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List of Publications by Year in descending order

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282 7,777 46
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291 291 all docs citations

291 times ranked 4740 citing authors

#	Article	IF	CITATIONS
1	The complete genome sequence of the avian pathogen Mycoplasma gallisepticum strain Rlow. Microbiology (United Kingdom), 2003, 149, 2307-2316.	0.7	204
2	Rotavirus serotypes 6 and 10 predominate in cattle. Journal of Clinical Microbiology, 1990, 28, 504-507.	1.8	166
3	Attenuated Vaccines Can Recombine to Form Virulent Field Viruses. Science, 2012, 337, 188-188.	6.0	154
4	The prevalence of enteric pathogens in diarrhoeic Thoroughbred foals in Britain and Ireland. Equine Veterinary Journal, 1991, 23, 405-409.	0.9	117
5	A novel group A rotavirus G serotype: serological and genomic characterization of equine isolate FI23. Journal of Clinical Microbiology, 1991, 29, 2043-2046.	1.8	116
6	A novel mechanism for control of antigenic variation in the haemagglutinin gene family of Mycoplasma synoviae. Molecular Microbiology, 2000, 35, 911-923.	1.2	113
7	Serological and genomic characterization of L338, a novel equine group A rotavirus G serotype. Journal of General Virology, 1991, 72, 1059-1064.	1.3	112
8	Chicken Anemia Virus VP2 Is a Novel Dual Specificity Protein Phosphatase. Journal of Biological Chemistry, 2002, 277, 39566-39573.	1.6	102
9	Multiplex Polymerase Chain Reaction Assay for Simultaneous Detection of Staphylococcus aureus and Streptococcal Causes of Bovine Mastitis. Journal of Dairy Science, 2001, 84, 1140-1148.	1.4	95
10	Molecular Epidemiology of Streptococcus uberis Isolates from Dairy Cows with Mastitis. Journal of Clinical Microbiology, 2001, 39, 1460-1466.	1.8	95
11	Variation between Pathogenic Serovars within Salmonella Pathogenicity Islands. Journal of Bacteriology, 2003, 185, 3624-3635.	1.0	95
12	Prevalence of Feline Chlamydia psittaci and Feline Herpesvirus 1 in Cats with Upper Respiratory Tract Disease. Journal of Veterinary Internal Medicine, 1999, 13, 153-162.	0.6	93
13	A type-specific serological test to distinguish antibodies to equine herpesviruses 4 and 1. Archives of Virology, 1995, 140, 245-258.	0.9	92
14	The central role of lipoproteins in the pathogenesis of mycoplasmoses. Veterinary Microbiology, 2011, 153, 44-50.	0.8	86
15	Mycoplasma synoviae has two distinct phase-variable major membrane antigens, one of which is a putative hemagglutinin. Infection and Immunity, 1997, 65, 2542-2547.	1.0	84
16	Lymphocytic infiltration in the chicken trachea in response to Mycoplasma gallisepticum infection. Microbiology (United Kingdom), 2000, 146, 1223-1229.	0.7	83
17	The organisation of the multigene family which encodes the major cell surface protein, pMGA, ofMycoplasma gallisepticum. FEBS Letters, 1994, 352, 347-352.	1.3	82
18	Development of a Recombinant Protein-Based Enzyme-Linked Immunosorbent Assay for Diagnosis of Mycoplasma bovis Infection in Cattle. Vaccine Journal, 2014, 21, 196-202.	3.2	77

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19	Barriers to and enablers of implementing antimicrobial stewardship programs in veterinary practices. Journal of Veterinary Internal Medicine, 2018, 32, 1092-1099.	0.6	77
20	Uropathogenic virulence factors in isolates of Escherichia coli from clinical cases of canine pyometra and feces of healthy bitches. Veterinary Microbiology, 2003, 94, 57-69.	0.8	75
