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List of Publications by Year in descending order

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| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Close genetic linkage between human and companion animal extraintestinal pathogenic <i>Escherichia coli</i> ST127. <i>Current Research in Microbial Sciences</i> , 2022, 3, 100106. | 1.4 | 9 |
| 2 | Antimicrobial stewardship in companion animal practice: an implementation trial in 135 general practice veterinary clinics. <i>JAC-Antimicrobial Resistance</i> , 2022, 4, dlac015. | 0.9 | 10 |
| 3 | Healthcare-associated infections caused by chlorhexidine-tolerant <i>Serratia marcescens</i> carrying a promiscuous IncHI2 multi-drug resistance plasmid in a veterinary hospital. <i>PLoS ONE</i> , 2022, 17, e0264848. | 1.1 | 6 |
| 4 | Genomic characterisation of an entomopathogenic strain of <i>Serratia ureilytica</i> in the critically endangered phasmid <i>Dryococelus australis</i> . <i>PLoS ONE</i> , 2022, 17, e0265967. | 1.1 | 0 |
| 5 | Efficient disruption of the function of the <i>mnuA</i> nuclease gene using the endogenous CRISPR/Cas system in <i>Mycoplasma gallisepticum</i> . <i>Veterinary Microbiology</i> , 2022, 269, 109436. | 0.8 | 7 |
| 6 | Evaluation of the MiIA ELISA for the diagnosis of herd infection with <i>Mycoplasma bovis</i> using bulk tank milk and estimation of the prevalence of <i>M. bovis</i> in Australia. <i>Veterinary Microbiology</i> , 2022, 270, 109454. | 0.8 | 3 |
| 7 | Genomic and Temporal Trends in Canine ExPEC Reflect Those of Human ExPEC. <i>Microbiology Spectrum</i> , 2022, 10, . | 1.2 | 11 |
| 8 | Bayesian latent class analysis to estimate the optimal cut-off for the MiIA ELISA for the detection of <i>Mycoplasma bovis</i> antibodies in sera, accounting for repeated measures. <i>Preventive Veterinary Medicine</i> , 2022, 205, 105694. | 0.7 | 4 |
| 9 | Water consumption and wastage behaviour in pigs: implications for antimicrobial administration and stewardship. <i>Animal</i> , 2022, 16, 100586. | 1.3 | 2 |
| 10 | Faecal microbiota and antimicrobial resistance gene profiles of healthy foals. <i>Equine Veterinary Journal</i> , 2021, 53, 806-816. | 0.9 | 6 |
| 11 | In-Water Antibiotic Dosing Practices on Pig Farms. <i>Antibiotics</i> , 2021, 10, 169. | 1.5 | 12 |
| 12 | Antimicrobial prescribing guidelines for poultry. <i>Australian Veterinary Journal</i> , 2021, 99, 181-235. | 0.5 | 21 |
| 13 | A <i>Mycoplasma gallisepticum</i> Glycerol ABC Transporter Involved in Pathogenicity. <i>Applied and Environmental Microbiology</i> , 2021, 87, . | 1.4 | 7 |
| 14 | Infectious bronchitis virus in Australia: a model of coronavirus evolution – a review. <i>Avian Pathology</i> , 2021, 50, 295-310. | 0.8 | 3 |
| 15 | Mucosal immune responses in the trachea after chronic infection with <i>Mycoplasma gallisepticum</i> in unvaccinated and vaccinated mature chickens. <i>Cellular Microbiology</i> , 2021, 23, e13383. | 1.1 | 5 |
| 16 | Effect of Drinking Water Distribution System Design on Antimicrobial Delivery to Pigs. <i>Animals</i> , 2021, 11, 2362. | 1.0 | 5 |
| 17 | Effects of immunosuppression on the efficacy of vaccination against <i>Mycoplasma gallisepticum</i> infection in chickens. <i>Veterinary Microbiology</i> , 2021, 260, 109182. | 0.8 | 11 |
| 18 | Antimicrobial stewardship in Australia: the role of qualitative research in programme development. <i>JAC-Antimicrobial Resistance</i> , 2021, 3, dlab166. | 0.9 | 8 |

