh. In 2016 the Ker Memorial Prize was awarded to Chris Johnston from Rick Maizels' lab in the Institute of Immunology and Infection Research, for his work on helminth-induced immunomodulation during sold organ transplantation. As part of his prize Chris presented his work at the *Edinburgh Infectious Diseases* annual symposium in June 2016 (see above), and received a cheque for £500.

#### Development of Massive Open Online Course on Antibiotics and AMR

A strand of activity to promote engagement with prospective students is via Massive Open Online Courses (MOOCs). Given the high global profile of antimicrobial resistance, and the broad depth of expertise in this subject in Edinburgh, Thamarai Schneiders in the Division of Infection and Pathway Medicine, supported by *Edinburgh Infectious Diseases*, has been working to develop a course in "Antibiotics and the rise of Antimicrobial Resistance". This course has received support from the Deanery of Biomedical Sciences and has been submitted to the University of Edinburgh MOOC office for approval. The course will be aimed at a general, educated audience, and is intended to highlight the importance of antimicrobial resistance to a world-wide audience, showcasing the contributions made by Edinburgh researchers. If approved, it is anticipated that the course will run in mid-2018.

## Communication, collaboration and engagement

One of the key ways in which *Edinburgh Infectious Diseases* adds value to the research environment at the University of Edinburgh and associated organisations, is to provide opportunities for our members to meet with each other, and learn about relevant research going on across the city. On a weekly basis we send a newsletter to all our members with details of relevant seminars and events, and throughout the year we host a variety of events aimed at sharing knowledge and providing fora where new ideas, interactions and collaborations can develop. We have also provided some financial

support for seminar series organised by the Edinburgh Microbiology Forum and the Edinburgh Virology Group.

Table 3: Edinburg	Table 3: Edinburgh Infectious Diseases events over the past year				
When	Event	Location			
February 2016	Common Ground 1 – Finding the	Institute for Advanced Studies in the			
	connections in social sciences	Humanities, UoE			
May 2016	Common Ground 2 – Finding the	Institute for Advanced Studies in the			
	connections in social sciences	Humanities, UoE			
June 2016	Annual Symposium 2016	The Hub, Castle Hill			
June 2016	Seminar by Michael Hsieh	Ashworth Laboratories, UoE			
September 2016	Mini-workshop with SynthSys	Roger Land Building, UoE			
November 2016	Public Winter Lecture	Playfair Library, UoE			
November 2016	Student forum with Wendy Barclay	Old Medical School, UoE			
February 2017	GCRF Metabolic Modelling for Infectious	Roslin Institute, UoE			
	Diseases workshop				
March 2017	ID4 seminar by David Dockrell	Queen's Medical Research Institute, UoE			
March 2017	Edinburgh AMR symposium	Playfair Library, UoE			
Upcoming					
April 2017	Global Challenges in Infectious Disease –	50 George Square, UoE			
	Social Science Showcase				
June 2017	Annual Symposium 2017	John McIntyre Conference Centre			

## Symposia and workshops organised by Edinburgh Infectious Diseases

Our flagship event is the Annual Symposium. The 2016 symposium was held at The Hub on the Royal Mile and was attended by 200 staff, postdocs and students. As ever we had an extremely varied programme with everything from *Theileria coinfections* and East Coast Fever (Mark Woolhouse, Centre for Immunity, Infection and Evolution), to squirrel pox in UK red squirrels (Colin McInnes, Moredun Research Institute), translational genomics in critical care medicine (Kenny Baillie, Roslin Institute) and controlling sleeping sickness in Southern Sudan (Pete Kingsley, School of Social and Political Science).





Left: A full house at The Hub, Castle Hill for our 5<sup>th</sup> Annual Symposium; right: Gordon Dougan, the Ker Memorial Lecturer, Chris Johnson, winner of the Ker Memorial Prize, and Ross Fitzgerald. (Photos Norrie Russell)

We were very pleased to have Prof Gordon Dougan FRS from the Sanger Institute as the Ker Memorial Lecturer who concluded the day with an excellent presentation on *Genetic analysis of host-pathogen interactions during infection*.

Another highlight of the day was the lively and well attended poster session. Over 40 posters were displayed, with the prize for best student poster awarded to Filipa Henderson Sousa from Edinburgh Napier University for her work on "Cationic Host Defence Peptides as novel therapeutics for rhinovirus and influenza virus infection", and the postdoc poster prize going to Prerna Vohra at the Roslin Institute for her work on "Understanding the dynamics of Salmonella infection in cattle".







Left and centre: Lively poster sessions at the annual symposium; right: poster prize winners Filipa Henderson Sousa and Prerna Vohra with Ross Fitzgerald. (Photos Norrie Russell)

## Adding value to existing University of Edinburgh activity

In addition to the main symposium, over the past year we have sought to identify particular topics of interest to our members and develop opportunities where these can be explored in more detail. We are also very aware of the need for our members to identify research programmes that are aligned with the Grand Challenges Research Fund.

One area where there is great potential for synergy in Edinburgh the use of synthetic biology to address questions in vaccine design and production. To help build links between infectious diseases researchers and colleagues in the Centre for Synthetic and Systems





Biology, we ran a successful mini-workshop in September 2016. More recently, members of *Edinburgh Infectious Diseases* also took part in a sandpit with synthetic biology researchers from several research organisations in China, to help developing collaborative projects that could be funded by the GCRF. In February 2017 we organised a workshop with Paul Michels (School of Biological Sciences) in February 2017 on how metabolic modelling can be used to inform control of infectious disease. Three visiting scientists from Brazil and Mexico joined

colleagues from the Universities of Edinburgh and Glasgow. The event was funded by a BBSRC Impact Accelerator Award to promote GCRF-relevant collaborations.

We also seek to add value to existing seminar programmes, hosting visiting speakers when

appropriate. This year Michael Hsieh from the Biomedical Research Institute, Maryland, gave a widely attended seminar on the ABCs of schistosome infection and met with a number of researchers to explore collaborations opportunities.

We have also connected with a broad audience at our public Winter Lecture, this year given by



Professor Wendy Barclay from Imperial College, London, on the highly topic subject of influenza vaccines. This was held in the Playfair library and attracted a wide audience of over 180 people.

## Public engagement

We continue to promote public engagement through supporting researchers to develop their own outreach projects, and through the existing resource of the "Bugs and Bones" in the Ashworth

Treasure Chest. These samples travelled throughout Edinburgh during the course of the year at a variety of science festivals. We have also used them to run workshops in primary schools in Midlothian and the Borders, on the spread and control on infectious disease. These activities have given us the opportunity to give training to undergraduate students on the Science Communications programme



about how to educate, enthuse and inspire the next generation of scientists. Additionally *Edinburgh Infectious Diseases* has been actively involved with the College of Science and Engineering Public Engagement team to develop a new strategy for public engagement that supports all staff and students to engage with all strata of our society with our research.

#### Building an online presence

We continue to develop our social media platforms and website in order to reach audiences out with Edinburgh. Twitter is an effective way of disseminating news about events and studentship opportunities, with many of these tweets being widely re-tweeted (@edin\_eid). During the past year our followers on Twitter have grown 50% to 2144, with our tweets being read more than 40,000 times a month.

The *Edinburgh Infectious Diseases* website remains a valuable medium for communication about the research of our members and the opportunities to work with us: http://www.eid.ed.ac.uk/.



During 2016 we had almost 85,000 page views, from a total of 27,640 individual viewers, 32% of whom were returning visitors. As shown on the map (left), visitors to the site are from around the world: in the last year 69% of visitors were from Europe, 12% from Asia, 10% from the Americas and 7% from Africa. 61% of visitors were between the ages of 18 and 35, which reflects the popularity of the pages for Wellcome Trust Four Year PhD Programme in

Hosts, Pathogens and Global Health, and other student-directed information.

The news and events sections of the website are regularly updated and are also highly visited; a summary of the news stories from the website is presented in Appendix 3. We also use our YouTube

channel to publish talks from our Symposia and Winter Lecture.

The launch of the AMR research strategy was accompanied by the development of a new website on the EdWeb platform supported by the University of Edinburgh web development team, http://www.ed.ac.uk/edinburgh-infectious-diseases/amr. Over the course of 2017 we will be extending this new website to include all the Edinburgh Infectious Diseases content.



## Appendix 1

## Publications from Edinburgh Infectious Diseases members in 2016

- 1. Adu, B., Cherif, M.K., Bosomprah, S., Diarra, A., Arthur, F.K., Dickson, E.K., Corradin, G., Cavanagh, D.R., Theisen, M., Sirima, S.B., et al. (2016). Antibody levels against GLURP R2, MSP1 block 2 hybrid and AS202.11 and the risk of malaria in children living in hyperendemic (Burkina Faso) and hypo-endemic (Ghana) areas. Malaria journal *15*, 123.
- 2. Ahlstrom, C., Barkema, H.W., Stevenson, K., Zadoks, R.N., Biek, R., Kao, R., Trewby, H., Haupstein, D., Kelton, D.F., Fecteau, G., et al. (2016). Genome-Wide Diversity and Phylogeography of Mycobacterium avium subsp. paratuberculosis in Canadian Dairy Cattle. PLoS One *11*, e0149017.
- 3. Al-Rubaye, D.S., Henihan, G., Al-Abasly, A.K., Seagar, A.L., Al-Attraqchi, A.A., Schulze, H., Hashim, D.S., Kamil, J.K., Laurenson, I.F., and Bachmann, T.T. (2016). Genotypic assessment of drug-resistant tuberculosis in Baghdad and other Iraqi provinces using low-cost and low-density DNA microarrays. Journal of medical microbiology *65*, 114-122.
- 4. Alderton, S., Macleod, E.T., Anderson, N.E., Schaten, K., Kuleszo, J., Simuunza, M., Welburn, S.C., and Atkinson, P.M. (2016). A Multi-Host Agent-Based Model for a Zoonotic, Vector-Borne Disease. A Case Study on Trypanosomiasis in Eastern Province, Zambia. PLoS Negl Trop Dis *10*, e0005252.
- 5. Alexander, K.A., Raggatt, L.J., Millard, S., Batoon, L., Chiu-Ku Wu, A., Chang, M.K., Hume, D.A., and Pettit, A.R. (2017). Resting and injury-induced inflamed periosteum contain multiple macrophage subsets that are located at sites of bone growth and regeneration. Immunol Cell Biol *95*, 7-16.
- 6. Aliberti, S., Hill, A.T., Mantero, M., Battaglia, S., Centanni, S., Cicero, S.L., Lacedonia, D., Saetta, M., Chalmers, J.D., and Blasi, F. (2016). Quality standards for the management of bronchiectasis in Italy: a national audit. Eur Respir J 48, 244-248.
- 7. Alibhai, J., Blanco, R.A., Barria, M.A., Piccardo, P., Caughey, B., Perry, V.H., Freeman, T.C., and Manson, J.C. (2016). Distribution of Misfolded Prion Protein Seeding Activity Alone Does Not Predict Regions of Neurodegeneration. PLoS Biol *14*, e1002579.
- 8. Allen, R.C., McNally, L., Popat, R., and Brown, S.P. (2016). Quorum sensing protects bacterial cooperation from exploitation by cheats. The ISME journal *10*(7), 1706-1716.
- 9. Alpren, C., Sloan, M., Boegler, K.A., Martin, D.W., Ervin, E., Washburn, F., Rickert, R., Singh, T., and Redd, J.T. (2016). Notes from The Field: Ebola Virus Disease Cluster Northern Sierra Leone, January 2016. MMWR. Morbidity and mortality weekly report *65*, 681-682.
- 10. Amanfo, S.A., Mduluza, T., Midzi, N., Cavanagh, D.R., and Mutapi, F. (2016). Seroepidemiology of Plasmodium species infections in Zimbabwean population. Malaria journal *15*, 267.
- 11. Amin, N., Allebrandt, K.V., van der Spek, A., Muller-Myhsok, B., Hek, K., Teder-Laving, M., Hayward, C., Esko, T., van Mill, J.G., Mbarek, H., et al. (2016). Genetic variants in RBFOX3 are associated with sleep latency. European journal of human genetics: EJHG *24*, 1488-1495.
- 12. Amyes, S.G. (2016). Editorial. International journal of infectious diseases: IJID: official publication of the International Society for Infectious Diseases.
- 13. Amyes, S.G. (2016). Corrigendum to 'Editorial' [International Journal of Infectious Diseases 50 (2016) 83-84]. International journal of infectious diseases: IJID: official publication of the International Society for Infectious Diseases *51*, 109.
- 14. Anastasi, E., MacArthur, I., Scortti, M., Alvarez, S., Giguere, S., and Vazquez-Boland, J.A. (2016). Pangenome and Phylogenomic Analysis of the Pathogenic Actinobacterium Rhodococcus equi. Genome biology and evolution *8*, 3140-3148.
- 15. Anderson, C., Khan, M.A., Wong, F., Solovieva, T., Oliveira, N.M., Baldock, R.A., Tickle, C., Burt, D.W., and Stern, C.D. (2016). A strategy to discover new organizers identifies a putative heart organizer. Nat Commun *7*, 12656.
- Anderson, N.E., Bessell, P.R., Mubanga, J., Thomas, R., Eisler, M.C., Fevre, E.M., and Welburn, S.C. (2016). Ecological Monitoring and Health Research in Luambe National Park, Zambia: Generation of Baseline Data Layers. EcoHealth 13, 511-524.
- 17. Andrews, C.P., Kruuk, L.E., and Smiseth, P.T. (2017). Evolution of elaborate parental care: phenotypic and genetic correlations between parent and offspring traits. Behavioral ecology: official journal of the International Society for Behavioral Ecology *28*, 39-48.

- 18. Anichini, G., Iqbal, M., Rafiq, N.M., Ironside, J.W., and Kamel, M. (2016). Sacrificing the superior petrosal vein during microvascular decompression. Is it safe? Learning the hard way. Case report and review of literature. Surgical neurology international *7*, S415-420.
- 19. Armstrong, S.D., Xia, D., Bah, G.S., Krishna, R., Ngangyung, H.F., LaCourse, E.J., McSorley, H.J., Kengne-Ouafo, J.A., Chounna-Ndongmo, P.W., Wanji, S., et al. (2016). Stage-specific proteomes from Onchocerca ochengi, sister species of the human river blindness parasite, uncover adaptations to a nodular lifestyle. Mol Cell Proteomics *15*, 2554-2575.
- 20. Auten, R., Schwarze, J., Ren, C., Davis, S., and Noah, T.L. (2016). Pediatric Pulmonology year in review 2015: Part 1. Pediatr Pulmonol *51*, 733-739.
- 21. Auty, H., Cleaveland, S., Malele, I., Masoy, J., Lembo, T., Bessell, P., Torr, S., Picozzi, K., and Welburn, S.C. (2016). Quantifying Heterogeneity in Host-Vector Contact: Tsetse (Glossina swynnertoni and G. pallidipes) Host Choice in Serengeti National Park, Tanzania. PLoS One *11*, e0161291.
- Baele, G., Suchard, M.A., Rambaut, A., and Lemey, P. (2016). Emerging concepts of data integration in pathogen phylodynamics. Systematic biology *66*, e47-e65.
- 23. Bah, S.Y., Dickinson, P., Forster, T., Kampmann, B., and Ghazal, P. (2016). Immune oxysterols: Role in mycobacterial infection and inflammation. The Journal of steroid biochemistry and molecular biology.
- 24. Bain, C.C., Hawley, C.A., Garner, H., Scott, C.L., Schridde, A., Steers, N.J., Mack, M., Joshi, A., Guilliams, M., Mowat, A.M., et al. (2016). Long-lived self-renewing bone marrow-derived macrophages displace embryo-derived cells to inhabit adult serous cavities. Nat Commun 7, ncomms11852.
- 25. Balsells, E., Filipescu, T., Kyaw, M.H., Wiuff, C., Campbell, H., and Nair, H. (2016). Infection prevention and control of Clostridium difficile: a global review of guidelines, strategies, and recommendations. J Glob Health *6*, 020410.
- 26. Banos, G., Winters, M., Mrode, R., Mitchell, A.P., Bishop, S.C., Woolliams, J.A., and Coffey, M.P. (2017). Genetic evaluation for bovine tuberculosis resistance in dairy cattle. J Dairy Sci *100*, 1272-1281.
- 27. Bardosh, K.L., Berbri, I.E., Ducrotoy, M., Bouslikhane, M., Ouafaa, F.F., and Welburn, S.C. (2016). ZOONOTIC ENCOUNTERS AT THE SLAUGHTERHOUSE: PATHWAYS AND POSSIBILITIES FOR THE CONTROL OF CYSTIC ECHINOCOCCOSIS IN NORTHERN MOROCCO. Journal of biosocial science 48 Suppl 1, S92-s115.
- 28. Barron, R.M., King, D., Jeffrey, M., McGovern, G., Agarwal, S., Gill, A.C., and Piccardo, P. (2016). PrP aggregation can be seeded by pre-formed recombinant PrP amyloid fibrils without the replication of infectious prions. Acta neuropathologica *132*, 611-624.
- 29. Bartley, P.M., Hamilton, C., Wilson, C., Innes, E.A., and Katzer, F. (2016). Erratum to: Detection of Babesia annae DNA in lung exudate samples from Red foxes (Vulpes vulpes) in Great Britain. Parasit Vectors *9*, 275.
- 30. Bartley, P.M., Hamilton, C., Wilson, C., Innes, E.A., and Katzer, F. (2016). Detection of Babesia annae DNA in lung exudate samples from Red foxes (Vulpes vulpes) in Great Britain. Parasit Vectors *9*, 84.
- 31. Berenos, C., Ellis, P.A., Pilkington, J.G., and Pemberton, J.M. (2016). Genomic Analysis Reveals Depression Due To Both Individual And Maternal Inbreeding In A Free-Living Mammal Population. Mol Ecol *25*, 3152-3168.
- 32. Berk, Z., Bishop, S.C., Forbes, A.B., and Kyriazakis, I. (2016). A simulation model to investigate interactions between first season grazing calves and Ostertagia ostertagi. Vet Parasitol *226*, 198-209.
- 33. Besier, R.B., Kahn, L.P., Sargison, N.D., and Van Wyk, J.A. (2016). Diagnosis, Treatment and Management of Haemonchus contortus in Small Ruminants. Advances in parasitology *93*, 181-238.
- 34. Besier, R.B., Kahn, L.P., Sargison, N.D., and Van Wyk, J.A. (2016). The Pathophysiology, Ecology and Epidemiology of Haemonchus contortus Infection in Small Ruminants. Advances in parasitology *93*, 95-143.
- 35. Bewley, M.A., Belchamber, K.B., Chana, K.K., Budd, R.C., Donaldson, G., Wedzicha, J.A., Brightling, C.E., Kilty, I., Donnelly, L.E., Barnes, P.J., et al. (2016). Differential Effects of p38, MAPK, PI3K or Rho Kinase Inhibitors on Bacterial Phagocytosis and Efferocytosis by Macrophages in COPD. PLoS One 11, e0163139.
- 36. Bhatia, R., Kavanagh, K., Cubie, H.A., Serrano, I., Wennington, H., Hopkins, M., Pan, J., Pollock, K.G., Palmer, T.J., and Cuschieri, K. (2016). Use of HPV testing for cervical screening in vaccinated women-Insights from the SHEVa (Scottish HPV Prevalence in Vaccinated Women) study. International journal of cancer *138*, 2922-2931.
- 37. Bielejec, F., Baele, G., Vrancken, B., Suchard, M.A., Rambaut, A., and Lemey, P. (2016). SpreaD3: Interactive Visualization of Spatiotemporal History and Trait Evolutionary Processes. Mol Biol Evol *33*, 2167-2169.

