

Gf Browning

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

269
papers

5,896
citations

41
h-index

58
g-index

291
ext. papers

6,721
ext. citations

3.5
avg, IF

5.54
L-index

#	Paper	IF	Citations
269	Close genetic linkage between human and companion animal extraintestinal pathogenic ST127.. <i>Current Research in Microbial Sciences</i> , 2022 , 3, 100106	3.3	3
268	Antimicrobial stewardship in companion animal practice: an implementation trial in 135 general practice veterinary clinics.. <i>JAC-Antimicrobial Resistance</i> , 2022 , 4, d1ac015	2.9	0
267	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	3.7	1
266	Genomic characterisation of an entomopathogenic strain of <i>Serratia ureilytica</i> in the critically endangered phasid <i>Dryococelus australis</i> .. <i>PLoS ONE</i> , 2022 , 17, e0265967	3.7	
265	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	3.3	1
264	Evaluation of the MilA 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 , 109454	3.3	0
263	Antimicrobial stewardship in Australia: the role of qualitative research in programme development. <i>JAC-Antimicrobial Resistance</i> , 2021 , 3, d1ab166	2.9	2
262	Faecal microbiota and antimicrobial resistance gene profiles of healthy foals. <i>Equine Veterinary Journal</i> , 2021 , 53, 806-816	2.4	2
261	Antimicrobial prescribing guidelines for poultry. <i>Australian Veterinary Journal</i> , 2021 , 99, 181-235	1.2	3
260	A Glycerol ABC Transporter Involved in Pathogenicity. <i>Applied and Environmental Microbiology</i> , 2021 , 87,	4.8	1
259	In-Water Antibiotic Dosing Practices on Pig Farms. <i>Antibiotics</i> , 2021 , 10,	4.9	3
258	Infectious bronchitis virus in Australia: a model of coronavirus evolution - a review. <i>Avian Pathology</i> , 2021 , 50, 295-310	2.4	
257	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	3.9	1
256	Effects of immunosuppression on the efficacy of vaccination against <i>Mycoplasma gallisepticum</i> infection in chickens. <i>Veterinary Microbiology</i> , 2021 , 260, 109182	3.3	3
255	Genomic comparisons of ST131 from Australia.. <i>Microbial Genomics</i> , 2021 , 7,	4.4	5
254	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,	4.9	5
253	Pathogenesis and tissue tropism of natural field recombinants of infectious laryngotracheitis virus. <i>Veterinary Microbiology</i> , 2020 , 243, 108635	3.3	5