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|----|---|-----|-----------|
| 19 | Pet Owners and Antibiotics: Knowledge, Opinions, Expectations, and Communication Preferences. <i>Antibiotics</i> , 2021, 10, 1326. | 1.5 | 9 |
| 20 | Water Distribution Systems in Pig Farm Buildings: Critical Elements of Design and Management. <i>Animals</i> , 2021, 11, 3268. | 1.0 | 6 |
| 21 | Genomic comparisons of <i>Escherichia coli</i> ST131 from Australia. <i>Microbial Genomics</i> , 2021, 7, . | 1.0 | 22 |
| 22 | Superinfection and recombination of infectious laryngotracheitis virus vaccines in the natural host. <i>Vaccine</i> , 2020, 38, 7508-7516. | 1.7 | 2 |
| 23 | Detection of naturally aerosolized <i>Actinobacillus pleuropneumoniae</i> on pig farms by cyclonic air sampling and qPCR. <i>Veterinary Microbiology</i> , 2020, 250, 108856. | 0.8 | 4 |
| 24 | Targeted mutagenesis of <i>Mycoplasma gallisepticum</i> using its endogenous CRISPR/Cas system. <i>Veterinary Microbiology</i> , 2020, 250, 108868. | 0.8 | 17 |
| 25 | Contagious Bovine and Caprine Pleuropneumonia: a research community's recommendations for the development of better vaccines. <i>Npj Vaccines</i> , 2020, 5, 66. | 2.9 | 23 |
| 26 | Duration of protective immunity induced by <i>Mycoplasma gallisepticum</i> strain ts-304 vaccine in chickens. <i>Veterinary Microbiology</i> , 2020, 251, 108883. | 0.8 | 11 |
| 27 | Colonization of a hand washing sink in a veterinary hospital by an <i>Enterobacter hormaechei</i> strain carrying multiple resistances to high importance antimicrobials. <i>Antimicrobial Resistance and Infection Control</i> , 2020, 9, 163. | 1.5 | 13 |
| 28 | <i>Mycoplasma bovis</i> mbfN Encodes a Novel LRR Lipoprotein That Undergoes Proteolytic Processing and Binds Host Extracellular Matrix Components. <i>Journal of Bacteriology</i> , 2020, 203, . | 1.0 | 3 |
| 29 | Use of cefovecin in dogs and cats attending first-opinion veterinary practices in Australia. <i>Veterinary Record</i> , 2020, 187, e95. | 0.2 | 14 |
| 30 | Use of Local Antibigram Data and Antimicrobial Importance Ratings to Select Optimal Empirical Therapies for Urinary Tract Infections in Dogs and Cats. <i>Antibiotics</i> , 2020, 9, 924. | 1.5 | 13 |
| 31 | Pathogenesis and tissue tropism of natural field recombinants of infectious laryngotracheitis virus. <i>Veterinary Microbiology</i> , 2020, 243, 108635. | 0.8 | 6 |
| 32 | <i>Mycoplasma bovis</i> Membrane Protein MilA Is a Multifunctional Lipase with Novel Lipid and Glycosaminoglycan Binding Activity. <i>Infection and Immunity</i> , 2020, 88, . | 1.0 | 10 |
| 33 | Antibiotic Resistance Genes in Antibiotic-Free Chicken Farms. <i>Antibiotics</i> , 2020, 9, 120. | 1.5 | 14 |
| 34 | Efficacy of citric acid and sodium hypochlorite as disinfectants against <i>Mycoplasma bovis</i> . <i>Veterinary Microbiology</i> , 2020, 243, 108630. | 0.8 | 10 |
| 35 | Genomic recombination between infectious laryngotracheitis vaccine strains occurs under a broad range of infection conditions in vitro and in ovo. <i>PLoS ONE</i> , 2020, 15, e0229082. | 1.1 | 3 |
| 36 | Differential Response of the Chicken Trachea to Chronic Infection with Virulent <i>Mycoplasma gallisepticum</i> Strain Ap3AS and Vaxsafe MG (Strain ts-304): a Transcriptional Profile. <i>Infection and Immunity</i> , 2020, 88, . | 1.0 | 14 |

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|----|---|-----|-----------|