- 38. Blackley, D.J., Wiley, M.R., Ladner, J.T., Fallah, M., Lo, T., Gilbert, M.L., Gregory, C., D'Ambrozio, J., Coulter, S., Mate, S., et al. (2016). Reduced evolutionary rate in reemerged Ebola virus transmission chains. Science advances *2*, e1600378.
- 39. Blayney, J.K., Davison, T., McCabe, N., Walker, S., Keating, K., Delaney, T., Greenan, C., Williams, A.R., McCluggage, W.G., Capes-Davis, A., et al. (2016). Prior knowledge transfer across transcriptional data sets and technologies using compositional statistics yields new mislabelled ovarian cell line. Nucleic Acids Res *44*, e137.
- 40. Bonsall, D., Gregory, W.F., Ip, C.L., Donfield, S., Iles, J., Ansari, M.A., Piazza, P., Trebes, A., Brown, A., Frater, J., et al. (2016). Evaluation of Viremia Frequencies of a Novel Human Pegivirus by Using Bioinformatic Screening and PCR. Emerg Infect Dis 22, 671-678.
- 41. Booth, J.W., Hamzah, L., Jose, S., Horsfield, C., O'Donnell, P., McAdoo, S., Kumar, E.A., Turner-Stokes, T., Khatib, N., Das, P., et al. (2016). Clinical characteristics and outcomes of HIV-associated immune complex kidney disease. Nephrology, dialysis, transplantation: official publication of the European Dialysis and Transplant Association European Renal Association 31, 2099-2107.
- 42. Borowska, D., Rothwell, L., Bailey, R.A., Watson, K., and Kaiser, P. (2016). Identification of stable reference genes for quantitative PCR in cells derived from chicken lymphoid organs. Vet Immunol Immunopathol *170*, 20-24.
- 43. Bourhy, H., Nakoune, E., Hall, M., Nouvellet, P., Lepelletier, A., Talbi, C., Watier, L., Holmes, E.C., Cauchemez, S., Lemey, P., et al. (2016). Revealing the Micro-scale Signature of Endemic Zoonotic Disease Transmission in an African Urban Setting. PLoS Pathog 12, e1005525.
- 44. Boyapati, R.K., Rossi, A.G., Satsangi, J., and Ho, G.T. (2016). Gut mucosal DAMPs in IBD: from mechanisms to therapeutic implications. Mucosal immunology *9*, 567-582.
- 45. Bradford, B.M., Brown, K.L., and Mabbott, N.A. (2016). Prion pathogenesis is unaltered following down-regulation of SIGN-R1. Virology *497*, 337-345.
- 46. Bridle, H.L., Heringa, M.B., and Schafer, A.I. (2016). Solid-phase microextraction to determine micropollutant-macromolecule partition coefficients. Nature protocols *11*, 1328-1344.
- 47. Brook, M., Tomlinson, G.H., Miles, K., Smith, R.W., Rossi, A.G., Hiemstra, P.S., van 't Wout, E.F., Dean, J.L., Gray, N.K., Lu, W., et al. (2016). Neutrophil-derived alpha defensins control inflammation by inhibiting macrophage mRNA translation. Proc Natl Acad Sci U S A *113*, 4350-4355.
- 48. Brown, A., Halliday, J.S., Swadling, L., Madden, R.G., Bendall, R., Hunter, J.G., Maggs, J., Simmonds, P., Smith, D.B., Vine, L., et al. (2016). Characterization of the Specificity, Functionality, and Durability of Host T-Cell Responses Against the Full-Length Hepatitis E Virus. Hepatology (Baltimore, Md.) *64*, 1934-1950.
- 49. Buda, R., Liu, Y., Yang, J., Hegde, S., Stevenson, K., Bai, F., and Pilizota, T. (2016). Dynamics of Escherichia coli's passive response to a sudden decrease in external osmolarity. Proc Natl Acad Sci U S A *113*, E5838-e5846.
- 50. Burgess, S.T., Nunn, F., Nath, M., Frew, D., Wells, B., Marr, E.J., Huntley, J.F., McNeilly, T.N., and Nisbet, A.J. (2016). A recombinant subunit vaccine for the control of ovine psoroptic mange (sheep scab). Vet Res 47, 26.
- 51. Burn, G.L., Cornish, G.H., Potrzebowska, K., Samuelsson, M., Griffie, J., Minoughan, S., Yates, M., Ashdown, G., Pernodet, N., Morrison, V.L., et al. (2016). Superresolution imaging of the cytoplasmic phosphatase PTPN22 links integrin-mediated T cell adhesion with autoimmunity. Science signaling *9*, ra99.
- 52. Burrells, A., Opsteegh, M., Pollock, K.G., Alexander, C.L., Chatterton, J., Evans, R., Walker, R., McKenzie, C.A., Hill, D., Innes, E.A., et al. (2016). The prevalence and genotypic analysis of Toxoplasma gondii from individuals in Scotland, 2006-2012. Parasit Vectors *9*, 324.
- 53. Callaby, R., Toye, P., Jennings, A., Thumbi, S.M., Coetzer, J.A., Conradie Van Wyk, I.C., Hanotte, O., Mbole-Kariuki, M.N., Bronsvoort, B.M., Kruuk, L.E., et al. (2016). Seroprevalence of respiratory viral pathogens of indigenous calves in Western Kenya. Research in veterinary science *108*, 120-124.
- 54. Campbell, H., Bont, L., and Nair, H. (2015). Respiratory syncytial virus (RSV) disease new data needed to guide future policy. J Glob Health *5*, 020101.
- 55. Canini, L., Woolhouse, M.E., Maines, T.R., and Carrat, F. (2016). Heterogeneous shedding of influenza by human subjects and its implications for epidemiology and control. Sci Rep *6*, 38749.
- 56. Carter, L.M., Pollitt, L.C., Wilson, L.G., and Reece, S.E. (2016). Ecological influences on the behaviour and fertility of malaria parasites. Malaria journal *15*, 220.
- 57. Cassady-Cain, R.L., Blackburn, E.A., Alsarraf, H., Dedic, E., Bease, A.G., Bottcher, B., Jorgensen, R., Wear, M., and Stevens, M.P. (2016). Biophysical characterization and activity of lymphostatin, a

- multifunctional virulence factor of attaching & effacing Escherichia coli. The Journal of biological chemistry *291*, 5803-5816.
- 58. Cassady-Cain, R.L., Blackburn, E.A., Bell, C.R., Elshina, E., Hope, J.C., and Stevens, M.P. (2017). Inhibition of Antigen-Specific and Nonspecific Stimulation of Bovine T and B Cells by Lymphostatin from Attaching and Effacing Escherichia coli. Infect Immun *85*.
- 59. Catafau, A.M., Bullich, S., Seibyl, J.P., Barthel, H., Ghetti, B., Leverenz, J., Ironside, J.W., Schulz-Schaeffer, W.J., Hoffmann, A., and Sabri, O. (2016). Cerebellar Amyloid-beta Plaques: How Frequent Are They, and Do They Influence 18F-Florbetaben SUV Ratios? Journal of nuclear medicine: official publication, Society of Nuclear Medicine *57*, 1740-1745.
- 60. Cescatti, M., Saverioni, D., Capellari, S., Tagliavini, F., Kitamoto, T., Ironside, J., Giese, A., and Parchi, P. (2016). Analysis of conformational stability of abnormal prion protein aggregates across the spectrum of Creutzfeldt-Jakob disease prions. J Virol *90*, 6244-6254.
- 61. Chalmers, J.D., Akram, A.R., Singanayagam, A., Wilcox, M.H., and Hill, A.T. (2016). Risk factors for Clostridium difficile infection in hospitalized patients with community-acquired pneumonia. The Journal of infection 73, 45-53.
- 62. Chalmers, J.D., Aliberti, S., Polverino, E., Vendrell, M., Crichton, M., Loebinger, M., Dimakou, K., Clifton, I., van der Eerden, M., Rohde, G., et al. (2016). The EMBARC European Bronchiectasis Registry: protocol for an international observational study. ERJ open research *2*, 00081.
- 63. Chalmers, J.D., McDonnell, M.J., Rutherford, R., Davidson, J., Finch, S., Crichton, M., Dupont, L., Hill, A.T., Fardon, T.C., De Soyza, A., et al. (2016). The generalizability of bronchiectasis randomized controlled trials: A multicentre cohort study. Respiratory medicine *112*, 51-58.
- 64. Checchi, M., Hewitt, P.E., Bennett, P., Ward, H.J., Will, R.G., Mackenzie, J.M., and Sinka, K. (2016). Tenyear follow-up of two cohorts with an increased risk of variant CJD: donors to individuals who later developed variant CJD and other recipients of these at-risk donors. Vox Sang 111, 325-332.
- 65. Chen, Q., Thomas, J.T., Gimenez-Lirola, L.G., Hardham, J.M., Gao, Q., Gerber, P.F., Opriessnig, T., Zheng, Y., Li, G., Gauger, P.C., et al. (2016). Evaluation of serological cross-reactivity and cross-neutralization between the United States porcine epidemic diarrhea virus prototype and S-INDEL-variant strains. BMC Vet Res 12, 70.
- 66. Chen, X., Liu, S., Rao, P., Bradshaw, J., and Weller, R. (2016). Topical application of superoxide dismutase mediated by HIV-TAT peptide attenuates UVB-induced damages in human skin. European journal of pharmaceutics and biopharmaceutics: official journal of Arbeitsgemeinschaft fur Pharmazeutische Verfahrenstechnik e.V 107, 286-294.
- 67. Chintoan-Uta, C., Cassady-Cain, R.L., and Stevens, M.P. (2016). Evaluation of flagellum-related proteins FliD and FspA as subunit vaccines against Campylobacter jejuni colonisation in chickens. Vaccine *34*, 1739-1743.
- 68. Choudhury, D., Tanner, M.G., McAughtrie, S., Yu, F., Mills, B., Choudhary, T.R., Seth, S., Craven, T.H., Stone, J.M., Mati, I.K., et al. (2017). Endoscopic sensing of alveolar pH. Biomedical optics express 8, 243-259
- 69. Christensen, L.L., Selman, C., Blount, J.D., Pilkington, J.G., Watt, K.A., Pemberton, J.M., Reid, J.M., and Nussey, D.H. (2016). Marker-dependent associations among oxidative stress, growth and survival during early life in a wild mammal. Proc Biol Sci 283, 283.
- 70. Chu, J.Y., Dransfield, I., Rossi, A.G., and Vermeren, S. (2016). Non-canonical PI3K-Cdc42-Pak-Mek-Erk Signaling Promotes Immune-Complex-Induced Apoptosis in Human Neutrophils. Cell reports *17*, 374-386.
- 71. Churchill, D., Waters, L., Ahmed, N., Angus, B., Boffito, M., Bower, M., Dunn, D., Edwards, S., Emerson, C., Fidler, S., et al. (2016). British HIV Association guidelines for the treatment of HIV-1-positive adults with antiretroviral therapy 2015. HIV Med *17 Suppl 4*, s2-s104.
- 72. Clavadetscher, J., Hoffmann, S., Lilienkampf, A., Mackay, L., Yusop, R.M., Rider, S.A., Mullins, J.J., and Bradley, M. (2016). Copper Catalysis in Living Systems and In Situ Drug Synthesis. Angewandte Chemie (International ed. in English) *55*, 15662-15666.
- 73. Claycomb, J., Abreu-Goodger, C., and Buck, A.H. (2017). RNA-Mediated Communication Between Helminths and their Hosts: The Missing Links. RNA Biol, 1-6.
- 74. Clement, M., Marsden, M., Stacey, M.A., Abdul-Karim, J., Gimeno Brias, S., Costa Bento, D., Scurr, M.J., Ghazal, P., Weaver, C.T., Carlesso, G., et al. (2016). Cytomegalovirus-Specific IL-10-Producing CD4+ T Cells Are Governed by Type-I IFN-Induced IL-27 and Promote Virus Persistence. PLoS Pathog 12, e1006050.

- 75. Coakley, G., Buck, A.H., and Maizels, R.M. (2016). Host parasite communications-Messages from helminths for the immune system: Parasite communication and cell-cell interactions. Mol Biochem Parasitol *208*, 33-40.
- 76. Concha-Marambio, L., Pritzkow, S., Moda, F., Tagliavini, F., Ironside, J.W., Schulz, P.E., and Soto, C. (2016). Detection of prions in blood from patients with variant Creutzfeldt-Jakob disease. Science translational medicine *8*, 370ra183.
- 77. Connelley, T.K., Li, X., MacHugh, N., Colau, D., Graham, S.P., van der Bruggen, P., Taracha, E.L., Gill, A., and Morrison, W.I. (2016). CD8 T-cell responses against the immunodominant Theileria parva peptide Tp249-59 are composed of two distinct populations specific for overlapping 11-mer and 10-mer epitopes. Immunology *149*, 172-185.
- 78. Conway Morris, A., Datta, D., Shankar-Hari, M., Weir, C.J., Rennie, J., Antonelli, J., Rossi, A.G., Warner, N., Keenan, J., Wang, A., et al. (2016). Predictive value of cell-surface markers in infections in critically ill patients: protocol for an observational study (ImmuNe FailurE in Critical Therapy (INFECT) Study). BMJ open 6, e011326.
- 79. Corbishley, A., Connelley, T.K., Wolfson, E.B., Ballingall, K., Beckett, A.E., Gally, D.L., and McNeilly, T.N. (2016). Identification of epitopes recognised by mucosal CD4(+) T-cell populations from cattle experimentally colonised with Escherichia coli O157:H7. Vet Res *47*, 90.
- 80. Cordeiro, O.G., Chypre, M., Brouard, N., Rauber, S., Alloush, F., Romera-Hernandez, M., Benezech, C., Li, Z., Eckly, A., Coles, M.C., et al. (2016). Integrin-Alpha IIb Identifies Murine Lymph Node Lymphatic Endothelial Cells Responsive to RANKL. PLoS One *11*, e0151848.
- 81. Costa, E., Buxton, J., Brown, J., Templeton, K.E., Breuer, J., and Johannessen, I. (2016). Fatal disseminated varicella zoster infection following zoster vaccination in an immunocompromised patient. BMJ case reports *2016*.
- 82. Coutinho, A.E., Kipari, T.M., Zhang, Z., Esteves, C.L., Lucas, C.D., Gilmour, J.S., Webster, S.P., Walker, B.R., Hughes, J., Savill, J.S., et al. (2016). 11beta-Hydroxysteroid Dehydrogenase type 1 is expressed in neutrophils and restrains an inflammatory response in male mice. Endocrinology *157*, 2928-2936.
- 83. Crilly, J.P., Nunn, F., Busin, V., Marr, E.J., Jennings, A.E., Burgess, S.T., and Sargison, N.D. (2016). Investigation and treatment of ovine psoroptic otoacariasis. Veterinary dermatology *27*, 206-e252.
- 84. Crilly, J.P., Soderquist, L., Holmstrom, A., and Sargison, N.D. (2016). Proof of concept of ovine artificial insemination by vaginal deposition of frozen-thawed semen under UK sheep-farming conditions. Vet Rec *178*, 532.
- 85. Cubie, H.A., Morton, D., Kawonga, E., Mautanga, M., Mwenitete, I., Teakle, N., Ngwira, B., Walker, H., Walker, G., Kafwafwa, S., et al. (2017). HPV prevalence in women attending cervical screening in rural Malawi using the cartridge-based Xpert(R) HPV assay. Journal of clinical virology: the official publication of the Pan American Society for Clinical Virology 87, 1-4.
- 86. Cui, J., Biernacka, K., Fan, J., Gerber, P.F., Stadejek, T., and Opriessnig, T. (2016). Circulation of Porcine Parvovirus Types 1 through 6 in Serum Samples Obtained from Six Commercial Polish Pig Farms. Transbound Emerg Dis.
- 87. Cui, J., Fan, J., Gerber, P.F., Biernacka, K., Stadejek, T., Xiao, C.T., and Opriessnig, T. (2016). First identification of porcine parvovirus 6 in Poland. Virus genes *53*, 100-104.
- 88. Cunningham, S., Nair, H., and Campbell, H. (2016). Deciphering clinical phenotypes in acute viral lower respiratory tract infection: Bronchiolitis is not an island. Thorax *71*, 679-680.
- 89. Currie, S.M., Gwyer Findlay, E., McFarlane, A.J., Fitch, P.M., Bottcher, B., Colegrave, N., Paras, A., Jozwik, A., Chiu, C., Schwarze, J., et al. (2016). Cathelicidins Have Direct Antiviral Activity against Respiratory Syncytial Virus In Vitro and Protective Function In Vivo in Mice and Humans. Journal of immunology (Baltimore, Md.: 1950) 196, 2699-2710.
- 90. Cuypers, B., Lecordier, L., Meehan, C.J., Van den Broeck, F., Imamura, H., Buscher, P., Dujardin, J.C., Laukens, K., Schnaufer, A., Dewar, C., et al. (2016). Apolipoprotein L1 Variant Associated with Increased Susceptibility to Trypanosome Infection. mBio 7, e02198-02115.
- 91. Danial, J., Ballard-Smith, S., Horsburgh, C., Crombie, C., Ovens, A., Templeton, K.E., Hardie, A., Cameron, F., Harvey, L., Stevenson, J., et al. (2016). Lessons learned from a prolonged and costly norovirus outbreak at a Scottish medicine of the elderly hospital: case study. The Journal of hospital infection *93*, 127-134.
- 92. Datta, D., Conway Morris, A., Antonelli, J., Warner, N., Brown, K.A., Wright, J., Simpson, A.J., Rennie, J., Hulme, G., Lewis, S.M., et al. (2016). Early PREdiction of Severe Sepsis (ExPRES-Sepsis) study: protocol for an observational derivation study to discover potential leucocyte cell surface biomarkers. BMJ open 6, e011335.