252	Mycoplasma bovis Membrane Protein MilA Is a Multifunctional Lipase with Novel Lipid and Glycosaminoglycan Binding Activity. <i>Infection and Immunity</i> , 2020 , 88,	3.7	5
251	Antibiotic Resistance Genes in Antibiotic-Free Chicken Farms. <i>Antibiotics</i> , 2020 , 9,	4.9	8
250	Efficacy of citric acid and sodium hypochlorite as disinfectants against Mycoplasma bovis. <i>Veterinary Microbiology</i> , 2020 , 243, 108630	3.3	7
249	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	3.7	3
248	Differential Response of the Chicken Trachea to Chronic Infection with Virulent Mycoplasma gallisepticum Strain Ap3AS and Vaxsafe MG (Strain ts-304): a Transcriptional Profile. <i>Infection and Immunity</i> , 2020 , 88,	3.7	7
247	Superinfection and recombination of infectious laryngotracheitis virus vaccines in the natural host. <i>Vaccine</i> , 2020 , 38, 7508-7516	4.1	0
246	Detection of naturally aerosolized Actinobacillus pleuropneumoniae on pig farms by cyclonic air sampling and qPCR. <i>Veterinary Microbiology</i> , 2020 , 250, 108856	3.3	1
245	Targeted mutagenesis of Mycoplasma gallisepticum using its endogenous CRISPR/Cas system. <i>Veterinary Microbiology</i> , 2020 , 250, 108868	3.3	5
244	Contagious Bovine and Caprine Pleuropneumonia: a research community's recommendations for the development of better vaccines. <i>Npj Vaccines</i> , 2020 , 5, 66	9.5	6
243	Duration of protective immunity induced by Mycoplasma gallisepticum strain ts-304 vaccine in chickens. <i>Veterinary Microbiology</i> , 2020 , 251, 108883	3.3	7
242	Colonization of a hand washing sink in a veterinary hospital by an Enterobacter hormaechei strain carrying multiple resistances to high importance antimicrobials. <i>Antimicrobial Resistance and Infection Control</i> , 2020 , 9, 163	6.2	5
241	Use of ceftiofur in dogs and cats attending first-opinion veterinary practices in Australia. <i>Veterinary Record</i> , 2020 , 187, e95	0.9	6
240	Mycoplasma gallisepticum strain ts-304 is a safe and effective live attenuated vaccine for use in chickens. <i>Veterinary Microbiology</i> , 2020 , 244, 108654	3.3	4
239	Antimicrobial prescribing guidelines for pigs. <i>Australian Veterinary Journal</i> , 2020 , 98, 105-134	1.2	2
238	Transcriptomic Analysis of Long-Term Protective Immunity Induced by Vaccination With Strain ts-304. <i>Frontiers in Immunology</i> , 2020 , 11, 628804	8.4	2
237	Review: Water medication of growing pigs: sources of between-animal variability in systemic exposure to antimicrobials. <i>Animal</i> , 2019 , 13, 3031-3040	3.1	9
236	Exploration of antibiotic resistance risks in a veterinary teaching hospital with Oxford Nanopore long read sequencing. <i>PLoS ONE</i> , 2019 , 14, e0217600	3.7	8
235	Genomic Island 1B Variant Found in a Sequence Type 117 Avian Pathogenic Escherichia coli Isolate. <i>MSphere</i> , 2019 , 4,	5	12

234	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	3.3	6
233	Vaccination with FAdV-8a induces protection against inclusion body hepatitis caused by homologous and heterologous strains. <i>Avian Pathology</i> , 2019 , 48, 396-405	2.4	4
232	Characterisation of the course of <i>Mycoplasma bovis</i> infection in naturally infected dairy herds. <i>Veterinary Microbiology</i> , 2019 , 231, 107-115	3.3	17
231	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	1.2	7
230	Does only the age of the hen matter in <i>Salmonella enterica</i> contamination of eggs?. <i>Food Microbiology</i> , 2019 , 77, 1-9	6	6
229	Survey of veterinary prescribing for poultry disease. <i>Australian Veterinary Journal</i> , 2019 , 97, 288	1.2	3
228	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	1.2	1
227	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	1.2	3
226	Comparative genomic analyses of 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>Avian Pathology</i> , 2019 , 48, 537-548	2.4	4
225	Recommended rejection of the names gen. nov., gen. nov., gen. nov., fam. nov., fam. nov., ord. nov., gen. nov., gen. nov. [Gupta, Sawnani, Adeolu, Alnajar and Oren 2018] and all proposed species comb. nov. placed therein. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019 , 69, 3650-3653	2.2	15
224	Whole genome sequence analysis of Australian avian pathogenic <i>Escherichia coli</i> that carry the class 1 integrase gene. <i>Microbial Genomics</i> , 2019 , 5,	4.4	31
223	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,	6.6	2
222	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	2.6	6
221	Barriers to and enablers of implementing antimicrobial stewardship programs in veterinary practices. <i>Journal of Veterinary Internal Medicine</i> , 2018 , 32, 1092-1099	3.1	45
220	Antimicrobial susceptibility testing by Australian veterinary diagnostic laboratories. <i>Australian Veterinary Journal</i> , 2018 , 96, 142-146	1.2	8
219	The major membrane nuclease MnuA degrades neutrophil extracellular traps induced by <i>Mycoplasma bovis</i> . <i>Veterinary Microbiology</i> , 2018 , 218, 13-19	3.3	23
218	Evaluation of <i>Mycoplasma gallisepticum</i> (MG) ts-304 vaccine as a live attenuated vaccine in turkeys. <i>Vaccine</i> , 2018 , 36, 2487-2493	4.1	8
217	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,	4.8	4