21	Genomic Heterogeneity of Equine Betaherpesviruses. Journal of General Virology, 1987, 68, 1441-1447.	1.3	74
22	Multigene Families Encoding the Major Hemagglutinins in Phylogenetically Distinct Mycoplasmas. Infection and Immunity, 1998, 66, 3470-3475.	1.0	73
23	Transcription in Mycoplasma pneumoniae. Nucleic Acids Research, 2000, 28, 4488-4496.	6.5	70
24	The cellular immune response in the tracheal mucosa to Mycoplasma gallisepticum in vaccinated and unvaccinated chickens in the acute and chronic stages of disease. Vaccine, 2006, 24, 2627-2633.	1.7	69
25	Detection of feline calicivirus, feline herpesvirus 1 and Chlamydia psittaci mucosal swabs by multiplex RT-PCR/PCR. Veterinary Microbiology, 2001, 81, 95-108.	0.8	67
26	pMGA Phenotypic Variation in Mycoplasma gallisepticum Occurs In Vivo and Is Mediated by Trinucleotide Repeat Length Variation. Infection and Immunity, 2000, 68, 6027-6033.	1.0	65
27	Equine rhinovirus 1 is more closely related to foot-and-mouth disease virus than to other picornaviruses Proceedings of the National Academy of Sciences of the United States of America, 1996, 93, 990-995.	3.3	63
28	Glycoprotein G is a virulence factor in infectious laryngotracheitis virus. Journal of General Virology, 2006, 87, 2839-2847.	1.3	63
29	Expression of the pMGA Genes of <i>Mycoplasma gallisepticum</i> Is Controlled by Variation in the GAA Trinucleotide Repeat Lengths within the 5′ Noncoding Regions. Infection and Immunity, 1998, 66, 5833-5841.	1.0	63
30	Human and bovine serotype G 8 rotaviruses may be derived by reassortment. Archives of Virology, 1992, 125, 121-128.	0.9	61
31	Associations between the Ecology of Virulent Rhodococcus equi and the Epidemiology of R. equi Pneumonia on Australian Thoroughbred Farms. Applied and Environmental Microbiology, 2006, 72, 6152-6160.	1.4	61
32	Expression of Two Members of the pMGA Gene Family of <i>Mycoplasma gallisepticum</i> Oscillates and Is Influenced by pMGA-Specific Antibodies. Infection and Immunity, 1998, 66, 2845-2853.	1.0	61
33	Association of iss and iucA, but Not tsh, with Plasmid-Mediated Virulence of Avian Pathogenic Escherichia coli. Infection and Immunity, 2004, 72, 6554-6560.	1.0	60
34	Size and genomic location of the pMGA multigene family of Mycoplasma gallisepticum. Microbiology (United Kingdom), 1996, 142, 1429-1435.	0.7	59
35	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
36	Molecular epidemiology of Salmonella Heidelberg in an equine hospital. Veterinary Microbiology, 2001, 80, 85-98.	0.8	55

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37	Mycoplasma hyopneumoniae mhp379 Is a Ca 2+ -Dependent, Sugar-Nonspecific Exonuclease Exposed on the Cell Surface. Journal of Bacteriology, 2007, 189, 3414-3424.	1.0	55
38	Restriction Fragment Length Polymorphisms of Virulence Plasmids in <i>Rhodococcus equi</i> Journal of Clinical Microbiology, 1999, 37, 3417-3420.	1.8	55
39	Asinine herpesvirus genomes: comparison with those of the equine herpesviruses. Archives of Virology, 1988, 101, 183-190.	0.9	54
40	Expression studies on four members of the pMGA multigene family in Mycoplasma gallisepticum 56. Microbiology (United Kingdom), 1995, 141, 3005-3014.	0.7	54
41	Rapid PCR detection of Salmonella in horse faecal samples. Veterinary Microbiology, 2001, 79, 63-74.	0.8	52
42	Whole genome sequence analysis of Australian avian pathogenic Escherichia coli that carry the class 1 integrase gene. Microbial Genomics, $2019,5,\ldots$	1.0	51