- 93. Davison, A., McDowell, G.S., Holden, J.M., Johnson, H.F., Koutsovoulos, G.D., Liu, M.M., Hulpiau, P., Van Roy, F., Wade, C.M., Banerjee, R., et al. (2016). Formin Is Associated with Left-Right Asymmetry in the Pond Snail and the Frog. Curr Biol *26*, 654-660.
- 94. de Vries, P.S., Chasman, D.I., Sabater-Lleal, M., Chen, M.H., Huffman, J.E., Steri, M., Tang, W., Teumer, A., Marioni, R.E., Grossmann, V., et al. (2016). A meta-analysis of 120 246 individuals identifies 18 new loci for fibrinogen concentration. Hum Mol Genet 25, 358-370.
- 95. Delannoy, C.M., Zadoks, R.N., Crumlish, M., Rodgers, D., Lainson, F.A., Ferguson, H.W., Turnbull, J., and Fontaine, M.C. (2016). Genomic comparison of virulent and non-virulent Streptococcus agalactiae in fish. Journal of fish diseases *39*, 13-29.
- 96. Denholm, S.J., McNeilly, T.N., Banos, G., Coffey, M.P., Russell, G.C., Bagnall, A., Mitchell, M.C., and Wall, E. (2017). Estimating genetic and phenotypic parameters of cellular immune-associated traits in dairy cows. J Dairy Sci *100*, 2850-2862.
- 97. Di Minno, G., Perno, C.F., Tiede, A., Navarro, D., Canaro, M., Guertler, L., and Ironside, J.W. (2016). Current concepts in the prevention of pathogen transmission via blood/plasma-derived products for bleeding disorders. Blood reviews *30*, 35-48.
- 98. Diack, A.B., Alibhai, J.D., Barron, R., Bradford, B., Piccardo, P., and Manson, J.C. (2015). Insights into Mechanisms of Chronic Neurodegeneration. International journal of molecular sciences *17*.
- 99. Diack, A.B., Alibhai, J.D., Barron, R., Bradford, B., Piccardo, P., and Manson, J.C. (2016). Insights into Mechanisms of Chronic Neurodegeneration. International journal of molecular sciences *17*.
- 100. Diallo, B., Sissoko, D., Loman, N.J., Bah, H.A., Bah, H., Worrell, M.C., Conde, L.S., Sacko, R., Mesfin, S., Loua, A., et al. (2016). Resurgence of Ebola Virus Disease in Guinea Linked to a Survivor With Virus Persistence in Seminal Fluid for More Than 500 Days. Clinical infectious diseases: an official publication of the Infectious Diseases Society of America *63*, 1353-1356.
- 101. Diehl, W.E., Lin, A.E., Grubaugh, N.D., Carvalho, L.M., Kim, K., Kyawe, P.P., McCauley, S.M., Donnard, E., Kucukural, A., McDonel, P., et al. (2016). Ebola Virus Glycoprotein with Increased Infectivity Dominated the 2013-2016 Epidemic. Cell *167*, 1088-1098.e1086.
- 102. Donaldson, D.S., and Mabbott, N.A. (2016). The influence of commensal and pathogenic gut microbiota on prion disease pathogenesis. J Gen Virol *97*, 725-738.
- 103. Donaldson, D.S., Sehgal, A., Rios, D., Williams, I.R., and Mabbott, N.A. (2016). Increased Abundance of M Cells in the Gut Epithelium Dramatically Enhances Oral Prion Disease Susceptibility. PLoS Pathog *12*, e1006075.
- 104. Donnelly, M.C., Imlach, S.N., Abravanel, F., Ramalingam, S., Johannessen, I., Petrik, J., Fraser, A.R., Campbell, J.D., Bramley, P., Dalton, H.R., et al. (2017). Sofosbuvir and Daclatasvir Anti-Viral Therapy Fails to Clear HEV Viremia and Restore Reactive T Cells in a HEV/HCV Co-Infected Liver Transplant Recipient. Gastroenterology 152, 300-301.
- 105. Dorward, D.A., Felton, J.M., Robb, C.T., Craven, T., Kipari, T., Walsh, T.S., Haslett, C., Kefala, K., Rossi, A.G., and Lucas, C.D. (2017). The cyclin-dependent kinase inhibitor AT7519 accelerates neutrophil apoptosis in sepsis-related acute respiratory distress syndrome. Thorax 72, 182-185.
- 106. Dransfield, I., and Farnworth, S. (2016). Axl and Mer Receptor Tyrosine Kinases: Distinct and Nonoverlapping Roles in Inflammation and Cancer? Advances in experimental medicine and biology 930, 113-132.
- 107. Drozdowska, J., Cousens, C., Finlayson, J., Collie, D., and Dagleish, M.P. (2016). Structural Development, Cellular Differentiation and Proliferation of the Respiratory Epithelium in the Bovine Fetal Lung. Journal of comparative pathology *154*, 42-56.
- 108. Duffin, R., O'Connor, R.A., Crittenden, S., Forster, T., Yu, C., Zheng, X., Smyth, D., Robb, C.T., Rossi, F., Skouras, C., et al. (2016). Prostaglandin E(2) constrains systemic inflammation through an innate lymphoid cell-IL-22 axis. Science *351*, 1333-1338.
- 109. Duffy, C., Venturato, A., Callanan, A., Lilienkampf, A., and Bradley, M. (2016). Arrays of 3D double-network hydrogels for the high-throughput discovery of materials with enhanced physical and biological properties. Acta biomaterialia *34*, 104-112.
- 110. Easton, S., Bartley, D.J., Hotchkiss, E., Hodgkinson, J.E., Pinchbeck, G.L., and Matthews, J.B. (2016). Use of a multiple choice questionnaire to assess UK prescribing channels' knowledge of helminthology and best practice surrounding anthelmintic use in livestock and horses. Prev Vet Med 128, 70-77.
- 111. Ekstrom, A.M., Clark, J., Byass, P., Lopez, A., De Savigny, D., Moyer, C.A., Campbell, H., Gage, A.J., Bocquier, P., AbouZahr, C., et al. (2016). INDEPTH Network: contributing to the data revolution. Lancet Diabetes Endocrinol *4*, 97.

- 112. El Tawil, S., Chohan, G., Mackenzie, J., Rowe, A., Weller, B., Will, R.G., and Knight, R. (2016). Isolated language impairment as the primary presentation of sporadic Creutzfeldt Jakob Disease. Acta neurologica Scandinavica *135*, 316-323.
- 113. Escamilla, A., Zafra, R., Perez, J., McNeilly, T.N., Pacheco, I.L., Buffoni, L., Martinez-Moreno, F.J., Molina-Hernandez, V., and Martinez-Moreno, A. (2016). Distribution of Foxp3+ T cells in the liver and hepatic lymph nodes of goats and sheep experimentally infected with Fasciola hepatica. Vet Parasitol *230*, 14-19.
- 114. Evenson, D., Gerber, P.F., Xiao, C.T., Halbur, P.G., Wang, C., Tian, D., Ni, Y.Y., Meng, X.J., and Opriessnig, T. (2016). A porcine reproductive and respiratory syndrome virus candidate vaccine based on the synthetic attenuated virus engineering approach is attenuated and effective in protecting against homologous virus challenge. Vaccine *34*, 5546-5553.
- 115. Eze, A.A., Gould, M.K., Munday, J.C., Tagoe, D.N., Stelmanis, V., Schnaufer, A., and De Koning, H.P. (2016). Reduced Mitochondrial Membrane Potential Is a Late Adaptation of Trypanosoma brucei brucei to Isometamidium Preceded by Mutations in the gamma Subunit of the F1Fo-ATPase. PLoS Negl Trop Dis 10, e0004791.
- 116. Fairfoul, G., McGuire, L.I., Pal, S., Ironside, J.W., Neumann, J., Christie, S., Joachim, C., Esiri, M., Evetts, S.G., Rolinski, M., et al. (2016). Alpha-synuclein RT-QuIC in the CSF of patients with alpha-synucleinopathies. Annals of clinical and translational neurology *3*, 812-818.
- 117. Fairlamb, A.H., Gow, N.A., Matthews, K.R., and Waters, A.P. (2016). Drug resistance in eukaryotic microorganisms. Nature microbiology *1*, 16092.
- 118. Fairlie, J., Holland, R., Pilkington, J.G., Pemberton, J.M., Harrington, L., and Nussey, D.H. (2016). Lifelong leukocyte telomere dynamics and survival in a free-living mammal. Aging cell *15*, 140-148.
- 119. Fan, J., Cui, J., Gerber, P.F., Biernacka, K., Stadejek, T., and Opriessnig, T. (2016). First Genome Sequences of Porcine Parvovirus 5 Strains Identified in Polish Pigs. Genome Announc 4.
- 120. Farre, M., Narayan, J., Slavov, G.T., Damas, J., Auvil, L., Li, C., Jarvis, E.D., Burt, D.W., Griffin, D.K., and Larkin, D.M. (2016). Novel Insights into Chromosome Evolution in Birds, Archosaurs, and Reptiles. Genome biology and evolution *8*, 2442-2451.
- 121. Fiegna, C., Clarke, C.L., Shaw, D.J., Baily, J.L., Clare, F.C., Gray, A., Garner, T.W., and Meredith, A.L. (2016). Pathological and phylogenetic characterization of Amphibiothecum sp. infection in an isolated amphibian (Lissotriton helveticus) population on the island of Rum (Scotland). Parasitology *22*, 1-13.
- 122. Fitchett, E.J., Seale, A.C., Vergnano, S., Sharland, M., Heath, P.T., Saha, S.K., Agarwal, R., Ayede, A.I., Bhutta, Z.A., Black, R., et al. (2016). Strengthening the Reporting of Observational Studies in Epidemiology for Newborn Infection (STROBE-NI): an extension of the STROBE statement for neonatal infection research. The Lancet. Infectious diseases *16*, e202-213.
- 123. Fitzgerald, J.R., and Holden, M.T. (2016). Genomics of Natural Populations of Staphylococcus aureus. Annual review of microbiology *8*, 459-478.
- 124. Fletcher, E., Pilizota, T., Davies, P.R., McVey, A., and French, C.E. (2016). Characterization of the effects of n-butanol on the cell envelope of E. coli. Applied microbiology and biotechnology *100*, 9653-9659.
- 125. Fontanella, E., Ma, Z., Zhang, Y., de Castro, A.M., Shen, H., Halbur, P.G., and Opriessnig, T. (2017). An interferon inducing porcine reproductive and respiratory syndrome virus vaccine candidate elicits protection against challenge with the heterologous virulent type 2 strain VR-2385 in pigs. Vaccine *35*, 125-131.
- 126. Forde, T., Biek, R., Zadoks, R., Workentine, M.L., De Buck, J., Kutz, S., Opriessnig, T., Trewby, H., van der Meer, F., and Orsel, K. (2016). Genomic analysis of the multi-host pathogen Erysipelothrix rhusiopathiae reveals extensive recombination as well as the existence of three generalist clades with wide geographic distribution. BMC Genomics *17*, 461.
- 127. French, A.S., Zadoks, R.N., Skuce, P.J., Mitchell, G., Gordon-Gibbs, D.K., Craine, A., Shaw, D., Gibb, S.W., and Taggart, M.A. (2016). Prevalence of Liver Fluke (Fasciola hepatica) in Wild Red Deer (Cervus elaphus): Coproantigen ELISA Is a Practicable Alternative to Faecal Egg Counting for Surveillance in Remote Populations. PLoS One *11*, e0162420.
- 128. Froy, H., Walling, C.A., Pemberton, J.M., Clutton-Brock, T.H., and Kruuk, L.E. (2016). Relative costs of offspring sex and offspring survival in a polygynous mammal. Biology letters *12*.
- 129. Fyfe, J., Picozzi, K., Waiswa, C., Bardosh, K.L., and Welburn, S.C. (2016). Impact of mass chemotherapy in domestic livestock for control of zoonotic T. b. rhodesiense human African trypanosomiasis in Eastern Uganda. Acta tropica *165*, 216-229.

- 130. Fyfe, J., Picozzi, K., Waiswa, C., Bardosh, K.L., and Welburn, S.C. (2017). Impact of mass chemotherapy in domestic livestock for control of zoonotic T. b. rhodesiense human African trypanosomiasis in Eastern Uganda. Acta tropica *165*, 216-229.
- 131. Gabaldon, T., Ginger, M.L., and Michels, P.A. (2016). Peroxisomes in parasitic protists. Mol Biochem Parasitol *209*, 35-45.
- 132. Gadalla, A.A., Schneider, P., Churcher, T.S., Nassir, E., Abdel-Muhsin, A.A., Ranford-Cartwright, L.C., Reece, S.E., and Babiker, H.A. (2016). Associations between Season and Gametocyte Dynamics in Chronic Plasmodium falciparum Infections. PLoS One *11*, e0166699.
- 133. Gadsby, N.J., Helgason, K.O., Dickson, E.M., Mills, J.M., Lindsay, D.S., Edwards, G.F., Hanson, M.F., and Templeton, K.E. (2016). Molecular diagnosis of Legionella infections--Clinical utility of front-line screening as part of a pneumonia diagnostic algorithm. The Journal of infection *72*, 161-170.
- 134. Gadsby, N.J., Russell, C.D., McHugh, M.P., Mark, H., Conway Morris, A., Laurenson, I.F., Hill, A.T., and Templeton, K.E. (2016). Reply to Zelyas and Robinson. Clinical infectious diseases: an official publication of the Infectious Diseases Society of America *63*, 142-143.
- 135. Gadsby, N.J., Russell, C.D., McHugh, M.P., Mark, H., Conway Morris, A., Laurenson, I.F., Hill, A.T., and Templeton, K.E. (2016). Comprehensive Molecular Testing for Respiratory Pathogens in Community-Acquired Pneumonia. Clinical infectious diseases: an official publication of the Infectious Diseases Society of America *62*, 817-823.
- 136. Garcia-Guzman, C., Fernandez, A., Avlonitis, N., Bradley, M., and Vendrell, M. (2016). Red-fluorescent Activatable Probes for the Detection of Hydrogen Peroxide in Living Cells. Combinatorial chemistry & high throughput screening *19*, 353-361.
- 137. Gaunt, E., Wise, H.M., Zhang, H., Lee, L.N., Atkinson, N.J., Nicol, M.Q., Highton, A.J., Klenerman, P., Beard, P.M., Dutia, B.M., et al. (2016). Elevation of CpG frequencies in influenza A genome attenuates pathogenicity but enhances host response to infection. Elife *5*, e12735.
- 138. Gause, W.C., and Maizels, R.M. (2016). Macrobiota helminths as active participants and partners of the microbiota in host intestinal homeostasis. Current opinion in microbiology *32*, 14-18.
- 139. Gerbe, F., Sidot, E., Smyth, D.J., Ohmoto, M., Matsumoto, I., Dardalhon, V., Cesses, P., Garnier, L., Pouzolles, M., Brulin, B., et al. (2016). Intestinal epithelial tuft cells initiate type 2 mucosal immunity to helminth parasites. Nature *529*, 226-230.
- 140. Gerber, P.F., Lelli, D., Zhang, J., Strandbygaard, B., Moreno, A., Lavazza, A., Perulli, S., Botner, A., Comtet, L., Roche, M., et al. (2016). Diagnostic evaluation of assays for detection of antibodies against porcine epidemic diarrhea virus (PEDV) in pigs exposed to different PEDV strains. Prev Vet Med 135, 87-94.
- 141. Gerber, P.F., Xiao, C.T., Lager, K., Crawford, K., Kulshreshtha, V., Cao, D., Meng, X.J., and Opriessnig, T. (2016). Increased frequency of porcine epidemic diarrhea virus shedding and lesions in suckling pigs compared to nursery pigs and protective immunity in nursery pigs after homologous re-challenge. Vet Res 47, 118.
- 142. Giakoumelou, S., Wheelhouse, N., Cuschieri, K., Entrican, G., Howie, S.E., and Horne, A.W. (2016). The role of infection in miscarriage. Human reproduction update *22*, 116-133.
- 143. Giannakopoulos, A., Valiakos, G., Papaspyropoulos, K., Dougas, G., Korou, L.M., Tasioudi, K.E., Fthenakis, G.C., Hutchings, M.R., Kaimaras, D., Tsokana, C.N., et al. (2016). Rabies outbreak in Greece during 2012-2014: use of Geographical Information System for analysis, risk assessment and control. Epidemiol Infect 144, 3068-3079.
- 144. Gibbons, C.L., van Bunnik, B.A., Blatchford, O., Robertson, C., Porphyre, T., Imrie, L., Wilson, J., Fitzgerald, J.R., Woolhouse, M.E., and Chase-Topping, M.E. (2016). Not just a matter of size: a hospital-level risk factor analysis of MRSA bacteraemia in Scotland. BMC Infect Dis 16, 222.
- 145. Goldmann, W., Marier, E., Stewart, P., Konold, T., Street, S., Langeveld, J., Windl, O., and Ortiz-Pelaez, A. (2016). Prion protein genotype survey confirms low frequency of scrapie-resistant K222 allele in British goat herds. Vet Rec *178*, 168.
- 146. Gonzalez-Fernandez, E., Avlonitis, N., Murray, A.F., Mount, A.R., and Bradley, M. (2016). Methylene blue not ferrocene: Optimal reporters for electrochemical detection of protease activity. Biosensors & bioelectronics *84*, 82-88.
- 147. Goplen, N.P., Saxena, V., Knudson, K.M., Schrum, A.G., Gil, D., Daniels, M.A., Zamoyska, R., and Teixeiro, E. (2016). IL-12 Signals through the TCR To Support CD8 Innate Immune Responses. Journal of immunology (Baltimore, Md.: 1950) 197, 2434-2443.

- 148. Grabert, K., Michoel, T., Karavolos, M.H., Clohisey, S., Baillie, J.K., Stevens, M.P., Freeman, T.C., Summers, K.M., and McColl, B.W. (2016). Microglial brain region-dependent diversity and selective regional sensitivities to aging. Nature neuroscience *19*, 504-516.
- 149. Graham, A.L., Nussey, D.H., Lloyd-Smith, J.O., Longbottom, D., Maley, M., Pemberton, J.M., Pilkington, J.G., Prager, K.C., Smith, L., Watt, K.A., et al. (2016). Exposure to viral and bacterial pathogens among Soay sheep (Ovis aries) of the St Kilda archipelago. Epidemiol Infect *19*, 353-361.
- 150. Granroth-Wilding, H.M., Daunt, F., Cunningham, E.J., and Burthe, S.J. (2016). Between-individual variation in nematode burden among juveniles in a wild host. Parasitology *144*, 1879-1888.
- 151. Grewal, J.S., McLuskey, K., Das, D., Myburgh, E., Wilkes, J., Brown, E., Lemgruber, L., Gould, M.K., Burchmore, R.J., Coombs, G.H., et al. (2016). PNT1 Is a C11 Cysteine Peptidase Essential for Replication of the Trypanosome Kinetoplast. The Journal of biological chemistry *291*, 9492-9500.
- 152. Griffith, D.M., Lewis, S., Rossi, A.G., Rennie, J., Salisbury, L., Merriweather, J.L., Templeton, K., and Walsh, T.S. (2016). Systemic inflammation after critical illness: relationship with physical recovery and exploration of potential mechanisms. Thorax *71*, 820-829.
- 153. Gripper, L.B., and Welburn, S.C. (2017). Neurocysticercosis infection and disease-A review. Acta tropica *166*, 218-224.
- 154. Guido, S., Katzer, F., Nanjiani, I., Milne, E., and Innes, E.A. (2016). Serology-Based Diagnostics for the Control of Bovine Neosporosis. Trends Parasitol *32*, 131-143.
- 155. Guo, Z., Gonzalez, J.F., Hernandez, J.N., McNeilly, T.N., Corripio-Miyar, Y., Frew, D., Morrison, T., Yu, P., and Li, R.W. (2016). Possible mechanisms of host resistance to Haemonchus contortus infection in sheep breeds native to the Canary Islands. Sci Rep *6*, 26200.
- 156. Haas, J., Schwarz, A., Korporal-Kuhnke, M., Jarius, S., and Wildemann, B. (2016). Myeloid dendritic cells exhibit defects in activation and function in patients with multiple sclerosis. Journal of neuroimmunology *301*, 53-60.
- 157. Hall, M.D., Woolhouse, M.E., and Rambaut, A. (2016). Using genomics data to reconstruct transmission trees during disease outbreaks. Rev Sci Tech *35*, 287-296.
- 158. Hall, M.D., Woolhouse, M.E., and Rambaut, A. (2016). The effects of sampling strategy on the quality of reconstruction of viral population dynamics using Bayesian skyline family coalescent methods: A simulation study. Virus evolution *2*, vew003.
- 159. Hamilton, C.A., Mahan, S., Entrican, G., and Hope, J.C. (2016). Interactions between natural killer cells and dendritic cells favour T helper1-type responses to BCG in calves. Vet Res *47*, 85.
- 160. Handel, I., Watt, K.A., Pilkington, J.G., Pemberton, J.M., Macrae, A., Scott, P., McNeilly, T.N., Berry, J.L., Clements, D.N., Nussey, D.H., et al. (2016). Vitamin D status predicts reproductive fitness in a wild sheep population. Sci Rep *6*, 18986.
- 161. Harder, M.J., Anliker, M., Hochsmann, B., Simmet, T., Huber-Lang, M., Schrezenmeier, H., Ricklin, D., Lambris, J.D., Barlow, P.N., and Schmidt, C.Q. (2016). Comparative Analysis of Novel Complement-Targeted Inhibitors, MiniFH, and the Natural Regulators Factor H and Factor H-like Protein 1 Reveal Functional Determinants of Complement Regulation. Journal of immunology (Baltimore, Md.: 1950) 196, 866-876.
- 162. Harijan, R.K., Mazet, M., Kiema, T.R., Bouyssou, G., Alexson, S.E., Bergmann, U., Moreau, P., Michels, P.A., Bringaud, F., and Wierenga, R.K. (2016). The SCP2-thiolase-like protein (SLP) of Trypanosoma brucei is an enzyme involved in lipid metabolism. Proteins *84*, 1075-1096.
- 163. Harrison, K., Haga, I.R., Pechenick Jowers, T., Jasim, S., Cintrat, J.C., Gillet, D., Schmitt-John, T., Digard, P., and Beard, P.M. (2016). Vaccinia virus uses retromer-independent cellular retrograde transport pathways to facilitate the wrapping of intracellular mature virions during viral morphogenesis. J Virol.
- 164. Harvey, S.R., MacPhee, C.E., Volkman, B.F., and Barran, P.E. (2016). The association and aggregation of the metamorphic chemokine lymphotactin with fondaparinux: from nm molecular complexes to mum molecular assemblies. Chem Commun (Camb) *52*, 394-397.
- 165. Helbert, M.R., Bangs, C., Bishop, M., Molesworth, A., and Ironside, J. (2016). No evidence of asymptomatic variant CJD infection in immunodeficiency patients treated with UK-sourced immunoglobulin. Vox Sang 110, 282-284.
- 166. Henihan, G., Schulze, H., Corrigan, D.K., Giraud, G., Terry, J.G., Hardie, A., Campbell, C.J., Walton, A.J., Crain, J., Pethig, R., et al. (2016). Label- and amplification-free electrochemical detection of bacterial ribosomal RNA. Biosensors & bioelectronics *81*, 487-494.
- 167. Hewitson, J.P., Nguyen, D.L., van Diepen, A., Smit, C.H., Koeleman, C.A., McSorley, H.J., Murray, J., Maizels, R.M., and Hokke, C.H. (2016). Novel O-linked methylated glycan antigens decorate secreted