| 37 | <i>Mycoplasma gallisepticum</i> strain ts-304 is a safe and effective live attenuated vaccine for use in chickens. <i>Veterinary Microbiology</i> , 2020, 244, 108654. | 0.8 | 9 |
| 38 | Antimicrobial prescribing guidelines for pigs. <i>Australian Veterinary Journal</i> , 2020, 98, 105-134. | 0.5 | 5 |
| 39 | Transcriptomic Analysis of Long-Term Protective Immunity Induced by Vaccination With <i>Mycoplasma gallisepticum</i> Strain ts-304. <i>Frontiers in Immunology</i> , 2020, 11, 628804. | 2.2 | 6 |
| 40 | Does only the age of the hen matter in <i>Salmonella enterica</i> contamination of eggs?. <i>Food Microbiology</i> , 2019, 77, 1-9. | 2.1 | 12 |
| 41 | Survey of veterinary prescribing for poultry disease. <i>Australian Veterinary Journal</i> , 2019, 97, 288-288. | 0.5 | 3 |
| 42 | Development of a veterinary antimicrobial stewardship online training program for Australian veterinarians: a national collaborative effort. <i>Australian Veterinary Journal</i> , 2019, 97, 290-291. | 0.5 | 1 |
| 43 | Appraisal of the Australian Veterinary Prescribing Guidelines for antimicrobial prophylaxis for surgery in dogs and cats. <i>Australian Veterinary Journal</i> , 2019, 97, 316-322. | 0.5 | 8 |
| 44 | Comparative genomic analyses of <i>Mycoplasma synoviae</i> vaccine strain MS-H and its wild-type parent strain 86079/7NS: implications for the identification of virulence factors and applications in diagnosis of <i>M. synoviae</i> . <i>Avian Pathology</i> , 2019, 48, 537-548. | 0.8 | 14 |
| 45 | Review: Water medication of growing pigs: sources of between-animal variability in systemic exposure to antimicrobials. <i>Animal</i> , 2019, 13, 3031-3040. | 1.3 | 18 |
| 46 | Exploration of antibiotic resistance risks in a veterinary teaching hospital with Oxford Nanopore long read sequencing. <i>PLoS ONE</i> , 2019, 14, e0217600. | 1.1 | 15 |
| 47 | <i>Salmonella</i> Genomic Island 1B Variant Found in a Sequence Type 117 Avian Pathogenic <i>Escherichia coli</i> Isolate. <i>MSphere</i> , 2019, 4, . | 1.3 | 18 |
| 48 | A combined metabolomic and bioinformatic approach to investigate the function of transport proteins of the important pathogen <i>Mycoplasma bovis</i> . <i>Veterinary Microbiology</i> , 2019, 234, 8-16. | 0.8 | 15 |
| 49 | Vaccination with FAdV-8a induces protection against inclusion body hepatitis caused by homologous and heterologous strains. <i>Avian Pathology</i> , 2019, 48, 396-405. | 0.8 | 7 |
| 50 | Characterisation of the course of <i>Mycoplasma bovis</i> infection in naturally infected dairy herds. <i>Veterinary Microbiology</i> , 2019, 231, 107-115. | 0.8 | 29 |
| 51 | Antimicrobial dosing for common equine drugs: a content review and practical advice for veterinarians in Australia. <i>Australian Veterinary Journal</i> , 2019, 97, 103-107. | 0.5 | 10 |
| 52 | Koala and Wombat Gammaherpesviruses Encode the First Known Viral NTPDase Homologs and Are Phylogenetically Divergent from All Known Gammaherpesviruses. <i>Journal of Virology</i> , 2019, 93, . | 1.5 | 2 |
| 53 | Development and application of high-resolution melting analysis for the classification of infectious laryngotracheitis virus strains and detection of recombinant progeny. <i>Archives of Virology</i> , 2019, 164, 427-438. | 0.9 | 8 |
| 54 | Recommended rejection of the names <i>Malacoplasma</i> gen. nov., <i>Mesomycoplasma</i> gen. nov., <i>Metamycoplasma</i> gen. nov., <i>Metamycoplasmataceae</i> fam. nov., <i>Mycoplasmoidaceae</i> fam. nov., <i>Mycoplasmoidales</i> ord. nov., <i>Mycoplasmoides</i> gen. nov., <i>Mycoplasmopsis</i> gen. nov. [Gupta, Sawnani, Adeolu, Alnajjar and Oren 2018] and all proposed species comb. nov. placed therein. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 69, 3650-3653. | 0.8 | 32 |