43	Challenges and recent advancements in infectious laryngotracheitis virus vaccines. Avian Pathology, 2013, 42, 195-205.	0.8	50
44	A high prevalence of beak and feather disease virus in non-psittacine Australian birds. Journal of Medical Microbiology, 2017, 66, 1005-1013.	0.7	50
45	Examination ofmycoplasma gallisepticumstrains using restriction endonuclease DNA analysis and DNAâ€DNA hybridisation. Avian Pathology, 1988, 17, 559-570.	0.8	49
46	The major membrane nuclease MnuA degrades neutrophil extracellular traps induced by Mycoplasma bovis. Veterinary Microbiology, 2018, 218, 13-19.	0.8	49
47	Identification of four VP4 serological types (P serotypes) of bovine rotavirus using viral reassortants. Journal of General Virology, 1992, 73, 2319-2325.	1.3	49
48	Prevalence of feline Chlamydia psittaci and feline herpesvirus 1 in cats with upper respiratory tract disease. Journal of Veterinary Internal Medicine, 1999, 13, 153-62.	0.6	49
49	Strain differentiation of isolates of streptococci from bovine mastitis by pulsed-field gel electrophoresis. Molecular and Cellular Probes, 1997, 11, 349-354.	0.9	47
50	Detection and strain differentiation of feline calicivirus in conjunctival swabs by RT-PCR of the hypervariable region of the capsid protein gene. Archives of Virology, 1998, 143, 1321-1334.	0.9	46
51	Multiplex polymerase chain reaction as a mastitis screening test for Staphylococcus aureus, Streptococcus agalactiae, Streptococcus dysgalactiae and Streptococcus uberis in bulk milk samples. Journal of Dairy Research, 2003, 70, 149-155.	0.7	46
52	Evaluation of immunological responses to a glycoprotein G deficient candidate vaccine strain of infectious laryngotracheitis virus. Vaccine, 2010, 28, 1325-1332.	1.7	45
53	Natural recombination in alphaherpesviruses: Insights into viral evolution through full genome sequencing and sequence analysis. Infection, Genetics and Evolution, 2017, 49, 174-185.	1.0	45
54	Infectious Bronchitis Viruses with a Novel Genomic Organization. Journal of Virology, 2008, 82, 2013-2024.	1.5	44

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55	Antimicrobial Prescribing in Dogs and Cats in Australia: Results of the Australasian Infectious Disease Advisory Panel Survey. Journal of Veterinary Internal Medicine, 2017, 31, 1100-1107.	0.6	44
56	Epidemiology of equine herpesvirus 2 (equine cytomegalovirus). Journal of Clinical Microbiology, 1987, 25, 13-16.	1.8	44
57	Colonization of the Respiratory Tract by a Virulent Strain of Avian Escherichia coli Requires Carriage of a Conjugative Plasmid. Infection and Immunity, 2000, 68, 1535-1541.	1.0	43
58	Development of a SYBR Green quantitative polymerase chain reaction assay for rapid detection and quantification of infectious laryngotracheitis virus. Avian Pathology, 2011, 40, 237-242.	0.8	43
59	Epidemiology of Rhodococcus equi Strains on Thoroughbred Horse Farms. Applied and Environmental Microbiology, 2001, 67, 2167-2175.	1.4	42
60	First complete genome sequence of infectious laryngotracheitis virus. BMC Genomics, 2011, 12, 197.	1.2	42
61	Comparison of the Polymerase Chain Reaction and Culture for the Detection of Feline <i>Chlamydia psittaci</i> in Untreated and Doxycyclineâ€Treated Experimentally Infected Cats. Journal of Veterinary Internal Medicine, 1999, 13, 146-152.	0.6	40
62	GenotypingCryptosporidium parvum by single-strand conformation polymorphism analysis of ribosomal and heat shock gene regions. Electrophoresis, 2001, 22, 433-437.	1.3	40