- immunodominant glycoproteins from the intestinal nematode Heligmosomoides polygyrus. Int J Parasitol 46, 157-170.
- 168. Hill, A.T. (2016). Macrolides for Clinically Significant Bronchiectasis in Adults: Who Should Receive this Treatment? Chest *150*, 1187-1193.
- 169. Hogan, C., Iles, J., Frost, E.H., Giroux, G., Cassar, O., Gessain, A., Dion, M.J., Ilunga, V., Rambaut, A., Yengo-Ki-Ngimbi, A.E., et al. (2016). Epidemic history and iatrogenic transmission of blood-borne viruses in mid-20th century Kinshasa. J Infect Dis *214*, 353-360.
- 170. Holmes, E.C., Dudas, G., Rambaut, A., and Andersen, K.G. (2016). The evolution of Ebola virus: Insights from the 2013-2016 epidemic. Nature *538*, 193-200.
- 171. Homem, R.A., Le Bihan, T., Yu, M., and Loake, G.J. (2016). Identification of S-Nitrosothiols by the Sequential Cysteine Blocking Technique. Methods Mol Biol *1424*, 163-174.
- 172. Hoodless, L.J., Lucas, C.D., Duffin, R., Denvir, M.A., Haslett, C., Tucker, C.S., and Rossi, A.G. (2016). Genetic and pharmacological inhibition of CDK9 drives neutrophil apoptosis to resolve inflammation in zebrafish in vivo. Sci Rep *5*, 36980.
- 173. Hoodless, L.J., Robb, C.T., Felton, J.M., Tucker, C.S., and Rossi, A.G. (2016). Models for the Study of the Cross Talk Between Inflammation and Cell Cycle. Methods Mol Biol *1336*, 179-209.
- 174. Hooper, T.N., Inglis, R., Lorusso, G., Ujma, J., Barran, P.E., Uhrin, D., Schnack, J., Piligkos, S., Evangelisti, M., and Brechin, E.K. (2016). Structurally Flexible and Solution Stable [Ln4TM8(OH)8(L)8(O2CR)8(MeOH)y](ClO4)4: A Playground for Magnetic Refrigeration. Inorganic chemistry 55, 10535-10546.
- 175. Hopkins, J.L., and Bromley, G.E. (2016). Preparing New Graduates for Interprofessional Teamwork: Effectiveness of a Nurse Residency Program. Journal of continuing education in nursing *47*, 140-148.
- Hu, T., Wu, Z., Vervelde, L., Rothwell, L., Hume, D.A., and Kaiser, P. (2016). Functional annotation of the T cell immunoglobulin mucin (TIM) family in birds. Immunology *148*, 287-303.
- 177. Huerta-Uribe, A., Marjenberg, Z.R., Yamaguchi, N., Fitzgerald, S., Connolly, J.P., Carpena, N., Uvell, H., Douce, G., Elofsson, M., Byron, O., et al. (2016). Identification and Characterization of Novel Compounds Blocking Shiga Toxin Expression in Escherichia coli O157:H7. Front Microbiol 7, 1930.
- Huisman, J., Kruuk, L.E., Ellis, P.A., Clutton-Brock, T., and Pemberton, J.M. (2016). Inbreeding depression across the lifespan in a wild mammal population. Proc Natl Acad Sci U S A *113*, 3585-3590.
- 179. Hwang, S.H., Thielbeer, F., Jeong, J., Han, Y., Chankeshwara, S.V., Bradley, M., and Cho, W.S. (2016). Dual contribution of surface charge and protein-binding affinity to the cytotoxicity of polystyrene nanoparticles in nonphagocytic A549 cells and phagocytic THP-1 cells. Journal of toxicology and environmental health. Part A 79, 925-937.
- 180. Ikonomovic, M.D., Buckley, C.J., Heurling, K., Sherwin, P., Jones, P.A., Zanette, M., Mathis, C.A., Klunk, W.E., Chakrabarty, A., Ironside, J., et al. (2016). Post-mortem histopathology underlying beta-amyloid PET imaging following flutemetamol F 18 injection. Acta neuropathologica communications *4*, 130.
- 181. Indrigo, E., Clavadetscher, J., Chankeshwara, S.V., Lilienkampf, A., and Bradley, M. (2016). Palladium-mediated in situ synthesis of an anticancer agent. Chem Commun (Camb) *52*, 14212-14214.
- 182. Itzhaki, R.F., Lathe, R., Balin, B.J., Ball, M.J., Bearer, E.L., Braak, H., Bullido, M.J., Carter, C., Clerici, M., Cosby, S.L., et al. (2016). Microbes and Alzheimer's Disease. Journal of Alzheimer's disease: JAD *51*, 979-984.
- 183. Jackson-Jones, L.H., Duncan, S.M., Magalhaes, M.S., Campbell, S.M., Maizels, R.M., McSorley, H.J., Allen, J.E., and Benezech, C. (2016). Fat-associated lymphoid clusters control local IgM secretion during pleural infection and lung inflammation. Nat Commun 7, 12651.
- 184. Jackson-Jones, L.H., Ruckerl, D., Svedberg, F., Duncan, S., Maizels, R.M., Sutherland, T.E., Jenkins, S.J., McSorley, H.J., Benezech, C., MacDonald, A.S., et al. (2016). IL-33 delivery induces serous cavity macrophage proliferation independent of interleukin-4 receptor alpha. European journal of immunology 46, 2311-2321.
- 185. Jacobs, M., Rodger, A., Bell, D.J., Bhagani, S., Cropley, I., Filipe, A., Gifford, R.J., Hopkins, S., Hughes, J., Jabeen, F., et al. (2016). Late Ebola virus relapse causing meningoencephalitis: a case report. Lancet 388, 498-503.
- 186. Jensen, K., Gallagher, I.J., Kaliszewska, A., Zhang, C., Abejide, O., Gallagher, M.P., Werling, D., and Glass, E.J. (2016). Live and inactivated Salmonella enterica serovar Typhimurium stimulate similar but distinct transcriptome profiles in bovine macrophages and dendritic cells. Vet Res *47*, 46.
- 187. Jeroncic, A., Memari, Y., Ritchie, G.R., Hendricks, A.E., Kolb-Kokocinski, A., Matchan, A., Vitart, V., Hayward, C., Kolcic, I., Glodzik, D., et al. (2016). Whole-exome sequencing in an isolated population from the Dalmatian island of Vis. European journal of human genetics: EJHG *24*, 1479-1487.

- 188. Jimenez, M., and Bridle, H. (2016). Microfluidics for effective concentration and sorting of waterborne protozoan pathogens. Journal of microbiological methods *126*, 8-11.
- 189. Jing, L., Laing, K.J., Dong, L., Russell, R.M., Barlow, R.S., Haas, J.G., Ramchandani, M.S., Johnston, C., Buus, S., Redwood, A.J., et al. (2016). Extensive CD4 and CD8 T Cell Cross-Reactivity between Alphaherpesviruses. Journal of immunology (Baltimore, Md.: 1950) 196, 2205-2218.
- 190. Jitprasutwit, N., Zainal-Abidin, N., Vander Broek, C., Kurian, D., Korbsrisate, S., Stevens, M.P., and Stevens, J.M. (2016). Identification of Candidate Host Cell Factors Required for Actin-Based Motility of Burkholderia pseudomallei. Journal of proteome research *15*, 4675-4685.
- 191. Johansen, M.V., Welburn, S.C., Dorny, P., and Brattig, N.W. (2017). Control of neglected zoonotic diseases. Acta tropica *165*, 1-2.
- 192. Johnston, C.J., Smyth, D.J., Dresser, D.W., and Maizels, R.M. (2016). TGF-beta in tolerance, development and regulation of immunity. Cellular immunology *299*, 14-22.
- 193. Johnston, S.E., Berenos, C., Slate, J., and Pemberton, J.M. (2016). Conserved Genetic Architecture Underlying Individual Recombination Rate Variation in a Wild Population of Soay sheep (Ovis aries). Genetics 203, 583-598.
- 194. Jones, H.R., Robb, C.T., Perretti, M., and Rossi, A.G. (2016). The role of neutrophils in inflammation resolution. Semin Immunol *28*, 137-145.
- 195. Jones, R., McDonald, K.E., Willson, J.A., Ghesquiere, B., Sammut, D., Daniel, E., Harris, A.J., Lewis, A., Thompson, A.A., Dickinson, R.S., et al. (2016). Mutations in succinate dehydrogenase B (SDHB) enhance neutrophil survival independent of HIF-1alpha expression. Blood *127*, 2641-2644.
- 196. Jorgensen, H.J., Nordstoga, A.B., Sviland, S., Zadoks, R.N., Solverod, L., Kvitle, B., and Mork, T. (2016). Streptococcus agalactiae in the environment of bovine dairy herds rewriting the textbooks? Vet Microbiol *184*, 64-72.
- 197. Jorgensen, H.J., Nordstoga, A.B., Sviland, S., Zadoks, R.N., Solverod, L., Kvitle, B., and Mork, T. (2016). Streptococcus agalactiae in the environment of bovine dairy herds--rewriting the textbooks? Vet Microbiol *184*, 64-72.
- 198. Joshi, P.K., Fischer, K., Schraut, K.E., Campbell, H., Esko, T., and Wilson, J.F. (2016). Variants near CHRNA3/5 and APOE have age- and sex-related effects on human lifespan. Nat Commun *7*, 11174.
- 199. Jubb, A.W., Young, R.S., Hume, D.A., and Bickmore, W.A. (2016). Enhancer Turnover Is Associated with a Divergent Transcriptional Response to Glucocorticoid in Mouse and Human Macrophages. Journal of immunology (Baltimore, Md.: 1950) *196*, 813-822.
- 200. Kasprzak, M.M., Houdijk, J.G., Liddell, S., Davis, K., Olukosi, O.A., Kightley, S., White, G.A., and Wiseman, J. (2016). Rapeseed napin and cruciferin are readily digested by poultry. Journal of animal physiology and animal nutrition.
- 201. Kaur, S., Raggatt, L.J., Batoon, L., Hume, D.A., Levesque, J.P., and Pettit, A.R. (2016). Role of bone marrow macrophages in controlling homeostasis and repair in bone and bone marrow niches. Seminars in cell & developmental biology *61*, 12-21.
- 202. Keogh, M.J., Wei, W., Wilson, I., Coxhead, J., Ryan, S., Rollinson, S., Griffin, H., Kurzawa-Akanbi, M., Santibanez-Koref, M., Talbot, K., et al. (2017). Genetic compendium of 1511 human brains available through the UK Medical Research Council Brain Banks Network Resource. Genome research *27*, 165-173.
- 203. Kerrin, A., Fitch, P., Errington, C., Kerr, D., Waxman, L., Riding, K., McCormack, J., Mehendele, F., McSorley, H., MacKenzie, K., et al. (2016). Differential lower airway dendritic cell patterns may reveal distinct endotypes of RSV bronchiolitis. Thorax.
- 204. Kieback, E., Hilgenberg, E., Stervbo, U., Lampropoulou, V., Shen, P., Bunse, M., Jaimes, Y., Boudinot, P., Radbruch, A., Klemm, U., et al. (2016). Thymus-Derived Regulatory T Cells Are Positively Selected on Natural Self-Antigen through Cognate Interactions of High Functional Avidity. Immunity 44, 1114-1126.
- 205. Kokki, I., Smith, D., Simmonds, P., Ramalingam, S., Wellington, L., Willocks, L., Johannessen, I., and Harvala, H. (2016). Hepatitis E virus is the leading cause of acute viral hepatitis in Lothian, Scotland. New microbes and new infections *10*, 6-12.
- 206. Kraemer, S.A., Morgan, A.D., Ness, R.W., Keightley, P.D., and Colegrave, N. (2016). Fitness effects of new mutations in Chlamydomonas reinhardtii across two stress gradients. Journal of evolutionary biology *29*, 583-593.
- 207. Krstajic, N., Akram, A.R., Choudhary, T.R., McDonald, N., Tanner, M.G., Pedretti, E., Dalgarno, P.A., Scholefield, E., Girkin, J.M., Moore, A., et al. (2016). Two-color widefield fluorescence microendoscopy enables multiplexed molecular imaging in the alveolar space of human lung tissue. Journal of biomedical optics *21*, 46009.

- 208. Ladyman, M.K., Walton, J.G., Lilienkampf, A., and Bradley, M. (2016). Fluorescent Formazans and Tetrazolium Salts Towards Fluorescent Cytotoxicity Assays. Combinatorial chemistry & high throughput screening *19*, 384-391.
- 209. Lafond, K.E., Nair, H., Rasooly, M.H., Valente, F., Booy, R., Rahman, M., Kitsutani, P., Yu, H., Guzman, G., Coulibaly, D., et al. (2016). Global Role and Burden of Influenza in Pediatric Respiratory Hospitalizations, 1982-2012: A Systematic Analysis. PLoS Med *13*, e1001977.
- 210. Lathe, R., and Haas, J.G. (2016). Distribution of cellular HSV-1 receptor expression in human brain. Journal of neurovirology.
- 211. Laxminarayan, R., Amabile-Cuevas, C.F., Cars, O., Evans, T., Heymann, D.L., Hoffman, S., Holmes, A., Mendelson, M., Sridhar, D., Woolhouse, M., et al. (2016). UN High-Level Meeting on antimicrobials—what do we need? Lancet *388*, 218-220.
- 212. Laxminarayan, R., Sridhar, D., Blaser, M., Wang, M., and Woolhouse, M. (2016). Achieving global targets for antimicrobial resistance. Science *353*, 874-875.
- 213. Leitch, C.S., Natafji, E., Yu, C., Abdul-Ghaffar, S., Madarasingha, N., Venables, Z.C., Chu, R., Fitch, P.M., Muinonen-Martin, A.J., Campbell, L.E., et al. (2016). Filaggrin-null mutations are associated with increased maturation markers on Langerhans cells. The Journal of allergy and clinical immunology *138*, 482-490.
- 214. Lemay, J.F., Marguerat, S., Larochelle, M., Liu, X., van Nues, R., Hunyadkurti, J., Hoque, M., Tian, B., Granneman, S., Bahler, J., et al. (2016). The Nrd1-like protein Seb1 coordinates cotranscriptional 3' end processing and polyadenylation site selection. Genes & development *30*, 1558-1572.
- 215. Lewis, S.C., Bhattacharya, S., Wu, O., Vincent, K., Jack, S.A., Critchley, H.O., Porter, M.A., Cranley, D., Wilson, J.A., and Horne, A.W. (2016). Gabapentin for the Management of Chronic Pelvic Pain in Women (GaPP1): A Pilot Randomised Controlled Trial. PLoS One *11*, e0153037.
- 216. Lewis, S.H., Salmela, H., and Obbard, D.J. (2016). Duplication and diversification of Dipteran Argonaute genes, and the evolutionary divergence of Piwi and Aubergine. Genome biology and evolution *8*, 507-518.
- 217. Lewis, S.H., Webster, C.L., Salmela, H., and Obbard, D.J. (2016). Repeated Duplication of Argonaute2 Is Associated with Strong Selection and Testis Specialization in Drosophila. Genetics *204*, 757-769.
- 218. Li, X., Song, P., Timofeeva, M., Meng, X., Rudan, I., Little, J., Satsangi, J., Campbell, H., and Theodoratou, E. (2016). Systematic meta-analyses and field synopsis of genetic and epigenetic studies in paediatric inflammatory bowel disease. Sci Rep *6*, 34076.
- 219. Liu, W., Sundararaman, S.A., Loy, D.E., Learn, G.H., Li, Y., Plenderleith, L.J., Ndjango, J.B., Speede, S., Atencia, R., Cox, D., et al. (2016). Multigenomic Delineation of Plasmodium Species of the Laverania Subgenus Infecting Wild-Living Chimpanzees and Gorillas. Genome biology and evolution 8, 1929-1939.
- 220. Logan, C.J., Kruuk, L.E., Stanley, R., Thompson, A.M., and Clutton-Brock, T.H. (2016). Endocranial volume is heritable and is associated with longevity and fitness in a wild mammal. R Soc Open Sci *3*, 160622.
- 221. Longdon, B., Day, J.P., Schulz, N., Leftwich, P.T., de Jong, M.A., Breuker, C.J., Gibbs, M., Obbard, D.J., Wilfert, L., Smith, S.C., et al. (2017). Vertically transmitted rhabdoviruses are found across three insect families and have dynamic interactions with their hosts. Proc Biol Sci 284.
- Longdon, B., Murray, G.G., Palmer, W.J., Day, J.P., Parker, D.J., Welch, J.J., Obbard, D.J., and Jiggins, F.M. (2015). The evolution, diversity, and host associations of rhabdoviruses. Virus evolution 1, vev014.
- 223. Lorenzo-Redondo, R., Fryer, H.R., Bedford, T., Kim, E.Y., Archer, J., Kosakovsky Pond, S.L., Chung, Y.S., Penugonda, S., Chipman, J.G., Fletcher, C.V., et al. (2016). Persistent HIV-1 replication maintains the tissue reservoir during therapy. Nature *530*, 51-56.
- 224. Lorusso, V., Wijnveld, M., Latrofa, M.S., Fajinmi, A., Majekodunmi, A.O., Dogo, A.G., Igweh, A.C., Otranto, D., Jongejan, F., Welburn, S.C., et al. (2016). Canine and ovine tick-borne pathogens in camels, Nigeria. Vet Parasitol *228*, 90-92.
- 225. Lorusso, V., Wijnveld, M., Majekodunmi, A.O., Dongkum, C., Fajinmi, A., Dogo, A.G., Thrusfield, M., Mugenyi, A., Vaumourin, E., Igweh, A.C., et al. (2016). Tick-borne pathogens of zoonotic and veterinary importance in Nigerian cattle. Parasit Vectors *9*, 217.
- Loy, D.E., Liu, W., Li, Y., Learn, G.H., Plenderleith, L.J., Sundararaman, S.A., Sharp, P.M., and Hahn, B.H. (2016). Out of Africa: origins and evolution of the human malaria parasites Plasmodium falciparum and Plasmodium vivax. Int J Parasitol *47*, 87-97.
- 227. Lu, L., Van Dung, N., Bryant, J.E., Carrique-Mas, J., Van Cuong, N., Anh, P.H., Rabaa, M.A., Baker, S., Simmonds, P., and Woolhouse, M.E. (2016). Evolution and phylogeographic dissemination of endemic porcine picornaviruses in Vietnam. Virus evolution *2*, vew001.