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|----|---|-----|-----------|
| 55 | Whole genome sequence analysis of Australian avian pathogenic Escherichia coli that carry the class 1 integrase gene. <i>Microbial Genomics</i> , 2019, 5, . | 1.0 | 51 |
| 56 | Barriers to and enablers of implementing antimicrobial stewardship programs in veterinary practices. <i>Journal of Veterinary Internal Medicine</i> , 2018, 32, 1092-1099. | 0.6 | 77 |
| 57 | Antimicrobial susceptibility testing by Australian veterinary diagnostic laboratories. <i>Australian Veterinary Journal</i> , 2018, 96, 142-146. | 0.5 | 8 |
| 58 | Autoimmune-Disease-Prone NOD Mice Help To Reveal a New Genetic Locus for Reducing Pulmonary Disease Caused by <i>Mycoplasma pulmonis</i> . <i>Infection and Immunity</i> , 2018, 86, . | 1.0 | 1 |
| 59 | The major membrane nuclease MnuA degrades neutrophil extracellular traps induced by <i>Mycoplasma bovis</i> . <i>Veterinary Microbiology</i> , 2018, 218, 13-19. | 0.8 | 49 |
| 60 | Evaluation of <i>Mycoplasma gallisepticum</i> (MG) ts-304 vaccine as a live attenuated vaccine in turkeys. <i>Vaccine</i> , 2018, 36, 2487-2493. | 1.7 | 13 |
| 61 | A Novel <i>Glaesserella</i> sp. Isolated from Pigs with Severe Respiratory Infections Has a Mosaic Genome with Virulence Factors Putatively Acquired by Horizontal Transfer. <i>Applied and Environmental Microbiology</i> , 2018, 84, . | 1.4 | 17 |
| 62 | Antimicrobials used for surgical prophylaxis by equine veterinary practitioners in Australia. <i>Equine Veterinary Journal</i> , 2018, 50, 65-72. | 0.9 | 23 |
| 63 | Determination of the minimum protective dose of a glycoprotein-G-deficient infectious laryngotracheitis virus vaccine delivered via eye-drop to week-old chickens. <i>PLoS ONE</i> , 2018, 13, e0207611. | 1.1 | 2 |
| 64 | <i>Mycoplasma bovis</i> antibody dynamics in naturally exposed dairy calves according to two diagnostic tests. <i>BMC Veterinary Research</i> , 2018, 14, 258. | 0.7 | 13 |
| 65 | Population wide assessment of antimicrobial use in dogs and cats using a novel data source – A cohort study using pet insurance data. <i>Veterinary Microbiology</i> , 2018, 225, 34-39. | 0.8 | 40 |
| 66 | Single Nucleotide Polymorphism Genotyping Analysis Shows That Vaccination Can Limit the Number and Diversity of Recombinant Progeny of Infectious Laryngotracheitis Viruses from the United States. <i>Applied and Environmental Microbiology</i> , 2018, 84, . | 1.4 | 1 |
| 67 | Two subspecies of bent-winged bats (<i>Miniopterus orianae bassanii</i> and <i>oceanensis</i>) in southern Australia have diverse fungal skin flora but not <i>Pseudogymnoascus destructans</i> . <i>PLoS ONE</i> , 2018, 13, e0204282. | 1.1 | 21 |
| 68 | Replication-independent reduction in the number and diversity of recombinant progeny viruses in chickens vaccinated with an attenuated infectious laryngotracheitis vaccine. <i>Vaccine</i> , 2018, 36, 5709-5716. | 1.7 | 3 |
| 69 | Innate immune genes in persistent mating-induced endometritis in horses. <i>Reproduction, Fertility and Development</i> , 2018, 30, 533. | 0.1 | 14 |
| 70 | Antimicrobial labelling in Australia: a threat to antimicrobial stewardship?. <i>Australian Veterinary Journal</i> , 2018, 96, 151-154. | 0.5 | 23 |