63	Population wide assessment of antimicrobial use in dogs and cats using a novel data source – A cohort study using pet insurance data. Veterinary Microbiology, 2018, 225, 34-39.	0.8	40
64	Prevalence of G and P serotypes among equine rotaviruses in the faeces of diarrhoeic foals. Archives of Virology, 1996, 141, 1077-1089.	0.9	38
65	Comparison of concentrations of Rhodococcus equiand virulent R. equi in air of stables and paddocks on horse breeding farms in a temperate climate. Equine Veterinary Journal, 2010, 38, 263-265.	0.9	38
66	Veterinary Students' Knowledge and Perceptions About Antimicrobial Stewardship and Biosecurityâ€"A National Survey. Antibiotics, 2018, 7, 34.	1.5	38
67	Comparison of <i>Chlamydia psittaci</i> from cats with upper respiratory tract disease by polymerase chain reaction analysis of the <i>ompA</i> gene. Veterinary Record, 1997, 140, 310-313.	0.2	36
68	Prevalence of the virulence-associated gene of Rhodococcus equi in isolates from infected foals. Journal of Clinical Microbiology, 1997, 35, 1642-1644.	1.8	36
69	Latency of equine herpesvirus 4 (equine rhinopneumonitis virus). Veterinary Record, 1988, 123, 518-519.	0.2	36
70	Differential sensitivity of culture and the polymerase chain reaction for detection of feline herpesvirus 1 in vaccinated and unvaccinated cats. Archives of Virology, 1997, 142, 65-74.	0.9	35
71	A glycoprotein I- and glycoprotein E-deficient mutant of infectious laryngotracheitis virus exhibits impaired cell-to-cell spread in cultured cells. Archives of Virology, 2006, 151, 1281-1289.	0.9	35
72	Glycoprotein G deficient infectious laryngotracheitis virus is a candidate attenuated vaccine. Vaccine, 2007, 25, 3561-3566.	1.7	35

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73	Detection of Virulent Rhodococcus equi in Exhaled Air Samples from Naturally Infected Foals. Journal of Clinical Microbiology, 2009, 47, 734-737.	1.8	35
74	Comparative Metabolomics of Mycoplasma bovis and Mycoplasma gallisepticum Reveals Fundamental Differences in Active Metabolic Pathways and Suggests Novel Gene Annotations. MSystems, 2017, 2, .	1.7	35
75	Rotavirus serotype G3 predominates in horses. Journal of Clinical Microbiology, 1992, 30, 59-62.	1.8	35
76	Embryonic age influences the capacity for cytokine induction in chicken thymocytes. Immunology, 2003, 110, 358-367.	2.0	34
77	Equine rotavirusesâ€"Current understanding and continuing challenges. Veterinary Microbiology, 2013, 167, 135-144.	0.8	34
78	Disruption of the Membrane Nuclease Gene (MBOVPG45_0215) of Mycoplasma bovis Greatly Reduces Cellular Nuclease Activity. Journal of Bacteriology, 2015, 197, 1549-1558.	1.0	34
79	Evidence for two serotype G3 subtypes among equine rotaviruses. Journal of Clinical Microbiology, 1992, 30, 485-491.	1.8	34
80	A gene family in Mycoplasma imitans closely related to the pMGA family of Mycoplasma gallisepticum. Microbiology (United Kingdom), 1999, 145, 2095-2103.	0.7	33
81	Development of a replicable oriC plasmid for Mycoplasma gallisepticum and Mycoplasma imitans, and gene disruption through homologous recombination in M. gallisepticum. Microbiology (United) Tj $ETQq1\ 1\ G$).7843 ₫ ærgBT	/Ossærlock 10
82	Comparison of the safety and protective efficacy of vaccination with glycoprotein-G-deficient infectious laryngotracheitis virus delivered via eye-drop, drinking water or aerosol. Avian Pathology, 2008, 37, 83-88.	0.8	33
83	Evaluation of an IgG Enzyme-Linked Immunosorbent Assay as a Serological Assay for Detection of Mycoplasma bovis Infection in Feedlot Cattle. Journal of Clinical Microbiology, 2016, 54, 1269-1275.	1.8	33