- 228. Lu, Y., Day, F.R., Gustafsson, S., Buchkovich, M.L., Na, J., Bataille, V., Cousminer, D.L., Dastani, Z., Drong, A.W., Esko, T., et al. (2016). New loci for body fat percentage reveal link between adiposity and cardiometabolic disease risk. Nat Commun 7, 10495.
- 229. Luis, T.C., Luc, S., Mizukami, T., Boukarabila, H., Thongjuea, S., Woll, P.S., Azzoni, E., Giustacchini, A., Lutteropp, M., Bouriez-Jones, T., et al. (2016). Initial seeding of the embryonic thymus by immune-restricted lympho-myeloid progenitors. Nat Immunol *17*, 1424-1435.
- 230. Lundie, R.J., Webb, L.M., Marley, A.K., Phythian-Adams, A.T., Cook, P.C., Jackson-Jones, L.H., Brown, S., Maizels, R.M., Boon, L., O'Keeffe, M., et al. (2016). A central role for hepatic conventional dendritic cells in supporting Th2 responses during helminth infection. Immunol Cell Biol *94*, 400-410.
- 231. Lupolova, N., Dallman, T.J., Matthews, L., Bono, J.L., and Gally, D.L. (2016). Support vector machine applied to predict the zoonotic potential of E. coli O157 cattle isolates. Proc Natl Acad Sci U S A *113*, 11312-11317.
- 232. Mabbott, N.A. (2016). From Scientific Curiosity to Public Enemy Number One in Six Short Months. PLoS Pathog *12*, e1005371.
- 233. Mackay, M., Perez-Lopez, A.M., Bradley, M., and Lilienkampf, A. (2016). Eliminating caspase-7 and cathepsin B cross-reactivity on fluorogenic caspase-3 substrates. Molecular bioSystems *12*, 693-696.
- 234. Mackley, E.C., Houston, S., Marriott, C.L., Halford, E.E., Lucas, B., Cerovic, V., Filbey, K.J., Maizels, R.M., Hepworth, M.R., Sonnenberg, G.F., et al. (2016). Corrigendum: CCR7-dependent trafficking of RORgamma(+) ILCs creates a unique microenvironment within mucosal draining lymph nodes. Nat Commun 7, 11186.
- 235. Mainda, G., Lupolova, N., Sikakwa, L., Bessell, P.R., Muma, J.B., Hoyle, D.V., McAteer, S.P., Gibbs, K., Williams, N.J., Sheppard, S.K., et al. (2016). Phylogenomic approaches to determine the zoonotic potential of Shiga toxin-producing Escherichia coli (STEC) isolated from Zambian dairy cattle. Sci Rep 6, 26589.
- 236. Maizels, R.M., and McSorley, H.J. (2016). Regulation of the host immune system by helminth parasites. The Journal of allergy and clinical immunology *138*, 666-675.
- 237. Mapholi, N.O., Maiwashe, A., Matika, O., Riggio, V., Bishop, S.C., MacNeil, M.D., Banga, C., Taylor, J.F., and Dzama, K. (2016). Genome-wide association study of tick resistance in South African Nguni cattle. Ticks and tick-borne diseases 7, 487-497.
- 238. Marshall, J., Rossez, Y., Mainda, G., Gally, D.L., Daniell, T.J., and Holden, N.J. (2016). Alternate thermoregulation and functional binding of Escherichia coli type 1 fimbriae in environmental and animal isolates. FEMS microbiology letters *363*.
- 239. Mason, A.S., Fulton, J.E., Hocking, P.M., and Burt, D.W. (2016). A new look at the LTR retrotransposon content of the chicken genome. BMC Genomics *17*, 688.
- 240. Matika, O., Riggio, V., Anselme-Moizan, M., Law, A.S., Pong-Wong, R., Archibald, A.L., and Bishop, S.C. (2016). Genome-wide association reveals QTL for growth, bone and in vivo carcass traits as assessed by computed tomography in Scottish Blackface lambs. Genetics, selection, evolution: GSE 48, 11.
- 241. Matthews, J.B., Geldhof, P., Tzelos, T., and Claerebout, E. (2016). Progress in the development of subunit vaccines for gastrointestinal nematodes of ruminants. Parasite immunology *38*, 744-753.
- 242. Matzinger, S.R., Opriessnig, T., Xiao, C.T., Catanzaro, N., Beach, N.M., Slade, D.E., Nitzel, G.P., and Meng, X.J. (2016). A chimeric virus created by DNA shuffling of the capsid genes of different subtypes of porcine circovirus type 2 (PCV2) in the backbone of the non-pathogenic PCV1 induces protective immunity against the predominant PCV2b and the emerging PCV2d in pigs. Virology 498, 82-93.
- 243. Mazeri, S., Sargison, N., Kelly, R.F., Bronsvoort, B.M., and Handel, I. (2016). Evaluation of the Performance of Five Diagnostic Tests for Fasciola hepatica Infection in Naturally Infected Cattle Using a Bayesian No Gold Standard Approach. PLoS One *11*, e0161621.
- 244. McGlasson, S.L., Semple, F., MacPherson, H., Gray, M., Davidson, D.J., and Dorin, J.R. (2017). Human beta-defensin 3 increases the TLR9-dependent response to bacterial DNA. European journal of immunology.
- 245. McLean, C.J., Marles-Wright, J., Custodio, R., Lowther, J., Kennedy, A.J., Pollock, J., Clarke, D.J., Brown, A.R., and Campopiano, D.J. (2017). Characterization of homologous sphingosine-1-phosphate lyase isoforms in the bacterial pathogen Burkholderia pseudomallei. Journal of lipid research *58*, 137-150.
- 246. McNally, L., and Brown, S.P. (2016). Microbiome: Ecology of stable gut communities. Nature microbiology *1*, 15016.
- 247. Meek, S., Thomson, A.J., Sutherland, L., Sharp, M.G., Thomson, J., Bishop, V., Meddle, S.L., Gloaguen, Y., Weidt, S., Singh-Dolt, K., et al. (2016). Reduced levels of dopamine and altered metabolism in brains of HPRT knock-out rats: a new rodent model of Lesch-Nyhan Disease. Sci Rep *6*, 25592.

- 248. Melzi, E., Caporale, M., Rocchi, M., Martin, V., Gamino, V., di Provvido, A., Marruchella, G., Entrican, G., Sevilla, N., and Palmarini, M. (2016). Follicular dendritic cell disruption as a novel mechanism of virus-induced immunosuppression. Proc Natl Acad Sci U S A *113*, E6238-e6247.
- 249. Mena, I., Nelson, M.I., Quezada-Monroy, F., Dutta, J., Cortes-Fernandez, R., Lara-Puente, J.H., Castro-Peralta, F., Cunha, L.F., Trovao, N.S., Lozano-Dubernard, B., et al. (2016). Origins of the 2009 H1N1 influenza pandemic in swine in Mexico. Elife 28.
- 250. Meng, L.S., Wang, Y.B., Loake, G.J., and Jiang, J.H. (2016). Seed Embryo Development Is Regulated via an AN3-MINI3 Gene Cascade. Frontiers in plant science *7*, 1645.
- 251. Mielczarek, W.S., Obaje, E.A., Bachmann, T.T., and Kersaudy-Kerhoas, M. (2016). Microfluidic blood plasma separation for medical diagnostics: is it worth it? Lab Chip *16*, 3441-3448.
- 252. Miesen, P., Ivens, A., Buck, A.H., and van Rij, R.P. (2016). Small RNA Profiling in Dengue Virus 2-Infected Aedes Mosquito Cells Reveals Viral piRNAs and Novel Host miRNAs. PLoS Negl Trop Dis *10*, e0004452.
- 253. Miller, B., Jimenez, M., and Bridle, H. (2016). Cascading and Parallelising Curvilinear Inertial Focusing Systems for High Volume, Wide Size Distribution, Separation and Concentration of Particles. Sci Rep *6*, 36386.
- 254. Minikel, E.V., Vallabh, S.M., Lek, M., Estrada, K., Samocha, K.E., Sathirapongsasuti, J.F., McLean, C.Y., Tung, J.Y., Yu, L.P., Gambetti, P., et al. (2016). Quantifying prion disease penetrance using large population control cohorts. Science translational medicine *8*, 322ra329.
- 255. Minutti, C.M., Knipper, J.A., Allen, J.E., and Zaiss, D.M. (2016). Tissue-specific contribution of macrophages to wound healing. Seminars in cell & developmental biology *61*, 3-11.
- 256. Mitchell, M.C., Tzelos, T., Handel, I., McWilliam, H.E., Hodgkinson, J.E., Nisbet, A.J., Kharchenko, V.O., Burgess, S.T., and Matthews, J.B. (2016). Development of a recombinant protein-based ELISA for diagnosis of larval cyathostomin infection. Parasitology *143*, 1055-1066.
- 257. Mole, D.J., Webster, S.P., Uings, I., Zheng, X., Binnie, M., Wilson, K., Hutchinson, J.P., Mirguet, O., Walker, A., Beaufils, B., et al. (2016). Kynurenine-3-monooxygenase inhibition prevents multiple organ failure in rodent models of acute pancreatitis. Nature medicine *22*, 202-209.
- 258. Molina, O., Carmena, M., Maudlin, I.E., and Earnshaw, W.C. (2016). PREditOR: a synthetic biology approach to removing heterochromatin from cells. Chromosome research: an international journal on the molecular, supramolecular and evolutionary aspects of chromosome biology *24*, 495-509.
- 259. Montazeri, Z., Theodoratou, E., Nyiraneza, C., Timofeeva, M., Chen, W., Svinti, V., Sivakumaran, S., Gresham, G., Cubitt, L., Carvajal-Carmona, L., et al. (2016). Systematic meta-analyses and field synopsis of genetic association studies in colorectal adenomas. International journal of epidemiology *45*, 186-205.
- 260. Moorad, J.A., and Nussey, D.H. (2016). Evolution of maternal effect senescence. Proc Natl Acad Sci U S A *113*, 362-367.
- 261. Moore, R.A., Choi, Y.P., Head, M.W., Ironside, J.W., Faris, R., Ritchie, D.L., Zanusso, G., and Priola, S.A. (2016). Relative Abundance of apoE and Abeta1-42 Associated with Abnormal Prion Protein Differs between Creutzfeldt-Jakob Disease Subtypes. Journal of proteome research *15*, 4518-4531.
- 262. Moore, R.A., Head, M.W., Ironside, J.W., Ritchie, D.L., Zanusso, G., Pyo Choi, Y., and Priola, S.A. (2016). The Distribution of Prion Protein Allotypes Differs Between Sporadic and latrogenic Creutzfeldt-Jakob Disease Patients. PLoS Pathog *12*, e1005416.
- 263. Morrison, L.J., Vezza, L., Rowan, T., and Hope, J.C. (2016). Animal African Trypanosomiasis: Time to Increase Focus on Clinically Relevant Parasite and Host Species. Trends Parasitol *23*, 599-607.
- 264. Morton, J.A., and Bridle, H. (2016). Student-led microfluidics lab practicals: Improving engagement and learning outcomes. Biomicrofluidics *10*, 034117.
- 265. Moura, A.S., Ledur, M.C., Boschiero, C., Nones, K., Pinto, L.F., Jaenisch, F.R., Burt, D.W., and Coutinho, L.L. (2016). Quantitative trait loci with sex-specific effects for internal organs weights and hematocrit value in a broiler-layer cross. Journal of applied genetics *57*, 215-224.
- 266. Murray, G.G., Wang, F., Harrison, E.M., Paterson, G.K., Mather, A.E., Harris, S.R., Holmes, M.A., Rambaut, A., and Welch, J.J. (2016). The effect of genetic structure on molecular dating and tests for temporal signal. Methods Ecol Evol *7*, 80-89.
- 267. Mutapi, F. (2016). Getting a GRiPP on everyday schistosomiasis: experience from Zimbabwe. Parasitology, 1-9.
- 268. Mutapi, F., Maizels, R., Fenwick, A., and Woolhouse, M. (2017). Human schistosomiasis in the post mass drug administration era. The Lancet. Infectious diseases *17*, e42-e48.
- 269. Nair, H. (2017). Simplified antibiotic regimens for community management of neonatal sepsis. Lancet Glob Health *5*, e118-e120.

- 270. Nair, H., Watts, A.T., Williams, L.J., Omer, S.B., Simpson, C.R., Willocks, L.J., Cameron, J.C., and Campbell, H. (2016). Pneumonia hospitalisations in Scotland following the introduction of pneumococcal conjugate vaccination in young children. BMC Infect Dis *16*, 390.
- 271. Nandi, S., Whyte, J., Taylor, L., Sherman, A., Nair, V., Kaiser, P., and McGrew, M.J. (2016). Cryopreservation of specialized chicken lines using cultured primordial germ cells. Poultry science *95*, 1905-1911.
- 272. Ness, R.W., Kraemer, S.A., Colegrave, N., and Keightley, P.D. (2016). Direct Estimate of the Spontaneous Mutation Rate Uncovers the Effects of Drift and Recombination in the Chlamydomonas reinhardtii Plastid Genome. Mol Biol Evol *33*, 800-808.
- 273. Neumann, K., Jain, S., Gambardella, A., Walker, S.E., Valero, E., Lilienkampf, A., and Bradley, M. (2017). Tetrazine-Responsive Self-immolative Linkers. Chembiochem: a European journal of chemical biology 18, 91-95.
- 274. Neumann, K., Jain, S., Geng, J., and Bradley, M. (2016). Nanoparticle "switch-on" by tetrazine triggering. Chem Commun (Camb) *52*, 11223-11226.
- 275. Nickbakhsh, S., Hall, M.D., Dorigatti, I., Lycett, S.J., Mulatti, P., Monne, I., Fusaro, A., Woolhouse, M.E., Rambaut, A., and Kao, R.R. (2016). Modelling the impact of co-circulating low pathogenic avian influenza viruses on epidemics of highly pathogenic avian influenza in poultry. Epidemics 17, 27-34.
- 276. Nicol, L., Gossner, A., Watkins, C., Chianini, F., Dalziel, R., and Hopkins, J. (2016). Variations in IL-23 and IL-25 receptor gene structure, sequence and expression associated with the two disease forms of sheep paratuberculosis. Vet Res *47*, 27.
- 277. Nicol, L., Wilkie, H., Gossner, A., Watkins, C., Dalziel, R., and Hopkins, J. (2016). Variations in T cell transcription factor gene structure and expression associated with the two disease forms of sheep paratuberculosis. Vet Res *47*, 83.
- 278. Nielsen, M.K., von Samson-Himmelstjerna, G., Pfister, K., Reinemeyer, C.R., Molento, M.B., Peregrine, A.S., Hodgkinson, J.E., Jacobsen, S., Kaplan, R.M., and Matthews, J.B. (2016). The appropriate antiparasitic treatment: Coping with emerging threats from old adversaries. Equine veterinary journal 48, 374-375.
- 279. Nightingale, S., Geretti, A.M., Beloukas, A., Fisher, M., Winston, A., Else, L., Nelson, M., Taylor, S., Ustianowski, A., Ainsworth, J., et al. (2016). Discordant CSF/plasma HIV-1 RNA in patients with unexplained low-level viraemia. Journal of neurovirology *22*, 852-860.
- 280. Nightingale, S., Michael, B.D., Fisher, M., Winston, A., Nelson, M., Taylor, S., Ustianowski, A., Ainsworth, J., Gilson, R., Haddow, L., et al. (2016). CSF/plasma HIV-1 RNA discordance even at low levels is associated with up-regulation of host inflammatory mediators in CSF. Cytokine *83*, 139-146.
- 281. Nisbet, A.J., McNeilly, T.N., Greer, A.W., Bartley, Y., Oliver, E.M., Smith, S., Palarea-Albaladejo, J., and Matthews, J.B. (2016). Protection of ewes against Teladorsagia circumcincta infection in the periparturient period by vaccination with recombinant antigens. Vet Parasitol *228*, 130-136.
- 282. Nisbet, A.J., Meeusen, E.N., Gonzalez, J.F., and Piedrafita, D.M. (2016). Immunity to Haemonchus contortus and Vaccine Development. Advances in parasitology *93*, 353-396.
- 283. Noho-Konteh, F., Adetifa, J.U., Cox, M., Hossin, S., Reynolds, J., Le, M.T., Sanyang, L.C., Drammeh, A., Plebanski, M., Forster, T., et al. (2016). Sex-Differential Non-Vaccine-Specific Immunological Effects of Diphtheria-Tetanus-Pertussis and Measles Vaccination. Clinical infectious diseases: an official publication of the Infectious Diseases Society of America *63*, 1213-1226.
- 284. Nolan, T.J., Gadsby, N.J., Hellyer, T.P., Templeton, K.E., McMullan, R., McKenna, J.P., Rennie, J., Robb, C.T., Walsh, T.S., Rossi, A.G., et al. (2016). Low-pathogenicity Mycoplasma spp. alter human monocyte and macrophage function and are highly prevalent among patients with ventilator-acquired pneumonia. Thorax *71*, 594-600.
- 285. Nowell, R.W., Laue, B.E., Sharp, P.M., and Green, S. (2016). Comparative genomics reveals genes significantly associated with woody hosts in the plant pathogen Pseudomonas syringae. Molecular plant pathology *17*, 1409-1424.
- 286. Nurnberger, B., Lohse, K., Fijarczyk, A., Szymura, J.M., and Blaxter, M.L. (2016). Para-allopatry in hybridizing fire-bellied toads (Bombina bombina and B. variegata): Inference from transcriptome-wide coalescence analyses. Evolution; international journal of organic evolution *70*, 2429.
- 287. O'Hara, L., Livigni, A., Theo, T., Boyer, B., Angus, T., Wright, D., Chen, S.H., Raza, S., Barnett, M.W., Digard, P., et al. (2016). Modelling the Structure and Dynamics of Biological Pathways. PLoS Biol *14*, e1002530.