| 71 | Veterinary Students’s™ Knowledge and Perceptions About Antimicrobial Stewardship and Biosecurity’s™ A National Survey. <i>Antibiotics</i> , 2018, 7, 34. | 1.5 | 38 |
| 72 | The Performance of Three Immune Assays to Assess the Serological Status of Cattle Experimentally Exposed to <i>Mycoplasma bovis</i> . <i>Veterinary Sciences</i> , 2018, 5, 27. | 0.6 | 9 |

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|----|---|-----|-----------|
| 73 | Genome analysis of <i>Mycoplasma synoviae</i> strain MS-H, the most common <i>M. synoviae</i> strain with a worldwide distribution. <i>BMC Genomics</i> , 2018, 19, 117. | 1.2 | 14 |
| 74 | Metabolite profiling of <i>Mycoplasma gallisepticum</i> mutants, combined with bioinformatic analysis, can reveal the likely functions of virulence-associated genes. <i>Veterinary Microbiology</i> , 2018, 223, 160-167. | 0.8 | 10 |
| 75 | First detection of bovine noroviruses and detection of bovine coronavirus in Australian dairy cattle. <i>Australian Veterinary Journal</i> , 2018, 96, 203-208. | 0.5 | 7 |
| 76 | Analysis of the <i>Mycoplasma bovis</i> lactate dehydrogenase reveals typical enzymatic activity despite the presence of an atypical catalytic site motif. <i>Microbiology (United Kingdom)</i> , 2018, 164, 186-193. | 0.7 | 4 |
| 77 | Antimicrobial Prescribing in Dogs and Cats in Australia: Results of the Australasian Infectious Disease Advisory Panel Survey. <i>Journal of Veterinary Internal Medicine</i> , 2017, 31, 1100-1107. | 0.6 | 44 |
| 78 | Antimicrobials used for surgical prophylaxis by companion animal veterinarians in Australia. <i>Veterinary Microbiology</i> , 2017, 203, 301-307. | 0.8 | 25 |
| 79 | Immune responses to vaccination and infection with <i>Mycoplasma gallisepticum</i> in turkeys. <i>Avian Pathology</i> , 2017, 46, 464-473. | 0.8 | 7 |
| 80 | The <i>oppD</i> Gene and Putative Peptidase Genes May Be Required for Virulence in <i>Mycoplasma gallisepticum</i> . <i>Infection and Immunity</i> , 2017, 85, . | 1.0 | 12 |
| 81 | Natural recombination in alphaherpesviruses: Insights into viral evolution through full genome sequencing and sequence analysis. <i>Infection, Genetics and Evolution</i> , 2017, 49, 174-185. | 1.0 | 45 |
| 82 | Safety and efficacy of a <i>Mycoplasma gallisepticum oppD</i> knockout mutant as a vaccine candidate. <i>Vaccine</i> , 2017, 35, 6248-6253. | 1.7 | 2 |
| 83 | Reproduction of respiratory mycoplasmosis in calves by exposure to an aerosolised culture of <i>Mycoplasma bovis</i> . <i>Veterinary Microbiology</i> , 2017, 210, 167-173. | 0.8 | 25 |
| 84 | Survey of Victorian small ruminant herds for mycoplasmas associated with contagious agalactia and molecular characterisation of <i>Mycoplasma mycoides</i> subspecies <i>capri</i> isolates from one herd. <i>Australian Veterinary Journal</i> , 2017, 95, 392-400. | 0.5 | 4 |
| 85 | Genetic Diversity of Infectious Laryngotracheitis Virus during In Vivo Coinfection Parallels Viral Replication and Arises from Recombination Hot Spots within the Genome. <i>Applied and Environmental Microbiology</i> , 2017, 83, . | 1.4 | 16 |
| 86 | Comparative Metabolomics of <i>Mycoplasma bovis</i> and <i>Mycoplasma gallisepticum</i> Reveals Fundamental Differences in Active Metabolic Pathways and Suggests Novel Gene Annotations. <i>MSystems</i> , 2017, 2, . | 1.7 | 35 |
| 87 | Cross-sectional study of antimicrobials used for surgical prophylaxis by bovine veterinary practitioners in Australia. <i>Veterinary Record</i> , 2017, 181, 426-426. | 0.2 | 14 |