84	A model for cystic endometrial hyperplasia/pyometra complex in the bitch. Theriogenology, 2006, 66, 1530-1536.	0.9	32
85	Genes Found Essential in Other Mycoplasmas Are Dispensable in Mycoplasma bovis. PLoS ONE, 2014, 9, e97100.	1.1	32
86	Recommended rejection of the names Malacoplasma gen. nov., Mesomycoplasma gen. nov., Metamycoplasma gen. nov., Metamycoplasmataceae fam. nov., Mycoplasmoidaceae fam. nov., Mycoplasmoidales ord. nov., Mycoplasmoides gen. nov., Mycoplasmopsis gen. nov. [Gupta, Sawnani, Adeolu, Alnajar and Oren 2018] and all proposed species comb. nov. placed therein. International	0.8	32
87	Journal of Systematic and Evolutionary Microbiology, 2019, 69, 3650-3653. Comparison of the Polymerase Chain Reaction and Culture for the Detection of Feline Chlamydia psittaci in Untreated and Doxycycline-Treated Experimentally Infected Cats. Journal of Veterinary Internal Medicine, 1999, 13, 146.	0.6	32
88	Site-directed mutagenesis of the VP2 gene of Chicken anemia virus affects virus replication, cytopathology and host-cell MHC class I expression. Journal of General Virology, 2006, 87, 823-831.	1.3	31
89	The immunoreactive 116 kDa surface protein of Mycoplasma pneumoniae is encoded in an operon. Microbiology (United Kingdom), 1997, 143, 3391-3402.	0.7	30
90	Genetic Mechanisms of Surface Variation. , 2002, , 417-443.		30

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91	Mycoplasma hyopneumoniae p65 Surface Lipoprotein Is a Lipolytic Enzyme with a Preference for Shorter-Chain Fatty Acids. Journal of Bacteriology, 2004, 186, 5790-5798.	1.0	30
92	Age related differences in the immune response to vaccination and infection with Mycoplasma gallisepticum. Vaccine, 2006, 24, 1687-1692.	1.7	30
93	Comparative analysis of the complete genome sequences of two Australian origin live attenuated vaccines of infectious laryngotracheitis virus. Vaccine, 2011, 29, 9583-9587.	1.7	30
94	Phylogenetic and Molecular Epidemiological Studies Reveal Evidence of Multiple Past Recombination Events between Infectious Laryngotracheitis Viruses. PLoS ONE, 2013, 8, e55121.	1.1	30
95	Characterisation of the course of Mycoplasma bovis infection in naturally infected dairy herds. Veterinary Microbiology, 2019, 231, 107-115.	0.8	29
96	Physical mapping of a genome of equine herpesvirus 2 (equine cytomegalovirus). Archives of Virology, 1989, 104, 77-86.	0.9	28
97	The conserved portion of the putative virulence region contributes to virulence of avian pathogenic Escherichia coli. Microbiology (United Kingdom), 2009, 155, 450-460.	0.7	28
98	Evidence of widespread natural recombination among field isolates of equine herpesvirus 4 but not among field isolates of equine herpesvirus 1. Journal of General Virology, 2016, 97, 747-755.	1.3	28
99	Physical mapping of the genomic heterogeneity of isolates of equine herpesvirus 2 (equine) Tj ETQq1 1 0.784314	4 rgBT /Ov	erlock 10 Tf
100	Development and application of an aerosol challenge method for reproduction of avian colibacillosis. Avian Pathology, 1998, 27, 505-511.	0.8	27
101	Outbreak of equine endometritis caused by a genotypically identical strain of <i>Pseudomonas aeruginosa</i> . Journal of Veterinary Diagnostic Investigation, 2011, 23, 1236-1239.	0.5	27
102	Genetic characterization of Cryptosporidium parvum from calves by mutation scanning and targeted sequencing – zoonotic implications. Electrophoresis, 2009, 30, 2640-2647.	1.3	26