216	Antimicrobials used for surgical prophylaxis by equine veterinary practitioners in Australia. <i>Equine Veterinary Journal</i> , 2018 , 50, 65-72	2.4	14
215	Veterinary Students Knowledge and Perceptions About Antimicrobial Stewardship and Biosecurity-A National Survey. <i>Antibiotics</i> , 2018 , 7,	4.9	23
214	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,	2.4	4
213	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	4.5	10
212	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	3.3	8
211	First detection of bovine noroviruses and detection of bovine coronavirus in Australian dairy cattle. <i>Australian Veterinary Journal</i> , 2018 , 96, 203-208	1.2	5
210	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	2.9	2
209	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, e0207617	3.7	1
208	<i>Mycoplasma bovis</i> antibody dynamics in naturally exposed dairy calves according to two diagnostic tests. <i>BMC Veterinary Research</i> , 2018 , 14, 258	2.7	7
207	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	3.3	26
206	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,	4.8	1
205	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	3.7	15
204	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	4.1	3
203	Innate immune genes in persistent mating-induced endometritis in horses. <i>Reproduction, Fertility and Development</i> , 2018 , 30, 533-545	1.8	6
202	Antimicrobial labelling in Australia: a threat to antimicrobial stewardship?. <i>Australian Veterinary Journal</i> , 2018 , 96, 151-154	1.2	15
201	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	3.1	33
200	Antimicrobials used for surgical prophylaxis by companion animal veterinarians in Australia. <i>Veterinary Microbiology</i> , 2017 , 203, 301-307	3.3	20
199	Immune responses to vaccination and infection with <i>Mycoplasma gallisepticum</i> in turkeys. <i>Avian Pathology</i> , 2017 , 46, 464-473	2.4	6

198	The Gene and Putative Peptidase Genes May Be Required for Virulence in <i>Mycoplasma gallisepticum</i> . <i>Infection and Immunity</i> , 2017 , 85,	3.7	11
197	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	4.5	24
196	Safety and efficacy of a <i>Mycoplasma gallisepticum</i> oppD knockout mutant as a vaccine candidate. <i>Vaccine</i> , 2017 , 35, 6248-6253	4.1	1
195	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	3.3	18
194	Survey of Victorian small ruminant herds for mycoplasmas associated with contagious agalactia and molecular characterisation of <i>Mycoplasma mycoides</i> subspecies capri isolates from one herd. <i>Australian Veterinary Journal</i> , 2017 , 95, 392-400	1.2	3
193	Genetic Diversity of Infectious Laryngotracheitis Virus during Coinfection Parallels Viral Replication and Arises from Recombination Hot Spots within the Genome. <i>Applied and Environmental Microbiology</i> , 2017 , 83,	4.8	13
192	Comparative Metabolomics of and Reveals Fundamental Differences in Active Metabolic Pathways and Suggests Novel Gene Annotations. <i>MSystems</i> , 2017 , 2,	7.6	17
191	Improvements in diagnosis of disease caused by <i>Mycoplasma bovis</i> in cattle. <i>Animal Production Science</i> , 2017 , 57, 1482	1.4	3
190	Cross-sectional study of antimicrobials used for surgical prophylaxis by bovine veterinary practitioners in Australia. <i>Veterinary Record</i> , 2017 , 181, 426	0.9	8
189	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	3.3	10
188	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, 512-519	1.6	5
187	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	3.7	15
186	A high prevalence of beak and feather disease virus in non-psittacine Australian birds. <i>Journal of Medical Microbiology</i> , 2017 , 66, 1005-1013	3.2	34
185	Investigation of a novel porcine bacterium by whole genome sequencing and mouse inoculation. <i>Animal Production Science</i> , 2017 , 57, 2494	1.4	
184	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, 1319-22	1.1	2
183	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	3.3	23
182	Oestrous cycle-dependent equine uterine immune response to induced infectious endometritis. <i>Veterinary Research</i> , 2016 , 47, 110	3.8	7
181	The first genome sequence of a metatherian herpesvirus: Macropodid herpesvirus 1. <i>BMC Genomics</i> , 2016 , 17, 70	4.5	5