| 88 | Identification of a new genetic marker in <i>Mycoplasma synoviae</i> vaccine strain MS-H and development of a strategy using polymerase chain reaction and high-resolution melting curve analysis for differentiating MS-H from field strains. <i>Veterinary Microbiology</i> , 2017, 210, 49-55. | 0.8 | 14 |
| 89 | Chronologic Analysis of Gross and Histologic Lesions Induced by Field Strains of FAdV-1, FAdV-8b, and FAdV-11 in Six-Week-Old Chickens. <i>Avian Diseases</i> , 2017, 61, 512. | 0.4 | 8 |
| 90 | Improvements in diagnosis of disease caused by <i>Mycoplasma bovis</i> in cattle. <i>Animal Production Science</i> , 2017, 57, 1482. | 0.6 | 7 |

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|-----|---|-----|-----------|
| 91 | A high prevalence of beak and feather disease virus in non-psittacine Australian birds. <i>Journal of Medical Microbiology</i> , 2017, 66, 1005-1013. | 0.7 | 50 |
| 92 | Development and application of a TaqMan single nucleotide polymorphism genotyping assay to study infectious laryngotracheitis virus recombination in the natural host. <i>PLoS ONE</i> , 2017, 12, e0174590. | 1.1 | 16 |
| 93 | Investigation of a novel porcine bacterium by whole genome sequencing and mouse inoculation. <i>Animal Production Science</i> , 2017, 57, 2494. | 0.6 | 0 |
| 94 | Low genetic diversity among historical and contemporary clinical isolates of felid herpesvirus 1. <i>BMC Genomics</i> , 2016, 17, 704. | 1.2 | 20 |
| 95 | Molecular epidemiology of an outbreak of clinical mastitis in sheep caused by <i>Mannheimia haemolytica</i> . <i>Veterinary Microbiology</i> , 2016, 191, 82-87. | 0.8 | 11 |
| 96 | Impacts of poultry vaccination on viruses of wild bird. <i>Current Opinion in Virology</i> , 2016, 19, 23-29. | 2.6 | 16 |
| 97 | Effect of ovarian hormones on the healthy equine uterus: a global gene expression analysis. <i>Reproduction, Fertility and Development</i> , 2016, 28, 1810. | 0.1 | 7 |
| 98 | Sequence and recombination analyses of archived field isolates of equine herpesviruses 1 and 4. <i>Journal of Equine Veterinary Science</i> , 2016, 39, S70-S71. | 0.4 | 0 |
| 99 | Evaluation of an IgG Enzyme-Linked Immunosorbent Assay as a Serological Assay for Detection of <i>Mycoplasma bovis</i> Infection in Feedlot Cattle. <i>Journal of Clinical Microbiology</i> , 2016, 54, 1269-1275. | 1.8 | 33 |
| 100 | Development of a blocking ELISA for detection of <i>Mycoplasma hyopneumoniae</i> infection based on a monoclonal antibody against protein P65. <i>Journal of Veterinary Medical Science</i> , 2016, 78, 1319-1322. | 0.3 | 2 |
| 101 | Full genome analysis of Australian infectious bronchitis viruses suggests frequent recombination events between vaccine strains and multiple phylogenetically distant avian coronaviruses of unknown origin. <i>Veterinary Microbiology</i> , 2016, 197, 27-38. | 0.8 | 25 |
| 102 | Oestrous cycle-dependent equine uterine immune response to induced infectious endometritis. <i>Veterinary Research</i> , 2016, 47, 110. | 1.1 | 15 |
| 103 | The first genome sequence of a metatherian herpesvirus: <i>Macropodid herpesvirus 1</i> . <i>BMC Genomics</i> , 2016, 17, 70. | 1.2 | 7 |
| 104 | Evidence of widespread natural recombination among field isolates of equine herpesvirus 4 but not among field isolates of equine herpesvirus 1. <i>Journal of General Virology</i> , 2016, 97, 747-755. | 1.3 | 28 |
| 105 | Effect of differing +2 amino acids on export of a heterologous PhoA lipoprotein in <i>Mycoplasma gallisepticum</i> . <i>Microbiology (United Kingdom)</i> , 2016, 162, 1300-1309. | 0.7 | 3 |