- 288. O'Neill, K.E., Bredenkamp, N., Tischner, C., Vaidya, H.J., Stenhouse, F.H., Peddie, C.D., Nowell, C.S., Gaskell, T., and Blackburn, C.C. (2016). Foxn1 Is Dynamically Regulated in Thymic Epithelial Cells during Embryogenesis and at the Onset of Thymic Involution. PLoS One *11*, e0151666.
- 289. Obieglo, K., Feng, X., Bollampalli, V.P., Dellacasa-Lindberg, I., Classon, C., Osterblad, M., Helmby, H., Hewitson, J.P., Maizels, R.M., Gigliotti Rothfuchs, A., et al. (2016). Chronic Gastrointestinal Nematode Infection Mutes Immune Responses to Mycobacterial Infection Distal to the Gut. Journal of immunology (Baltimore, Md.: 1950) *196*, 2262-2271.
- 290. Ochoa-Villarreal, M., Howat, S., Hong, S., Jang, M.O., Jin, Y.W., Lee, E.K., and Loake, G.J. (2016). Plant cell culture strategies for the production of natural products. BMB reports *49*, 149-158.
- 291. Opi, D.H., Uyoga, S., Orori, E.N., Williams, T.N., and Rowe, J.A. (2016). Red blood cell complement receptor one level varies with Knops blood group, alpha(+)thalassaemia and age among Kenyan children. Genes and immunity *17*, 171-178.
- 292. Opriessnig, T., Xiao, C.T., Halbur, P.G., Gerber, P.F., Matzinger, S.R., and Meng, X.J. (2017). A commercial porcine circovirus (PCV) type 2a-based vaccine reduces PCV2d viremia and shedding and prevents PCV2d transmission to naive pigs under experimental conditions. Vaccine *35*, 248-254.
- 293. Orlando, G., Law, P.J., Palin, K., Tuupanen, S., Gylfe, A., Hanninen, U.A., Cajuso, T., Tanskanen, T., Kondelin, J., Kaasinen, E., et al. (2016). Variation at 2q35 (PNKD and TMBIM1) influences colorectal cancer risk and identifies a pleiotropic effect with inflammatory bowel disease. Hum Mol Genet *25*, 2349-2359.
- 294. Orsini, L., Gilbert, D., Podicheti, R., Jansen, M., Brown, J.B., Solari, O.S., Spanier, K.I., Colbourne, J.K., Rush, D., Decaestecker, E., et al. (2016). Daphnia magna transcriptome by RNA-Seq across 12 environmental stressors. Scientific data *3*, 160030.
- 295. Oshita, M., Yokoyama, T., Takei, Y., Takeuchi, A., Ironside, J.W., Kitamoto, T., and Morita, M. (2016). Efficient propagation of variant Creutzfeldt-Jakob disease prion protein using the cell-protein misfolding cyclic amplification technique with samples containing plasma and heparin. Transfusion *56*, 223-230.
- 296. Palani Velu, P., Bedi, P., Turnbull, K., and Hill, A.T. (2016). Risk Factors for Requiring Intravenous Antibiotic Therapy in Hospital and 30-Day Readmission for Exacerbations of Bronchiectasis. QJM: monthly journal of the Association of Physicians 109, 723-729.
- 297. Palmer, W.H., and Obbard, D.J. (2016). Variation and Evolution in the Glutamine-Rich Repeat Region of Drosophila Argonaute-2. G3 (Bethesda, Md.) *6*, 2563-2572.
- 298. Pan, Q., Cui, B., Deng, F., Quan, J., Loake, G.J., and Shan, W. (2016). RTP1 encodes a novel endoplasmic reticulum (ER)-localized protein in Arabidopsis and negatively regulates resistance against biotrophic pathogens. The New phytologist *209*, 1641-1654.
- 299. Parikh, S.R., Campbell, H., Beebeejaun, K., Ribeiro, S., Gray, S.J., Borrow, R., Ramsay, M.E., and Ladhani, S.N. (2016). Meningococcal Group W Disease in Infants and Potential Prevention by Vaccination. Emerg Infect Dis *22*, 1505-1507.
- 300. Parker, R.E., Laut, C., Gaddy, J.A., Zadoks, R.N., Davies, H.D., and Manning, S.D. (2016). Association between genotypic diversity and biofilm production in group B Streptococcus. BMC microbiology *16*, 86.
- 301. Pattaro, C., Teumer, A., Gorski, M., Chu, A.Y., Li, M., Mijatovic, V., Garnaas, M., Tin, A., Sorice, R., Li, Y., et al. (2016). Genetic associations at 53 loci highlight cell types and biological pathways relevant for kidney function. Nat Commun 7, 10023.
- 302. Pattle, S.B., Utjesanovic, N., Togo, A., Wells, L., Conn, B., Monaghan, H., Junor, E., Johannessen, I., Cuschieri, K., and Talbot, S. (2017). Copy number gain of 11q13.3 genes associates with pathological stage in hypopharyngeal squamous cell carcinoma. Genes, chromosomes & cancer *56*, 185-198.
- 303. Paultre, D.S., Gustin, M.P., Molnar, A., and Oparka, K.J. (2016). Lost in transit: long-distance trafficking and phloem unloading of protein signals in Arabidopsis homografts. The Plant cell.
- 304. Pavitt, A.T., Pemberton, J.M., Kruuk, L.E., and Walling, C.A. (2016). Testosterone and cortisol concentrations vary with reproductive status in wild female red deer. Ecology and evolution *6*, 1163-1172.
- 305. Peachey, L.E., Pinchbeck, G.L., Matthews, J.B., Burden, F.A., Behnke, J.M., and Hodgkinson, J.E. (2016). Papaya latex supernatant has a potent effect on the free-living stages of equid cyathostomins in vitro. Vet Parasitol *228*, 23-29.
- 306. Pelly, V.S., Kannan, Y., Coomes, S.M., Entwistle, L.J., Ruckerl, D., Seddon, B., MacDonald, A.S., McKenzie, A., and Wilson, M.S. (2016). IL-4-producing ILC2s are required for the differentiation of T2 cells following Heligmosomoides polygyrus infection. Mucosal immunology *9*, 1407-1417.

- 307. Pemberton, J.M., Ellis, P.E., Pilkington, J.G., and Berenos, C. (2017). Inbreeding depression by environment interactions in a free-living mammal population. Heredity (Edinb) *118*, 64-77.
- 308. Peng, B., Klausen, C., Campbell, L., Leung, P.C., Horne, A.W., and Bedaiwy, M.A. (2016). Gonadotropin-releasing hormone and gonadotropin-releasing hormone receptor are expressed at tubal ectopic pregnancy implantation sites. Fertility and sterility *105*, 1620-1627.
- 309. Pengelly, R.J., Gheyas, A.A., Kuo, R., Mossotto, E., Seaby, E.G., Burt, D.W., Ennis, S., and Collins, A. (2016). Commercial chicken breeds exhibit highly divergent patterns of linkage disequilibrium. Heredity (Edinb) 117, 375-382.
- 310. Perez-Lopez, A.M., Soria-Gila, M.L., Marsden, E.R., Lilienkampf, A., and Bradley, M. (2016). Fluorogenic Substrates for In Situ Monitoring of Caspase-3 Activity in Live Cells. PLoS One *11*, e0153209.
- 311. Petrik, J. (2016). Immunomodulatory effects of exosomes produced by virus-infected cells. Transfusion and apheresis science: official journal of the World Apheresis Association: official journal of the European Society for Haemapheresis 55, 84-91.
- 312. Petrik, J., and Seghatchian, J. (2016). Big things from small packages: The multifaceted roles of extracellular vesicles in the components quality, therapy and infection. Transfusion and apheresis science: official journal of the World Apheresis Association: official journal of the European Society for Haemapheresis 55, 4-8.
- 313. Pittini, A., Casaravilla, C., Allen, J.E., and Diaz, A. (2016). Pharmacological inhibition of PI3K class III enhances the production of pro- and anti-inflammatory cytokines in dendritic cells stimulated by TLR agonists. International immunopharmacology *36*, 213-217.
- 314. Porphyre, T., Boden, L.A., Correia-Gomes, C., Auty, H.K., Gunn, G.J., and Woolhouse, M.E. (2016). Using national movement databases to help inform responses to swine disease outbreaks in Scotland: the impact of uncertainty around incursion time. Sci Rep *6*, 20258.
- 315. Prentice, J.C., Marion, G., Hutchings, M.R., McNeilly, T.N., and Matthews, L. (2017). Complex responses to movement-based disease control: when livestock trading helps. Journal of the Royal Society, Interface 14.
- 316. Pridans, C., Davis, G.M., Sauter, K.A., Lisowski, Z.M., Corripio-Miyar, Y., Raper, A., Lefevre, L., Young, R., McCulloch, M.E., Lillico, S., et al. (2016). A Csf1r-EGFP Transgene Provides a Novel Marker for Monocyte Subsets in Sheep. Journal of immunology (Baltimore, Md.: 1950) 197, 2297-2305.
- 317. Pritchard, T.C., Coffey, M.P., Bond, K.S., Hutchings, M.R., and Wall, E. (2017). Phenotypic effects of subclinical paratuberculosis (Johne's disease) in dairy cattle. J Dairy Sci *100*, 679-690.
- 318. Psifidi, A., Banos, G., Matika, O., Desta, T.T., Bettridge, J., Hume, D.A., Dessie, T., Christley, R., Wigley, P., Hanotte, O., et al. (2016). Genome-wide association studies of immune, disease and production traits in indigenous chicken ecotypes. Genetics, selection, evolution: GSE 48, 74.
- 319. Psifidi, A., Fife, M., Howell, J., Matika, O., van Diemen, P.M., Kuo, R., Smith, J., Hocking, P.M., Salmon, N., Jones, M.A., et al. (2016). The genomic architecture of resistance to Campylobacter jejuni intestinal colonisation in chickens. BMC Genomics *17*, 293.
- 320. Puenpa, J., Vongpunsawad, S., Osterback, R., Waris, M., Eriksson, E., Albert, J., Midgley, S., Fischer, T.K., Eis-Hubinger, A.M., Cabrerizo, M., et al. (2016). Molecular epidemiology and the evolution of human coxsackievirus A6. J Gen Virol *97*, 3225-3231.
- 321. Quick, J., Loman, N.J., Duraffour, S., Simpson, J.T., Severi, E., Cowley, L., Bore, J.A., Koundouno, R., Dudas, G., Mikhail, A., et al. (2016). Real-time, portable genome sequencing for Ebola surveillance. Nature *530*, 228-232.
- 322. Rambaut, A., Dudas, G., de Carvalho, L.M., Park, D.J., Yozwiak, N.L., Holmes, E.C., and Andersen, K.G. (2016). Comment on "Mutation rate and genotype variation of Ebola virus from Mali case sequences". Science *353*, 658.
- 323. Ramiro, R.S., Pollitt, L.C., Mideo, N., and Reece, S.E. (2016). Facilitation through altered resource availability in a mixed-species rodent malaria infection. Ecology letters *19*, 1041-1050.
- 324. Ratmann, O., Hodcroft, E.B., Pickles, M., Cori, A., Hall, M., Lycett, S., Colijn, C., Dearlove, B., Didelot, X., Frost, S., et al. (2017). Phylogenetic Tools for Generalized HIV-1 Epidemics: Findings from the PANGEA-HIV Methods Comparison. Mol Biol Evol *34*, 185-203.
- 325. Regan, C.E., Pilkington, J.G., Berenos, C., Pemberton, J.M., Smiseth, P.T., and Wilson, A.J. (2017). Accounting for female space sharing in St. Kilda Soay sheep (Ovis aries) results in little change in heritability estimates. Journal of evolutionary biology 30, 96-111.
- 326. Reid, W.D., Close, A.J., Humphrey, S., Chaloner, G., Lacharme-Lora, L., Rothwell, L., Kaiser, P., Williams, N.J., Humphrey, T.J., Wigley, P., et al. (2016). Cytokine responses in birds challenged with the human food-borne pathogen Campylobacter jejuni implies a Th17 response. R Soc Open Sci 3, 150541.

- 327. Ritchie, D.L., Barria, M.A., Peden, A.H., Yull, H.M., Kirkpatrick, J., Adlard, P., Ironside, J.W., and Head, M.W. (2016). UK latrogenic Creutzfeldt-Jakob disease: investigating human prion transmission across genotypic barriers using human tissue-based and molecular approaches. Acta neuropathologica 133, 579-595.
- 328. Rivero, L.A., Concepcion, J.L., Quintero-Troconis, E., Quinones, W., Michels, P.A., and Acosta, H. (2016). Trypanosoma evansi contains two auxiliary enzymes of glycolytic metabolism: Phosphoenolpyruvate carboxykinase and pyruvate phosphate dikinase. Exp Parasitol *165*, 7-15.
- Robb, C.T., Regan, K.H., Dorward, D.A., and Rossi, A.G. (2016). Key mechanisms governing resolution of lung inflammation. Seminars in immunopathology *38*, 425-448.
- 330. Robertson, A.L., Ogryzko, N.V., Henry, K.M., Loynes, C.A., Foulkes, M.J., Meloni, M.M., Wang, X., Ford, C., Jackson, M., Ingham, P.W., et al. (2016). Identification of benzopyrone as a common structural feature in compounds with anti-inflammatory activity in a zebrafish phenotypic screen. Disease models & mechanisms *9*, 621-632.
- 331. Robertson, K.A., and Ghazal, P. (2016). Interferon Control of the Sterol Metabolic Network: Bidirectional Molecular Circuitry-Mediating Host Protection. Front Immunol *7*, 634.
- 332. Robertson, K.A., Hsieh, W.Y., Forster, T., Blanc, M., Lu, H., Crick, P.J., Yutuc, E., Watterson, S., Martin, K., Griffiths, S.J., et al. (2016). An Interferon Regulated MicroRNA Provides Broad Cell-Intrinsic Antiviral Immunity through Multihit Host-Directed Targeting of the Sterol Pathway. PLoS Biol *14*, e1002364.
- 333. Robinson, G., Caldwell, G.S., Wade, M.J., Free, A., Jones, C.L., and Stead, S.M. (2016). Profiling bacterial communities associated with sediment-based aquaculture bioremediation systems under contrasting redox regimes. Sci Rep *6*, 38850.
- 334. Robinson, T.P., Bu, D.P., Carrique-Mas, J., Fevre, E.M., Gilbert, M., Grace, D., Hay, S.I., Jiwakanon, J., Kakkar, M., Kariuki, S., et al. (2016). Antibiotic resistance is the quintessential One Health issue. Transactions of the Royal Society of Tropical Medicine and Hygiene *110*, 377-380.
- 335. Robinson, T.P., Bu, D.P., Carrique-Mas, J., Fevre, E.M., Gilbert, M., Grace, D., Hay, S.I., Jiwakanon, J., Kakkar, M., Kariuki, S., et al. (2016). Antibiotic resistance: mitigation opportunities in livestock sector development. Animal *11*, 1-3.
- 336. Robledo, D., Taggart, J.B., Ireland, J.H., McAndrew, B.J., Starkey, W.G., Haley, C.S., Hamilton, A., Guy, D.R., Mota-Velasco, J.C., Gheyas, A.A., et al. (2016). Gene expression comparison of resistant and susceptible Atlantic salmon fry challenged with Infectious Pancreatic Necrosis virus reveals a marked contrast in immune response. BMC Genomics 17, 279.
- 337. Roehe, R., Dewhurst, R.J., Duthie, C.A., Rooke, J.A., McKain, N., Ross, D.W., Hyslop, J.J., Waterhouse, A., Freeman, T.C., Watson, M., et al. (2016). Bovine Host Genetic Variation Influences Rumen Microbial Methane Production with Best Selection Criterion for Low Methane Emitting and Efficiently Feed Converting Hosts Based on Metagenomic Gene Abundance. PLoS genetics 12, e1005846.
- 338. Rogers, A.J., Huang, Y.W., Heffron, C.L., Opriessnig, T., Patterson, A.R., and Meng, X.J. (2016). Prevalence of the Novel Torque Teno Sus Virus Species k2b from Pigs in the United States and Lack of Association with Post-Weaning Multisystemic Wasting Syndrome or Mulberry Heart Disease. Transbound Emerg Dis.
- 339. Rogers, P.A., Adamson, G.D., Al-Jefout, M., Becker, C.M., D'Hooghe, T.M., Dunselman, G.A., Fazleabas, A., Giudice, L.C., Horne, A.W., Hull, M.L., et al. (2016). Research Priorities for Endometriosis: Recommendations From a Global Consortium of Investigators in Endometriosis. Reproductive sciences (Thousand Oaks, Calif.).
- 340. Rosario, R., Filis, P., Tessyman, V., Kinnell, H., Childs, A.J., Gray, N.K., and Anderson, R.A. (2016). FMRP Associates with Cytoplasmic Granules at the Onset of Meiosis in the Human Oocyte. PLoS One *11*, e0163987.
- 341. Rudan, I., and Sridhar, D. (2016). Structure, function and five basic needs of the global health research system. J Glob Health *6*, 010505.
- Rudan, I., Yoshida, S., Chan, K.Y., Cousens, S., Sridhar, D., Bahl, R., and Martines, J. (2016). Setting health research priorities using the CHNRI method: I. Involving funders. J Glob Health *6*, 010301.
- 343. Rund, S.S., O'Donnell, A.J., Gentile, J.E., and Reece, S.E. (2016). Daily Rhythms in Mosquitoes and Their Consequences for Malaria Transmission. Insects 7.
- 344. Russell, C.D., Koch, O., Laurenson, I.F., O'Shea, D.T., Sutherland, R., and Mackintosh, C.L. (2015). Diagnosis and features of hospital-acquired pneumonia: a retrospective cohort study. The Journal of hospital infection.

- 345. Russell, C.D., Koch, O., Laurenson, I.F., O'Shea, D.T., Sutherland, R., and Mackintosh, C.L. (2016). Diagnosis and features of hospital-acquired pneumonia: a retrospective cohort study. The Journal of hospital infection *92*, 273-279.
- Rydevik, G., Innocent, G.T., Marion, G., Davidson, R.S., White, P.C., Billinis, C., Barrow, P., Mertens, P.P., Gavier-Widen, D., and Hutchings, M.R. (2016). Using Combined Diagnostic Test Results to Hindcast Trends of Infection from Cross-Sectional Data. PLoS computational biology *12*, e1004901.
- 347. Saccareau, M., Moreno, C.R., Kyriazakis, I., Faivre, R., and Bishop, S.C. (2016). Modelling gastrointestinal parasitism infection in a sheep flock over two reproductive seasons: in silico exploration and sensitivity analysis. Parasitology *143*, 1509-1531.
- 348. Sadeghi, M., Kapusinszky, B., Yugo, D.M., Phan, T.G., Deng, X., Kanevsky, I., Opriessnig, T., Woolums, A.R., Hurley, D.J., Meng, X.J., et al. (2017). Virome of US bovine calf serum. Biologicals: journal of the International Association of Biological Standardization.
- 349. Sahi, G., Hedley, P.E., Morris, J., Loake, G.J., and MacFarlane, S.A. (2016). Molecular and Biochemical Examination of Spraing Disease in Potato Tuber in Response to Tobacco rattle virus Infection. Molecular plant-microbe interactions: MPMI *29*, 822-828.
- 350. Sanderson, P.A., Critchley, H.O., Williams, A.R., Arends, M.J., and Saunders, P.T. (2016). New concepts for an old problem: the diagnosis of endometrial hyperplasia. Human reproduction update.
- 351. Sanghera, S., Barton, P., Bhattacharya, S., Horne, A.W., and Roberts, T.E. (2016). Pharmaceutical treatments to prevent recurrence of endometriosis following surgery: a model-based economic evaluation. BMJ open *6*, e010580.
- 352. Saraswat, L., Ayansina, D.T., Cooper, K.G., Bhattacharya, S., Miligkos, D., Horne, A.W., and Bhattacharya, S. (2016). Pregnancy outcomes in women with endometriosis: a national record linkage study. BJOG: an international journal of obstetrics and gynaecology *124*, 444-452.
- 353. Sargison, N., Francis, E., Davison, C., Bronsvoort, B.M., Handel, I., and Mazeri, S. (2016). Observations on the biology, epidemiology and economic relevance of rumen flukes (Paramphistomidae) in cattle kept in a temperate environment. Vet Parasitol *219*, 7-16.
- 354. Sauter, K.A., Waddell, L.A., Lisowski, Z.M., Young, R., Lefevre, L., Davis, G.M., Clohisey, S.M., McCulloch, M., Magowan, E., Mabbott, N.A., et al. (2016). Macrophage colony-stimulating factor (CSF1) controls monocyte production and maturation and the steady-state size of the liver in pigs. American journal of physiology. Gastrointestinal and liver physiology *311*, G533-547.
- 355. Schwarz-Linek, J., Arlt, J., Jepson, A., Dawson, A., Vissers, T., Miroli, D., Pilizota, T., Martinez, V.A., and Poon, W.C. (2016). Escherichia coli as a model active colloid: A practical introduction. Colloids and surfaces. B, Biointerfaces 137, 2-16.
- 356. Schwarze, J., Fitch, P.M., Heimweg, J., Errington, C., Matsuda, R., de Bruin, H.G., van den Berge, M., van Oosterhout, A.J., and Heijink, I.H. (2016). Viral mimic poly-(I:C) attenuates airway epithelial T-cell suppressive capacity: implications for asthma. Eur Respir J 48, 1785-1788.
- 357. Seeker, L.A., Holland, R., Underwood, S., Fairlie, J., Psifidi, A., Ilska, J.J., Bagnall, A., Whitelaw, B., Coffey, M., Banos, G., et al. (2016). Method Specific Calibration Corrects for DNA Extraction Method Effects on Relative Telomere Length Measurements by Quantitative PCR. PLoS One *11*, e0164046.
- 358. Sehgal, A., Kobayashi, A., Donaldson, D.S., and Mabbott, N.A. (2017). c-Rel is dispensable for the differentiation and functional maturation of M cells in the follicle-associated epithelium. Immunobiology *222*, 316-326.
- 359. Seibyl, J., Catafau, A.M., Barthel, H., Ishii, K., Rowe, C.C., Leverenz, J.B., Ghetti, B., Ironside, J.W., Takao, M., Akatsu, H., et al. (2016). Impact of training method on the robustness of the visual assessment of 18F-florbetaben PET scans: results from a Phase 3 trial. Journal of nuclear medicine: official publication, Society of Nuclear Medicine *57*, 900-906.
- 360. Selega, A., Sirocchi, C., Iosub, I., Granneman, S., and Sanguinetti, G. (2017). Robust statistical modeling improves sensitivity of high-throughput RNA structure probing experiments. Nature methods *14*, 83-89
- 361. Serao, N.V., Kemp, R.A., Mote, B.E., Willson, P., Harding, J.C., Bishop, S.C., Plastow, G.S., and Dekkers, J.C. (2016). Genetic and genomic basis of antibody response to porcine reproductive and respiratory syndrome (PRRS) in gilts and sows. Genetics, selection, evolution: GSE 48, 51.
- 362. Seth, S., Akram, A.R., McCool, P., Westerfeld, J., Wilson, D., McLaughlin, S., Dhaliwal, K., and Williams, C.K. (2016). Assessing the utility of autofluorescence-based pulmonary optical endomicroscopy to predict the malignant potential of solitary pulmonary nodules in humans. Sci Rep *6*, 31372.