180	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	4.9	21
179	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	2.9	2
178	Low genetic diversity among historical and contemporary clinical isolates of felid herpesvirus 1. <i>BMC Genomics</i> , 2016 , 17, 704	4.5	13
177	Molecular epidemiology of an outbreak of clinical mastitis in sheep caused by <i>Mannheimia haemolytica</i> . <i>Veterinary Microbiology</i> , 2016 , 191, 82-7	3.3	8
176	Impacts of poultry vaccination on viruses of wild bird. <i>Current Opinion in Virology</i> , 2016 , 19, 23-9	7.5	14
175	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, 1269-75	9.7	23
174	Disruption of the membrane nuclease gene (MBOVPG45_0215) of <i>Mycoplasma bovis</i> greatly reduces cellular nuclease activity. <i>Journal of Bacteriology</i> , 2015 , 197, 1549-58	3.5	17
173	Chronological analysis of gross and histological lesions induced by field strains of fowl adenovirus serotypes 1, 8b and 11 in one-day-old chickens. <i>Avian Pathology</i> , 2015 , 44, 106-13	2.4	41
172	Genetic diversity of <i>Mycoplasma arginini</i> isolates based on multilocus sequence typing. <i>Veterinary Microbiology</i> , 2015 , 180, 123-8	3.3	10
171	Novel assay to quantify recombination in a calicivirus. <i>Veterinary Microbiology</i> , 2015 , 177, 25-31	3.3	5
170	Human Wound Infection with <i>Mannheimia glucosida</i> following Lamb Bite. <i>Journal of Clinical Microbiology</i> , 2015 , 53, 3374-6	9.7	5
169	Evidence of apoptosis induced by viral protein 2 of chicken anaemia virus. <i>Archives of Virology</i> , 2015 , 160, 2557-63	2.6	9
168	The upper respiratory tract is a natural reservoir of haemolytic <i>Mannheimia</i> species associated with ovine mastitis. <i>Veterinary Microbiology</i> , 2015 , 181, 308-12	3.3	4
167	Development and application of molecular methods (PCR) for detection of Tasmanian Atlantic salmon reovirus. <i>Journal of Fish Diseases</i> , 2015 , 38, 739-54	2.6	8
166	Marsupial and monotreme serum immunoglobulin binding by proteins A, G and L and anti-kangaroo antibody. <i>Journal of Immunological Methods</i> , 2015 , 427, 94-9	2.5	4
165	Deep sequencing of the uterine immune response to bacteria during the equine oestrous cycle. <i>BMC Genomics</i> , 2015 , 16, 934	4.5	20
164	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	3.7	14
163	Protection Induced in Broiler Chickens following Drinking-Water Delivery of Live Infectious Laryngotracheitis Vaccines against Subsequent Challenge with Recombinant Field Virus. <i>PLoS ONE</i> , 2015 , 10, e0137719	3.7	4