| 106 | Marsupial and monotreme serum immunoglobulin binding by proteins A, G and L and anti-kangaroo antibody. <i>Journal of Immunological Methods</i> , 2015, 427, 94-99. | 0.6 | 7 |
| 107 | Deep sequencing of the uterine immune response to bacteria during the equine oestrous cycle. <i>BMC Genomics</i> , 2015, 16, 934. | 1.2 | 22 |
| 108 | The Effect of an Alternate Start Codon on Heterologous Expression of a PhoA Fusion Protein in <i>Mycoplasma gallisepticum</i> . <i>PLoS ONE</i> , 2015, 10, e0127911. | 1.1 | 16 |

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|-----|---|-----|-----------|
| 109 | Protection Induced in Broiler Chickens following Drinking-Water Delivery of Live Infectious Laryngotracheitis Vaccines against Subsequent Challenge with Recombinant Field Virus. PLoS ONE, 2015, 10, e0137719. | 1.1 | 8 |
| 110 | Analysis of the complete genomic sequences of two virus subpopulations of the Australian infectious bronchitis virus vaccine VicS. Avian Pathology, 2015, 44, 182-191. | 0.8 | 13 |
| 111 | Development of a <i>Mycoplasma gallisepticum</i> infection model in turkeys. Avian Pathology, 2015, 44, 35-42. | 0.8 | 17 |
| 112 | Disruption of the Membrane Nuclease Gene (MBOVPG45_0215) of <i>Mycoplasma bovis</i> Greatly Reduces Cellular Nuclease Activity. Journal of Bacteriology, 2015, 197, 1549-1558. | 1.0 | 34 |
| 113 | Chronological analysis of gross and histological lesions induced by field strains of fowl adenovirus serotypes 1, 8b and 11 in one-day-old chickens. Avian Pathology, 2015, 44, 106-113. | 0.8 | 56 |
| 114 | Genetic diversity of <i>Mycoplasma arginini</i> isolates based on multilocus sequence typing. Veterinary Microbiology, 2015, 180, 123-128. | 0.8 | 11 |
| 115 | Novel assay to quantify recombination in a calicivirus. Veterinary Microbiology, 2015, 177, 25-31. | 0.8 | 8 |
| 116 | Human Wound Infection with <i>Mannheimia glucosida</i> following Lamb Bite. Journal of Clinical Microbiology, 2015, 53, 3374-3376. | 1.8 | 8 |
| 117 | Evidence of apoptosis induced by viral protein 2 of chicken anaemia virus. Archives of Virology, 2015, 160, 2557-2563. | 0.9 | 11 |
| 118 | The upper respiratory tract is a natural reservoir of haemolytic <i>Mannheimia</i> species associated with ovine mastitis. Veterinary Microbiology, 2015, 181, 308-312. | 0.8 | 5 |
| 119 | Development and application of molecular methods (qPCR) for detection of Tasmanian Atlantic salmon reovirus. Journal of Fish Diseases, 2015, 38, 739-754. | 0.9 | 8 |
| 120 | Development and Host Compatibility of Plasmids for Two Important Ruminant Pathogens, <i>Mycoplasma bovis</i> and <i>Mycoplasma agalactiae</i> . PLoS ONE, 2015, 10, e0119000. | 1.1 | 8 |
| 121 | Growth Kinetics and Transmission Potential of Existing and Emerging Field Strains of Infectious Laryngotracheitis Virus. PLoS ONE, 2015, 10, e0120282. | 1.1 | 24 |
| 122 | Genes Found Essential in Other Mycoplasmas Are Dispensable in <i>Mycoplasma bovis</i> . PLoS ONE, 2014, 9, e97100. | 1.1 | 32 |
| 123 | Development of a Recombinant Protein-Based Enzyme-Linked Immunosorbent Assay for Diagnosis of <i>Mycoplasma bovis</i> Infection in Cattle. Vaccine Journal, 2014, 21, 196-202. | 3.2 | 77 |
| 124 | Avian pathogenic <i>Escherichia coli</i> Δ tonB mutants are safe and protective live-attenuated vaccine candidates. Veterinary Microbiology, 2014, 173, 289-298. | 0.8 | 9 |