- 363. Sharma, A., Hussain, A., Mun, B.G., Imran, Q.M., Falak, N., Lee, S.U., Kim, J.Y., Hong, J.K., Loake, G.J., Ali, A., et al. (2016). Comprehensive analysis of plant rapid alkalization factor (RALF) genes. Plant physiology and biochemistry: PPB / Societe française de physiologie vegetale *106*, 82-90.
- 364. Sharma, J.R., Khatri, R., and Harper, I. (2016). Understanding Health Research Ethics in Nepal. Developing world bioethics *16*, 140-147.
- 365. Sharma, V., Kaur, T., Bridle, H., and Ghosh, M. (2017). Antimicrobial efficacy and safety of mucoadhesive exopolymer produced by Acinetobacter haemolyticus. International journal of biological macromolecules *94*, 187-193.
- 366. Sharp, G.C., Hutchinson, J.L., Hibbert, N., Freeman, T.C., Saunders, P.T., and Norman, J.E. (2016). Transcription Analysis of the Myometrium of Labouring and Non-Labouring Women. PLoS One *11*, e0155413.
- 367. Shepherd, L., Borges, A., Ledergerber, B., Domingo, P., Castagna, A., Rockstroh, J., Knysz, B., Tomazic, J., Karpov, I., Kirk, O., et al. (2016). Infection-related and -unrelated malignancies, HIV and the aging population. HIV Med *17*, 590-600.
- 368. Simbari, F., McCaskill, J., Coakley, G., Millar, M., Maizels, R.M., Fabrias, G., Casas, J., and Buck, A.H. (2016). Plasmalogen enrichment in exosomes secreted by a nematode parasite versus those derived from its mouse host: implications for exosome stability and biology. Journal of extracellular vesicles *5*, 30741.
- 369. Sinclair, R., Melville, L., Sargison, F., Kenyon, F., Nussey, D., Watt, K., and Sargison, N. (2016). Gastrointestinal nematode species diversity in Soay sheep kept in a natural environment without active parasite control. Vet Parasitol *227*, 1-7.
- 370. Smith, D.B., Gaunt, E.R., Digard, P., Templeton, K., and Simmonds, P. (2016). Detection of influenza C virus but not influenza D virus in Scottish respiratory samples. Journal of clinical virology: the official publication of the Pan American Society for Clinical Virology *74*, 50-53.
- 371. Smith, D.B., Paddy, J.O., and Simmonds, P. (2016). The use of human sewage screening for community surveillance of hepatitis E virus in the UK. Journal of medical virology *88*, 915-918.
- 372. Smith, D.B., Simmonds, P., Izopet, J., Oliveira-Filho, E.F., Ulrich, R.G., Johne, R., Koenig, M., Jameel, S., Harrison, T.J., Meng, X.J., et al. (2016). Proposed reference sequences for Hepatitis E virus subtypes. J Gen Virol *97*, 537-542.
- 373. Smith, J., Gheyas, A., and Burt, D.W. (2016). Animal genomics and infectious disease resistance in poultry. Rev Sci Tech *35*, 105-119.
- 374. Smith, K.A., Filbey, K.J., Reynolds, L.A., Hewitson, J.P., Harcus, Y., Boon, L., Sparwasser, T., Hammerling, G., and Maizels, R.M. (2016). Low-level regulatory T-cell activity is essential for functional type-2 effector immunity to expel gastrointestinal helminths. Mucosal immunology *9*, 428-443.
- 375. Smits, H.H., Hiemstra, P.S., Prazeres da Costa, C., Ege, M., Edwards, M., Garn, H., Howarth, P.H., Jartti, T., de Jong, E.C., Maizels, R.M., et al. (2016). Microbes and asthma: Opportunities for intervention. The Journal of allergy and clinical immunology *137*, 690-697.
- 376. Sood, S., Brownlie, R.J., Garcia, C., Cowan, G., Salmond, R.J., Sakaguchi, S., and Zamoyska, R. (2016). Loss of the Protein Tyrosine Phosphatase PTPN22 Reduces Mannan-Induced Autoimmune Arthritis in SKG Mice. Journal of immunology (Baltimore, Md.: 1950) 197, 429-440.
- 377. Soukarieh, F., Nowicki, M.W., Bastide, A., Poyry, T., Jones, C., Dudek, K., Patwardhan, G., Meullenet, F., Oldham, N.J., Walkinshaw, M.D., et al. (2016). Design of nucleotide-mimetic and non-nucleotide inhibitors of the translation initiation factor eIF4E: Synthesis, structural and functional characterisation. European journal of medicinal chemistry *124*, 200-217.
- 378. Sridhar, D. (2016). Making the SDGs useful: a Herculean task. Lancet 388, 1453-1454.
- 379. Sridhar, D., Kickbusch, I., Moon, S., Dzau, V., Heymann, D., Jha, A.K., Saavendra, J., Stocking, B., Woskie, L., and Piot, P. (2016). Facing forward after Ebola: questions for the next director general of the World Health Organization. Bmj *353*, i2666.
- 380. Stacy, A., McNally, L., Darch, S.E., Brown, S.P., and Whiteley, M. (2016). The biogeography of polymicrobial infection. Nature reviews. Microbiology *14*, 93-105.
- 381. Steer, B., Strehle, M., Sattler, C., Bund, D., Flach, B., Stoeger, T., Haas, J.G., and Adler, H. (2016). The small noncoding RNAs (sncRNAs) of murine gammaherpesvirus 68 (MHV-68) are involved in regulating the latent-to-lytic switch in vivo. Sci Rep *6*, 32128.
- 382. Stephen, J., Bravo, E.L., Colligan, D., Fraser, A.R., Petrik, J., and Campbell, J.D. (2016). Mesenchymal stromal cells as multifunctional cellular therapeutics a potential role for extracellular vesicles. Transfusion and apheresis science: official journal of the World Apheresis Association: official journal of the European Society for Haemapheresis 55, 62-69.

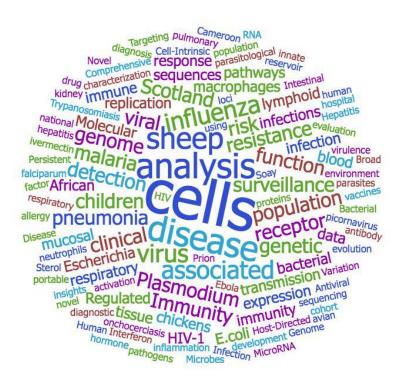
- 383. Stevens, M.P. (2016). Modulation of the Interaction of Enteric Bacteria with Intestinal Mucosa by Stress-Related Catecholamines. Advances in experimental medicine and biology *874*, 143-166.
- 384. Stevenson, K., McVey, A.F., Clark, I.B., Swain, P.S., and Pilizota, T. (2016). General calibration of microbial growth in microplate readers. Sci Rep *6*, 38828.
- 385. Subiros-Funosas, R., Mendive-Tapia, L., Sot, J., Pound, J.D., Barth, N., Varela, Y., Goni, F.M., Paterson, M., Gregory, C.D., Albericio, F., et al. (2017). A Trp-BODIPY cyclic peptide for fluorescence labelling of apoptotic bodies. Chem Commun (Camb) *53*, 945-948.
- 386. Subramaniam, S., Cao, D., Tian, D., Cao, Q.M., Overend, C., Yugo, D.M., Matzinger, S.R., Rogers, A.J., Heffron, C.L., Catanzaro, N., et al. (2017). Efficient priming of CD4 T cells by Langerin-expressing dendritic cells targeted with porcine epidemic diarrhea virus spike protein domains in pigs. Virus research 227, 212-219.
- 387. Sundararaman, S.A., Plenderleith, L.J., Liu, W., Loy, D.E., Learn, G.H., Li, Y., Shaw, K.S., Ayouba, A., Peeters, M., Speede, S., et al. (2016). Genomes of cryptic chimpanzee Plasmodium species reveal key evolutionary events leading to human malaria. Nat Commun *7*, 11078.
- 388. Surve, S.V., Jensen, B.C., Heestand, M., Mazet, M., Smith, T.K., Bringaud, F., Parsons, M., and Schnaufer, A. (2017). NADH dehydrogenase of Trypanosoma brucei is important for efficient acetate production in bloodstream forms. Mol Biochem Parasitol *211*, 57-61.
- 389. Swain, P.S., Stevenson, K., Leary, A., Montano-Gutierrez, L.F., Clark, I.B., Vogel, J., and Pilizota, T. (2016). Inferring time derivatives including cell growth rates using Gaussian processes. Nat Commun 7, 13766.
- 390. Syed, S.N., Ducrotoy, M.J., and Bachmann, T.T. (2016). Antimicrobial resistance diagnostics: time to call in the young? The Lancet. Infectious diseases *16*, 519-521.
- 391. Szitenberg, A., Cha, S., Opperman, C.H., Bird, D.M., Blaxter, M.L., and Lunt, D.H. (2016). Genetic drift, not life history or RNAi, determine long term evolution of transposable elements. Genome biology and evolution *8*, 2964-2978.
- 392. Taylor, M., Sargison, N., Howe, M., and Scott, P. (2016). Dosing ewes at lambing time. Vet Rec *178*, 377-378.
- 393. Thackeray, S.J., Henrys, P.A., Hemming, D., Bell, J.R., Botham, M.S., Burthe, S., Helaouet, P., Johns, D.G., Jones, I.D., Leech, D.I., et al. (2016). Phenological sensitivity to climate across taxa and trophic levels. Nature *535*, 241-245.
- 394. Thamsborg, S.M., Ketzis, J., Horii, Y., and Matthews, J.B. (2016). Strongyloides spp. infections of veterinary importance. Parasitology, 1-11.
- 395. Thomson, E., Ip, C.L., Badhan, A., Christiansen, M.T., Adamson, W., Ansari, M.A., Bibby, D., Breuer, J., Brown, A., Bowden, R., et al. (2016). Comparison of next generation sequencing technologies for the comprehensive assessment of full-length hepatitis C viral genomes. Journal of clinical microbiology *54*, 2470-2484.
- 396. Thurlow, S.E., Kilgour, D.P., Campopiano, D.J., Mackay, C.L., Langridge-Smith, P.R., Clarke, D.J., and Campbell, C.J. (2016). Determination of Protein Thiol Reduction Potential by Isotope Labeling and Intact Mass Measurement. Anal Chem 88, 2727-2733.
- 397. Trainor-Moss, S., and Mutapi, F. (2016). Schistosomiasis therapeutics: whats in the pipeline? Expert review of clinical pharmacology *9*, 157-160.
- 398. Trikin, R., Doiron, N., Hoffmann, A., Haenni, B., Jakob, M., Schnaufer, A., Schimanski, B., Zuber, B., and Ochsenreiter, T. (2016). TAC102 Is a Novel Component of the Mitochondrial Genome Segregation Machinery in Trypanosomes. PLoS Pathog *12*, e1005586.
- 399. Trubitsyn, D., Abreu, F., Ward, F.B., Taylor, T., Hattori, M., Kondo, S., Trivedi, U., Staniland, S., Lins, U., and Bazylinski, D.A. (2016). Draft Genome Sequence of Magnetovibrio blakemorei Strain MV-1, a Marine Vibrioid Magnetotactic Bacterium. Genome Announc 4.
- 400. Tsai, H.Y., Hamilton, A., Tinch, A.E., Guy, D.R., Bron, J.E., Taggart, J.B., Gharbi, K., Stear, M., Matika, O., Pong-Wong, R., et al. (2016). Genomic prediction of host resistance to sea lice in farmed Atlantic salmon populations. Genetics, selection, evolution: GSE 48, 47.
- 401. Tsairidou, S., Brotherstone, S., Coffey, M., Bishop, S.C., and Woolliams, J.A. (2016). Quantitative genetic analysis of the bTB diagnostic single intradermal comparative cervical test (SICCT). Genetics, selection, evolution: GSE 48, 90.
- 402. Tuck, L.R., Altenbach, K., Ang, T.F., Crawshaw, A.D., Campopiano, D.J., Clarke, D.J., and Marles-Wright, J. (2016). Insight into Coenzyme A cofactor binding and the mechanism of acyl-transfer in an acylating aldehyde dehydrogenase from Clostridium phytofermentans. Sci Rep *6*, 22108.

- 403. Turnbull, M.L., Wise, H.M., Nicol, M.Q., Smith, N., Dunfee, R.L., Beard, P.M., Jagger, B.W., Ligertwood, Y., Hardisty, G.R., Xiao, H., et al. (2016). The Role of the B-Allele of the Influenza A Virus Segment 8 in Setting Mammalian Host Range and Pathogenicity. J Virol 90, 9263-9284.
- 404. Turner, D.G., Leech, M.D., O'Connor, R.A., and Anderton, S.M. (2016). Methods for Ex Vivo Analysis of Immune Cell Function from the Central Nervous System. Methods Mol Biol *1304*, 81-90.
- 405. Tzelos, T., Matthews, J.B., Buck, A.H., Simbari, F., Frew, D., Inglis, N.F., McLean, K., Nisbet, A.J., Whitelaw, C.B., Knox, D.P., et al. (2016). A preliminary proteomic characterisation of extracellular vesicles released by the ovine parasitic nematode, Teladorsagia circumcincta. Vet Parasitol *221*, 84-92.
- 406. Ugrina, I., Campbell, H., and Vuckovic, F. (2017). Laboratory Experimental Design for a Glycomic Study. Methods Mol Biol *1503*, 13-19.
- 407. Ulyanchenko, S., O'Neill, K.E., Medley, T., Farley, A.M., Vaidya, H.J., Cook, A.M., Blair, N.F., and Blackburn, C.C. (2016). Identification of a Bipotent Epithelial Progenitor Population in the Adult Thymus. Cell reports *14*, 2819-2832.
- 408. Urwin, P.J., Mackenzie, J.M., Llewelyn, C.A., Will, R.G., and Hewitt, P.E. (2016). Creutzfeldt-Jakob disease and blood transfusion: updated results of the UK Transfusion Medicine Epidemiology Review Study. Vox Sang *110*, 310-316.
- 409. Vaidya, H.J., Briones Leon, A., and Blackburn, C.C. (2016). FOXN1 in thymus organogenesis and development. European journal of immunology *46*, 1826-1837.
- 410. Vale, P.F., and Jardine, M.D. (2016). Infection avoidance behaviour: viral exposure reduces the motivation to forage in female Drosophila melanogaster. Fly, 0.
- 411. Vale, P.F., McNally, L., Doeschl-Wilson, A., King, K.C., Popat, R., Domingo-Sananes, M.R., Allen, J.E., Soares, M.P., and Kummerli, R. (2016). Beyond killing: can we find new ways to manage infection? Evolution, medicine, and public health *2016*, 148-157.
- 412. Van Dung, N., Anh, P.H., Van Cuong, N., Hoa, N.T., Carrique-Mas, J., Hien, V.B., Sharp, C., Rabaa, M., Berto, A., Campbell, J., et al. (2016). Large-scale screening and characterization of enteroviruses and kobuviruses infecting pigs in Vietnam. J Gen Virol *97*, 378-388.
- 413. van Leeuwen, E.M., Sabo, A., Bis, J.C., Huffman, J.E., Manichaikul, A., Smith, A.V., Feitosa, M.F., Demissie, S., Joshi, P.K., Duan, Q., et al. (2016). Meta-analysis of 49 549 individuals imputed with the 1000 Genomes Project reveals an exonic damaging variant in ANGPTL4 determining fasting TG levels. Journal of medical genetics *53*, 441-449.
- 414. Velu, P.P., Fernandes, S.E., Laurenson, I.F., and Noble, D.D. (2016). Pulmonary Mycobacterium marinum infection: 'fish tank granuloma' of the lung. Scottish medical journal *61*, 203-206.
- 415. Venkateswaran, S., Gwynne, P.J., Wu, M., Hardman, A., Lilienkampf, A., Pernagallo, S., Blakely, G., Swann, D.G., Bradley, M., and Gallagher, M.P. (2016). High-throughput Identification of Bacteria Repellent Polymers for Medical Devices. Journal of visualized experiments: JoVE.
- Venkateswaran, S., Henrique Dos Santos, O.D., Scholefield, E., Lilienkampf, A., Gwynne, P.J., Swann, D.G., Dhaliwal, K., Gallagher, M.P., and Bradley, M. (2016). Fortified interpenetrating polymers bacteria resistant coatings for medical devices. Journal of materials chemistry. B, Materials for biology and medicine 4, 5405-5411.
- 417. Venkateswaran, S., Luque-Gonzalez, M.A., Tabraue-Chavez, M., Fara, M.A., Lopez-Longarela, B., Cano-Cortes, V., Lopez-Delgado, F.J., Sanchez-Martin, R.M., Ilyine, H., Bradley, M., et al. (2016). Novel bead-based platform for direct detection of unlabelled nucleic acids through Single Nucleobase Labelling. Talanta *161*, 489-496.
- 418. Venturato, A., MacFarlane, G., Geng, J., and Bradley, M. (2016). Understanding Polymer-Cell Attachment. Macromolecular bioscience *16*, 1864-1872.
- 419. Vermeren, S., Miles, K., Chu, J.Y., Salter, D., Zamoyska, R., and Gray, M. (2016). PTPN22 Is a Critical Regulator of Fcgamma Receptor-Mediated Neutrophil Activation. Journal of immunology (Baltimore, Md.: 1950) 197, 4771-4779.
- 420. Vincenti, J.E., Murphy, L., Grabert, K., McColl, B.W., Cancellotti, E., Freeman, T.C., and Manson, J.C. (2015). Defining the microglia response during the time course of chronic neurodegeneration. J Virol.
- 421. Vincenti, J.E., Murphy, L., Grabert, K., McColl, B.W., Cancellotti, E., Freeman, T.C., and Manson, J.C. (2016). Defining the Microglia Response during the Time Course of Chronic Neurodegeneration. J Virol 90, 3003-3017.
- 422. Vitak, N., Hume, D.A., Chappell, K.J., Sester, D.P., and Stacey, K.J. (2016). Induction of interferon and cell death in response to cytosolic DNA in chicken macrophages. Dev Comp Immunol *59*, 145-152.