162	Analysis of the complete genomic sequences of two virus subpopulations of the Australian infectious bronchitis virus vaccine VicS. <i>Avian Pathology</i> , 2015 , 44, 182-91	2.4	10
161	Development of a <i>Mycoplasma gallisepticum</i> infection model in turkeys. <i>Avian Pathology</i> , 2015 , 44, 35-42.	4.4	12
160	Development and host compatibility of plasmids for two important ruminant pathogens, <i>Mycoplasma bovis</i> and <i>Mycoplasma agalactiae</i> . <i>PLoS ONE</i> , 2015 , 10, e0119000	3.7	4
159	Growth kinetics and transmission potential of existing and emerging field strains of infectious laryngotracheitis virus. <i>PLoS ONE</i> , 2015 , 10, e0120282	3.7	23
158	Effect of ovarian hormones on the healthy equine uterus: a global gene expression analysis. <i>Reproduction, Fertility and Development</i> , 2015 ,	1.8	6
157	Avian pathogenic <i>Escherichia coli</i> Φ onB mutants are safe and protective live-attenuated vaccine candidates. <i>Veterinary Microbiology</i> , 2014 , 173, 289-98	3.3	4
156	Evaluation of a novel strain of infectious bronchitis virus emerged as a result of spike gene recombination between two highly diverged parent strains. <i>Avian Pathology</i> , 2014 , 43, 249-57	2.4	14
155	The spatial and temporal variation of the distribution and prevalence of Atlantic salmon reovirus (TSRV) infection in Tasmania, Australia. <i>Preventive Veterinary Medicine</i> , 2014 , 116, 214-9	3.1	4
154	Sequence diversity, cytotoxicity and antigenic similarities of the leukotoxin of isolates of <i>Mannheimia</i> species from mastitis in domestic sheep. <i>Veterinary Microbiology</i> , 2014 , 174, 172-9	3.3	8
153	Cross-protective immune responses between genotypically distinct lineages of infectious laryngotracheitis viruses. <i>Avian Diseases</i> , 2014 , 58, 147-52	1.6	2
152	Genes found essential in other mycoplasmas are dispensable in <i>Mycoplasma bovis</i> . <i>PLoS ONE</i> , 2014 , 9, e97100	3.7	21
151	Development of a recombinant protein-based enzyme-linked immunosorbent assay for diagnosis of <i>Mycoplasma bovis</i> infection in cattle. <i>Vaccine Journal</i> , 2014 , 21, 196-202		60
150	The <i>Mycoplasma gallisepticum</i> virulence factor lipoprotein MslA is a novel polynucleotide binding protein. <i>Infection and Immunity</i> , 2013 , 81, 3220-6	3.7	23
149	Differential transcription patterns in wild-type and glycoprotein G-deleted infectious laryngotracheitis viruses. <i>Avian Pathology</i> , 2013 , 42, 253-9	2.4	6
148	Membrane proteins of <i>Mycoplasma bovis</i> and their role in pathogenesis. <i>Research in Veterinary Science</i> , 2013 , 95, 321-5	2.5	17
147	The role of Type 1, P and S fimbriae in binding of <i>Escherichia coli</i> to the canine endometrium. <i>Veterinary Microbiology</i> , 2013 , 164, 399-404	3.3	9
146	Equine rotaviruses--current understanding and continuing challenges. <i>Veterinary Microbiology</i> , 2013 , 167, 135-44	3.3	26
145	MalF is essential for persistence of <i>Mycoplasma gallisepticum</i> in vivo. <i>Microbiology (United Kingdom)</i> , 2013 , 159, 1459-1470	2.9	15