| 125 | Evaluation of a novel strain of infectious bronchitis virus emerged as a result of spike gene recombination between two highly diverged parent strains. Avian Pathology, 2014, 43, 249-257. | 0.8 | 17 |
| 126 | The spatial and temporal variation of the distribution and prevalence of Atlantic salmon reovirus (TSRV) infection in Tasmania, Australia. Preventive Veterinary Medicine, 2014, 116, 214-219. | 0.7 | 5 |

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|-----|---|-----|-----------|
| 127 | Sequence diversity, cytotoxicity and antigenic similarities of the leukotoxin of isolates of Mannheimia species from mastitis in domestic sheep. <i>Veterinary Microbiology</i> , 2014, 174, 172-179. | 0.8 | 10 |
| 128 | Expression of equine β -defensin 110, 123 and 124 genes in the stallion reproductive tract. <i>Journal of Equine Veterinary Science</i> , 2014, 34, 69. | 0.4 | 1 |
| 129 | The Mycoplasma gallisepticum Virulence Factor Lipoprotein MslA Is a Novel Polynucleotide Binding Protein. <i>Infection and Immunity</i> , 2013, 81, 3220-3226. | 1.0 | 26 |
| 130 | Differential transcription patterns in wild-type and glycoprotein G-deleted infectious laryngotracheitis viruses. <i>Avian Pathology</i> , 2013, 42, 253-259. | 0.8 | 7 |
| 131 | Membrane proteins of Mycoplasma bovis and their role in pathogenesis. <i>Research in Veterinary Science</i> , 2013, 95, 321-325. | 0.9 | 21 |
| 132 | Cross-Protective Immune Responses Between Genotypically Distinct Lineages of Infectious Laryngotracheitis Viruses. <i>Avian Diseases</i> , 2013, 58, 147. | 0.4 | 3 |
| 133 | The role of Type 1, P and S fimbriae in binding of Escherichia coli to the canine endometrium. <i>Veterinary Microbiology</i> , 2013, 164, 399-404. | 0.8 | 11 |
| 134 | Equine rotaviruses—Current understanding and continuing challenges. <i>Veterinary Microbiology</i> , 2013, 167, 135-144. | 0.8 | 34 |
| 135 | MalF is essential for persistence of Mycoplasma gallisepticum in vivo. <i>Microbiology (United Kingdom)</i> , 2013, 159, 1459-1470. | 0.7 | 26 |
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| 273 | Physical mapping of the genomic heterogeneity of isolates of equine herpesvirus 2 (equine) Tj ETQq1 1 0.784314 rgBT /Overlock 10 T | 0.9 | 27 |
| 274 | Latency of equine herpesvirus 4. <i>Veterinary Record</i> , 1989, 124, 203-203. | 0.2 | 1 |
| 275 | Asinine herpesvirus genomes: comparison with those of the equine herpesviruses. <i>Archives of Virology</i> , 1988, 101, 183-190. | 0.9 | 54 |
| 276 | Examination of mycoplasma gallisepticum strains using restriction endonuclease DNA analysis and DNA-DNA hybridisation. <i>Avian Pathology</i> , 1988, 17, 559-570. | 0.8 | 49 |
| 277 | Latency of equine herpesvirus 4 (equine rhinopneumonitis virus). <i>Veterinary Record</i> , 1988, 123, 518-519. | 0.2 | 36 |
| 278 | Genomic Heterogeneity of Equine Betaherpesviruses. <i>Journal of General Virology</i> , 1987, 68, 1441-1447. | 1.3 | 74 |
| 279 | Epidemiology of equine herpesvirus 2 (equine cytomegalovirus). <i>Journal of Clinical Microbiology</i> , 1987, 25, 13-16. | 1.8 | 44 |
| 280 | Spastic paresis in a Poll Hereford heifer. <i>Australian Veterinary Journal</i> , 1986, 63, 367-369. | 0.5 | 8 |
| 281 | Equine herpesvirus genomes: Heterogeneity of naturally occurring type 4 isolates and of a type 1 isolate after heterologous cell passage. <i>Archives of Virology</i> , 1986, 91, 375-381. | 0.9 | 18 |
| 282 | Bovine encephalitis herpesvirus is different from bovine herpesvirus 1 and from other ruminant herpesviruses. <i>Australian Veterinary Journal</i> , 1985, 62, 149-150. | 0.5 | 0 |