- 423. Voss, J.J., Ford, C.A., Petrova, S., Melville, L., Paterson, M., Pound, J.D., Holland, P., Giotti, B., Freeman, T.C., and Gregory, C.D. (2017). Modulation of macrophage antitumor potential by apoptotic lymphoma cells. Cell death and differentiation.
- 424. Vuckovic, F., Theodoratou, E., Thaci, K., Timofeeva, M., Vojta, A., Stambuk, J., Pucic-Bakovic, M., Derek, L., Servis, D., Rudd, P., et al. (2016). IgG glycome in colorectal cancer. Clinical cancer research: an official journal of the American Association for Cancer Research *22*, 3078-3086.
- Walker, D., Abbondati, E., Cox, A.L., Mitchell, G.B., Pizzi, R., Sharp, C.P., and Philbey, A.W. (2016). Infectious canine hepatitis in red foxes (Vulpes vulpes) in wildlife rescue centres in the UK. Vet Rec *178*, 421.
- 426. Walker, D., Fee, S.A., Hartley, G., Learmount, J., O'Hagan, M.J., Meredith, A.L., de, C.B.B.M., Porphyre, T., Sharp, C.P., and Philbey, A.W. (2016). Serological and molecular epidemiology of canine adenovirus type 1 in red foxes (Vulpes vulpes) in the United Kingdom. Sci Rep *6*, 36051.
- 427. Wami, W.M., Nausch, N., Midzi, N., Gwisai, R., Mduluza, T., Woolhouse, M.E., and Mutapi, F. (2016). Comparative Assessment of Health Benefits of Praziquantel Treatment of Urogenital Schistosomiasis in Preschool and Primary School-Aged Children. BioMed research international *2016*, 9162631.
- 428. Wang, B., McHugh, B.J., Qureshi, A., Campopiano, D.J., Clarke, D.J., Fitzgerald, J.R., Dorin, J.R., Weller, R., and Davidson, D.J. (2017). IL-1beta-Induced Protection of Keratinocytes against Staphylococcus aureus-Secreted Proteases Is Mediated by Human beta-Defensin 2. J Invest Dermatol *137*, 95-105.
- 429. Wang, G., Duan, Z., Sheng, Y., Neumann, K., Deng, L., Li, J., Bradley, M., and Zhang, R. (2016). Tuning the emission properties of a fluorescent polymer using a polymer microarray approach identification of an optothermo responsive polymer. Chem Commun (Camb) *52*, 10521-10524.
- 430. Wang, Y., Klaric, L., Yu, X., Thaqi, K., Dong, J., Novokmet, M., Wilson, J., Polasek, O., Liu, Y., Kristic, J., et al. (2016). The Association Between Glycosylation of Immunoglobulin G and Hypertension: A Multiple Ethnic Cross-Sectional Study. Medicine *95*, e3379.
- 431. Ward, M.J., Goncheva, M., Richardson, E., McAdam, P.R., Raftis, E., Kearns, A., Daum, R.S., David, M.Z., Lauderdale, T.L., Edwards, G.F., et al. (2016). Identification of source and sink populations for the emergence and global spread of the East-Asia clone of community-associated MRSA. Genome Biol *17*, 160.
- 432. Wardrop, N.A., Thomas, L.F., Cook, E.A., de Glanville, W.A., Atkinson, P.M., Wamae, C.N., and Fevre, E.M. (2016). The Sero-epidemiology of Coxiella burnetii in Humans and Cattle, Western Kenya: Evidence from a Cross-Sectional Study. PLoS Negl Trop Dis *10*, e0005032.
- 433. Warren, W.C., Hillier, L.W., Tomlinson, C., Minx, P., Kremitzki, M., Graves, T., Markovic, C., Bouk, N., Pruitt, K.D., Thibaud-Nissen, F., et al. (2017). A New Chicken Genome Assembly Provides Insight into Avian Genome Structure. G3 (Bethesda, Md.) 7, 109-117.
- 434. Waters, S.A., McAteer, S.P., Kudla, G., Pang, I., Deshpande, N.P., Amos, T.G., Leong, K.W., Wilkins, M.R., Strugnell, R., Gally, D.L., et al. (2016). Small RNA interactome of pathogenic E. coli revealed through crosslinking of RNase E. The EMBO journal *36*, 374-387.
- 435. Watson, R.L., Bird, E.J., Underwood, S., Wilbourn, R.V., Fairlie, J., Watt, K., Salvo-Chirnside, E., Pilkington, J.G., Pemberton, J.M., McNeilly, T.N., et al. (2016). Sex differences in leucocyte telomere length in a free-living mammal. Mol Ecol.
- 436. Watson, R.L., McNeilly, T.N., Watt, K.A., Pemberton, J.M., Pilkington, J.G., Waterfall, M., Hopper, P.R., Cooney, D., Zamoyska, R., and Nussey, D.H. (2016). Cellular and humoral immunity in a wild mammal: Variation with age & sex and association with overwinter survival. Ecology and evolution *6*, 8695-8705.
- 437. Watt, K.A., Nussey, D.H., Maclellan, R., Pilkington, J.G., and McNeilly, T.N. (2016). Fecal antibody levels as a noninvasive method for measuring immunity to gastrointestinal nematodes in ecological studies. Ecology and evolution *6*, 56-67.
- 438. Webster, C.L., Longdon, B., Lewis, S.H., and Obbard, D.J. (2016). Twenty-Five New Viruses Associated with the Drosophilidae (Diptera). Evolutionary bioinformatics online *12*, 13-25.
- Weiss, E., Zgaga, L., Read, S., Wild, S., Dunlop, M.G., Campbell, H., McQuillan, R., and Wilson, J.F. (2016). Farming, Foreign Holidays, and Vitamin D in Orkney. PLoS One *11*, e0155633.
- 440. Welburn, S.C., Bardosh, K.L., and Coleman, P.G. (2016). Novel Financing Model for Neglected Tropical Diseases: Development Impact Bonds Applied to Sleeping Sickness and Rabies Control. PLoS Negl Trop Dis 10, e0005000.
- 441. Welburn, S.C., Molyneux, D.H., and Maudlin, I. (2015). Beyond Tsetse Implications for Research and Control of Human African Trypanosomiasis Epidemics. Trends Parasitol.
- Welburn, S.C., Molyneux, D.H., and Maudlin, I. (2016). Beyond Tsetse--Implications for Research and Control of Human African Trypanosomiasis Epidemics. Trends Parasitol *32*, 230-241.

- 443. Wells, B., Thomson, S., Ensor, H., Innes, E.A., and Katzer, F. (2016). Development of a sensitive method to extract and detect low numbers of Cryptosporidium oocysts from adult cattle faecal samples. Vet Parasitol *227*, 26-29.
- 444. Whitaker, L.H., Reid, J., Choa, A., McFee, S., Seretny, M., Wilson, J., Elton, R.A., Vincent, K., and Horne, A.W. (2016). An Exploratory Study into Objective and Reported Characteristics of Neuropathic Pain in Women with Chronic Pelvic Pain. PLoS One *11*, e0151950.
- Whitehouse, I.J., Brown, D., Baybutt, H., Diack, A.B., Kellett, K.A., Piccardo, P., Manson, J.C., and Hooper, N.M. (2016). Ablation of Prion Protein in Wild Type Human Amyloid Precursor Protein (APP) Transgenic Mice Does Not Alter The Proteolysis of APP, Levels of Amyloid-beta or Pathologic Phenotype. PLoS One 11, e0159119.
- Widder, S., Allen, R.J., Pfeiffer, T., Curtis, T.P., Wiuf, C., Sloan, W.T., Cordero, O.X., Brown, S.P., Momeni,
   B., Shou, W., et al. (2016). Challenges in microbial ecology: building predictive understanding of community function and dynamics. The ISME journal 1-, 2557-2568.
- 447. Wilkie, H., Gossner, A., Bishop, S., and Hopkins, J. (2016). Correction: Variations in T cell transcription factor sequence and expression associated with resistance to the sheep nematode Teladorsagia circumcincta. PLoS One *11*, e0152396.
- 448. Wilkie, H., Gossner, A., Bishop, S., and Hopkins, J. (2016). Variations in T Cell Transcription Factor Sequence and Expression Associated with Resistance to the Sheep Nematode Teladorsagia circumcincta. PLoS One *11*, e0149644.
- 449. Wilkie, H., Nicol, L., Gossner, A., and Hopkins, J. (2016). Mucosal Expression of T Cell Gene Variants Is Associated with Differential Resistance to Teladorsagia circumcincta. PLoS One *11*, e0168194.
- 450. Will, R.G., and Ironside, J.W. (2017). Sporadic and Infectious Human Prion Diseases. Cold Spring Harbor perspectives in medicine 7.
- 451. Williams, J.L., Hall, S.J., Del Corvo, M., Ballingall, K.T., Colli, L., Ajmone Marsan, P., and Biscarini, F. (2016). Inbreeding and purging at the genomic Level: the Chillingham cattle reveal extensive, non-random SNP heterozygosity. Anim Genet *47*, 19-27.
- 452. Williams, T., Nair, H., Simpson, J., and Embleton, N. (2016). Use of Donor Human Milk and Maternal Breastfeeding Rates: A Systematic Review. Journal of human lactation: official journal of International Lactation Consultant Association *32*, 212-220.
- 453. Wilson, R., Aksamit, T., Aliberti, S., De Soyza, A., Elborn, J.S., Goeminne, P., Hill, A.T., Menendez, R., and Polverino, E. (2016). Challenges in managing Pseudomonas aeruginosa in non-cystic fibrosis bronchiectasis. Respiratory medicine *117*, 179-189.
- 454. Witteveldt, J., Martin-Gans, M., and Simmonds, P. (2016). Enhancement of the replication of HCV replicons of genotypes 1-4 by manipulation of CpG and UpA dinucleotide frequencies and use of cell lines expressing SECL14L2 application for antiviral resistance testing. Antimicrobial agents and chemotherapy.
- 455. Woolhouse, M., Waugh, C., Perry, M.R., and Nair, H. (2016). Global disease burden due to antibiotic resistance state of the evidence. J Glob Health *6*, 010306.
- Woolhouse, M.E., Brierley, L., McCaffery, C., and Lycett, S. (2016). Assessing the Epidemic Potential of RNA and DNA Viruses. Emerg Infect Dis *22*, 2037-2044.
- 457. Wright, H.W., Bartley, K., Huntley, J.F., and Nisbet, A.J. (2016). Characterisation of tropomyosin and paramyosin as vaccine candidate molecules for the poultry red mite, Dermanyssus gallinae. Parasit Vectors *9*, 544.
- 458. Wu, Z., Hu, T., Rothwell, L., Vervelde, L., Hume, D.A., and Kaiser, P. (2016). Analysis of the function of IL-10 in chickens using specific neutralising antibodies and a sensitive capture ELISA. Dev Comp Immunol *63*, 206-212.
- 459. Wylie, C.E., Shaw, D.J., Verheyen, K.L., and Newton, J.R. (2016). Decision-tree analysis of clinical data to aid diagnostic reasoning for equine laminitis: a cross-sectional study. Vet Rec *178*, 420.
- 460. Xiao, C.T., Harmon, K.M., Halbur, P.G., and Opriessnig, T. (2016). PCV2d-2 is the predominant type of PCV2 DNA in pig samples collected in the U.S. during 2014-2016. Vet Microbiol *197*, 72-77.
- 461. Yebra, G., Kalish, M.L., and Leigh Brown, A.J. (2016). Reconstructing the HIV-1 CRF02\_AG and CRF06 cpx epidemics in Burkina Faso and West Africa using early samples. Infect Genet Evol 46.
- 462. Young, V.J., Ahmad, S.F., Brown, J.K., Duncan, W.C., and Horne, A.W. (2016). ID2 mediates the transforming growth factor-beta1-induced Warburg-like effect seen in the peritoneum of women with endometriosis. Mol Hum Reprod *22*, 648-654.

- 463. Yu, X., Wang, Y., Kristic, J., Dong, J., Chu, X., Ge, S., Wang, H., Fang, H., Gao, Q., Liu, D., et al. (2016). Profiling IgG N-glycans as potential biomarker of chronological and biological ages: A community-based study in a Han Chinese population. Medicine *95*, e4112.
- 464. Yun, B.W., Skelly, M.J., Yin, M., Yu, M., Mun, B.G., Lee, S.U., Hussain, A., Spoel, S.H., and Loake, G.J. (2016). Nitric oxide and S-nitrosoglutathione function additively during plant immunity. The New phytologist *211*, 516-526.
- 465. Zahoor, I., Mitchell, M.A., Hall, S., Beard, P.M., Gous, R.M., De Koning, D.J., and Hocking, P.M. (2016). Predicted optimum ambient temperatures for broiler chickens to dissipate metabolic heat do not affect performance or improve breast muscle quality. British poultry science *57*, 134-141.
- 466. Zhang, S., Sammon, P.M., King, I., Andrade, A.L., Toscano, C.M., Araujo, S.N., Sinha, A., Madhi, S.A., Khandaker, G., Yin, J.K., et al. (2016). Cost of management of severe pneumonia in young children: systematic analysis. J Glob Health *6*, 010408.
- 467. Zhao, C., Fernandez, A., Avlonitis, N., Vande Velde, G., Bradley, M., Read, N.D., and Vendrell, M. (2016). Searching for the Optimal Fluorophore to Label Antimicrobial Peptides. ACS combinatorial science *18*, 689-696.
- 468. Zhao, Y., He, M., Ding, J., Xi, Q., Loake, G.J., and Zheng, W. (2016). Regulation of Anticancer Styrylpyrone Biosynthesis in the Medicinal Mushroom Inonotus obliquus Requires Thioredoxin Mediated Transnitrosylation of S-nitrosoglutathione Reductase. Sci Rep *6*, 37601.
- 469. Zheng, X., Liu, G., Opriessnig, T., Wang, Z., Yang, Z., and Jiang, Y. (2016). Rapid detection and grouping of porcine bocaviruses by an EvaGreen based multiplex real-time PCR assay using melting curve analysis. Molecular and cellular probes *30*, 195-204.
- 470. Zimmermann, S., Hall, L., Riley, S., Sorensen, J., Amaro, R.E., and Schnaufer, A. (2016). A novel high-throughput activity assay for the Trypanosoma brucei editosome enzyme REL1 and other RNA ligases. Nucleic Acids Res *44*, e24.



## Appendix 2

## Summary of *Edinburgh Infectious Diseases* accounts for FY2015/16

Details	Income (£)	Expenditure (£)
Events		4005.11
Meeting support		1137.1
Network support		1600
Office support		1055.57
Printing		451
Prizes		200
Public Engagement		71.81
Salaries		38436.85
Sponsorship	-3000	
Symposium 2015		3818.74
Symposium 2016		9826.09
UoE	-45600	
Vacation Studentship		1988.12
Total in		-48600
Total out		62590.39
Carried forward	-30024	
Total		-16033.61

## Appendix 3

## News stories from around Edinburgh Infectious Diseases

The full stories can be read at http://www.eid.ed.ac.uk/newsdb

- £10m investment in Roslin Technologies set to boost impact of animal science innovations
- Lung probe developed in Edinburgh spots infections and aims to cut antibiotic overuse



- Members of Moredun Research Institute help launch the SEFARI Centre for Knowledge Exchange & Impact
- Work from the Roslin Institute shows gut macrophage dysregulation key to Inflammatory Bowel Disease
- New study from the School of Biological Sciences finds spread of ages is key to impact of disease
- New musical theatre show to spread message about antimicrobial resistance in schools
- Meriem El Karoui has won £955K from The Wellcome Trust to study how antibiotic resistance emerges
- Professor Dave Robertson has been appointed as Head of the College of Science & Engineering at the University of Edinburgh
- Gene-edited pigs produced at the Roslin Institute show signs of resistance to major viral disease
- Eleanor Riley to be new Director of the Roslin Institute

- Immune cell study from Edinburgh University's Centre for Inflammation Research prompts rethink on how to tackle infections
- New bacteria study from University of Edinburgh finds killing off rivals makes for happy families
- Major drug initiatives are best way to curb threat from parasites shows new study from the University of Edinburgh
- Researchers in the University of Edinburgh's Usher Institute to lead €29m research grant that will assess risk posed by deadly lung infections
- Gut cells are gatekeepers of infectious brain diseases, finds new study from the Roslin Institute's Neil Mabbott
- Infections that pose greatest pandemic risk identified by experts from the Roslin Institute and School of Biological Studies
- Red squirrels in Britain are stricken by medieval strain of leprosy, shows a new study from Edinburgh researchers



- Antibiotics Awareness Week 14 20 November 2016
- Breakthrough in the control of Amoebic Gill Disease by researchers at the Roslin Institute
- Immune cell insight offers hope for tackling the deadly lung condition Acute Respiratory Distress Syndrome

- Migration routes hold key to bird flu spread, global study finds
- New Ker Memorial PhD Studentship available at the University of Edinburgh for 2017
- Andrew Rambaut from Institute of Evolutionary Biology awarded the 2016 Chancellor's Research Award
- Researchers at University of Edinburgh find skin's own defences point towards new eczema therapies
- A global call for global zoonotic TB control and eradication
- Africa Week at the University of Edinburgh highlights infectious disease research in the School of Biological Sciences
- Edinburgh researchers find there is no silver bullet for forests under climate change
- Great day of Biology on Doors Open Day at the University of Edinburgh
- An infectious diseases undergraduate student talks about his experiences on a project at Leiden University Medical Center



- Vaccines; a new weapon in the arsenal to fight against the worms that turned?
- Global superbug response needs \$5bn each year, experts predict
- Bugs' flair for foraging inspires quest for new smart therapies at the University of Edinburgh
- New malaria study by University of Edinburgh researchers shows how multiple infections make disease worse

- Food security focus for major research and teaching initiative at the University of Edinburgh
- New collaboration between Edinburgh researchers and Immunovaccine Inc. to study malaria vaccine
- Researchers at Royal (Dick) School for Veterinary Studies launch red squirrel study to assess leprosy infection
- International Day of Immunology 2016: The Edinburgh Immunology Group
- New study shows Nigerian cattle carry zoonotic pathogens with implications for human health



- Arthritis and other inflammatory conditions could be helped by new insights from the Centre for Inflammation Research
- Roslin Institute scientists part of landmark WHO consultation to galvanise action against Zoonotic Tuberculosis
- New paper published: Beyond killing: can we find new ways to manage infection?
- New PhD position available on Wellcome Trust 4 year Hosts, Pathogens and Global Health PhD programme
- The Roslin Institute receives Scottish Government funding for research into infectious livestock diseases
- Edinburgh researchers part of study shedding light on how malaria parasites jumped from chimps to humans
- Treatment and prevention of life-threatening infections could be improved by research that reveals how bacteria are kept in check
- Roslin Technologies launches to boost animal sciences company creation

- Researchers from the School of Biological Sciences win Royal Society Brian Mercer Feasibility Award
- Bid to beat infant RSV infection boosted by immune study from Centre for Inflammation Research
- Luciana Brondi featured on Sputnik radio programme discussing the current Zika virus outbreak
- Pigs' genetic code altered in bid to tackle infection with deadly African Swine Fever virus
- Ivan Morrison at The Roslin Institute awarded over £0.7M to develop new vaccines against bovine Tuberculosis
- New lecture series on Neglected Tropical Diseases launches for February and March 2016
- New research from the University of Edinburgh argues pregnant mums can hold key to controlling sleeping sickness
- Decades-long quest to beat river blindness edges towards vaccine
- New study charts the hotspots where bat viruses are more likely to infect humans
- Mother appetites can keep size of wild animal groups in check, finds Edinburgh Infectious
  Diseases member Tom Little