144	Isolation and characterization of a novel herpesvirus from a free-ranging eastern grey kangaroo (<i>Macropus giganteus</i>). <i>Journal of Wildlife Diseases</i> , 2013 , 49, 143-51	1.3	15
143	Challenges and recent advancements in infectious laryngotracheitis virus vaccines. <i>Avian Pathology</i> , 2013 , 42, 195-205	2.4	39
142	Methicillin-resistant <i>Staphylococcus aureus</i> : an issue for veterinary hospitals. <i>Australian Veterinary Journal</i> , 2013 , 91, 215-9	1.2	4
141	Phylogenetic and molecular epidemiological studies reveal evidence of multiple past recombination events between infectious laryngotracheitis viruses. <i>PLoS ONE</i> , 2013 , 8, e55121	3.7	28
140	Histochemical and morphometric characterization of broncho-pneumonia in calves caused by infection with <i>Mycoplasma bovis</i> . <i>Veterinary Microbiology</i> , 2012 , 158, 220-4	3.3	8
139	Air sampling in the breathing zone of neonatal foals for prediction of subclinical <i>Rhodococcus equi</i> infection. <i>Equine Veterinary Journal</i> , 2012 , 44, 203-6	2.4	4
138	Uropathogenic virulence factor FimH facilitates binding of uteropathogenic <i>Escherichia coli</i> to canine endometrium. <i>Comparative Immunology, Microbiology and Infectious Diseases</i> , 2012 , 35, 461-7	2.6	18
137	Safety and vaccine efficacy of a glycoprotein G deficient strain of infectious laryngotracheitis virus delivered in ovo. <i>Vaccine</i> , 2012 , 30, 7193-8	4.1	13
136	A novel transposon construct expressing PhoA with potential for studying protein expression and translocation in <i>Mycoplasma gallisepticum</i> . <i>BMC Microbiology</i> , 2012 , 12, 138	4.5	9
135	Effect of simulated stages of the canine oestrous cycle on <i>Escherichia coli</i> binding to canine endometrium. <i>Reproduction in Domestic Animals</i> , 2012 , 47 Suppl 6, 331-4	1.6	1
134	Attenuated vaccines can recombine to form virulent field viruses. <i>Science</i> , 2012 , 337, 188	33.3	126
133	Kinetics of transcription of infectious laryngotracheitis virus genes. <i>Comparative Immunology, Microbiology and Infectious Diseases</i> , 2012 , 35, 103-15	2.6	15
132	TonB is essential for virulence in avian pathogenic <i>Escherichia coli</i> . <i>Comparative Immunology, Microbiology and Infectious Diseases</i> , 2012 , 35, 129-38	2.6	16
131	Molecular epidemiology of <i>Mannheimia haemolytica</i> and <i>Mannheimia glucosida</i> associated with ovine mastitis. <i>Journal of Veterinary Diagnostic Investigation</i> , 2012 , 24, 730-4	1.5	6
130	Detection of a second novel gammaherpesvirus in a free-ranging koala (<i>Phascolarctos cinereus</i>). <i>Journal of Wildlife Diseases</i> , 2012 , 48, 226-9	1.3	11
129	Development and immunogenicity of recombinant GapA(+) <i>Mycoplasma gallisepticum</i> vaccine strain ts-11 expressing infectious bronchitis virus-S1 glycoprotein and chicken interleukin-6. <i>Vaccine</i> , 2011 , 29, 3197-205	4.1	10
128	Horizontal transmission dynamics of a glycoprotein G deficient candidate vaccine strain of infectious laryngotracheitis virus and the effect of vaccination on transmission of virulent virus. <i>Vaccine</i> , 2011 , 29, 5699-704	4.1	20
127	Comparative analysis of the complete genome sequences of two Australian origin live attenuated vaccines of infectious laryngotracheitis virus. <i>Vaccine</i> , 2011 , 29, 9583-7	4.1	27

126	The role of Mannheimia species in ovine mastitis. <i>Veterinary Microbiology</i> , 2011 , 153, 67-72	3.3	23
125	The central role of lipoproteins in the pathogenesis of mycoplasmoses. <i>Veterinary Microbiology</i> , 2011 , 153, 44-50	3.3	60
124	Infectious bronchitis viruses with naturally occurring genomic rearrangement and gene deletion. <i>Archives of Virology</i> , 2011 , 156, 245-52	2.6	18
123	First complete genome sequence of infectious laryngotracheitis virus. <i>BMC Genomics</i> , 2011 , 12, 197	4.5	38
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