103	The role of Mannheimia species in ovine mastitis. Veterinary Microbiology, 2011, 153, 67-72.	0.8	26
104	Comparative <i>in vivo</i> safety and efficacy of a glycoprotein G-deficient candidate vaccine strain of infectious laryngotracheitis virus delivered via eye drop. Avian Pathology, 2011, 40, 411-417.	0.8	26
105	The Mycoplasma gallisepticum Virulence Factor Lipoprotein MslA Is a Novel Polynucleotide Binding Protein. Infection and Immunity, 2013, 81, 3220-3226.	1.0	26
106	MalF is essential for persistence of Mycoplasma gallisepticum in vivo. Microbiology (United Kingdom), 2013, 159, 1459-1470.	0.7	26
107	Full-length infectious clone of a pathogenic Australian isolate of chicken anaemia virus. Australian Veterinary Journal, 2000, 78, 637-640.	0.5	25
108	Differential expression of lipoprotein genes in Mycoplasma pneumoniae after contact with human lung epithelial cells, and under oxidative and acidic stress. BMC Microbiology, 2008, 8, 124.	1.3	25

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109	Full genome analysis of Australian infectious bronchitis viruses suggests frequent recombination events between vaccine strains and multiple phylogenetically distant avian coronaviruses of unknown origin. Veterinary Microbiology, 2016, 197, 27-38.	0.8	25
110	Antimicrobials used for surgical prophylaxis by companion animal veterinarians in Australia. Veterinary Microbiology, 2017, 203, 301-307.	0.8	25
111	Reproduction of respiratory mycoplasmosis in calves by exposure to an aerosolised culture of Mycoplasma bovis. Veterinary Microbiology, 2017, 210, 167-173.	0.8	25
112	Plasmid-Borne Virulence-Associated Genes Have a Conserved Organization in Virulent Strains of Avian Pathogenic <i>Escherichia coli</i> Journal of Clinical Microbiology, 2009, 47, 2513-2519.	1.8	24
113	GapA+ Mycoplasma gallisepticum ts- 11 has improved vaccine characteristics. Microbiology (United) Tj ETQq 1	1 0.784314 i	rgBT /Overlo
114	Growth Kinetics and Transmission Potential of Existing and Emerging Field Strains of Infectious Laryngotracheitis Virus. PLoS ONE, 2015, 10, e0120282.	1.1	24
115	Uropathogenic virulence factor FimH facilitates binding of uteropathogenic Escherichia coli to canine endometrium. Comparative Immunology, Microbiology and Infectious Diseases, 2012, 35, 461-467.	0.7	23
116	Kinetics of transcription of infectious laryngotracheitis virus genes. Comparative Immunology, Microbiology and Infectious Diseases, 2012, 35, 103-115.	0.7	23
117	Antimicrobials used for surgical prophylaxis by equine veterinary practitioners in Australia. Equine Veterinary Journal, 2018, 50, 65-72.	0.9	23
118	Antimicrobial labelling in Australia: a threat to antimicrobial stewardship?. Australian Veterinary Journal, 2018, 96, 151-154.	0.5	23
119	Contagious Bovine and Caprine Pleuropneumonia: a research community's recommendations for the development of better vaccines. Npj Vaccines, 2020, 5, 66.	2.9	23
120	Identification and differentiation of avirulent and virulent Rhodococcus equi using selective media and colony blotting DNA hybridization to determine their concentrations in the environment. Veterinary Microbiology, 2004, 100, 121-127.	0.8	22
121	The vlhA loci of Mycoplasma synoviae are confined to a restricted region of the genome. Microbiology (United Kingdom), 2005, 151, 935-940.	0.7	22
122	IFN-Î ³ Enhances Immune Responses to E. coliInfection in the Chicken. Journal of Interferon and Cytokine Research, 2007, 27, 937-946.	0.5	22
123	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. Vaccine, 2011, 29, 5699-5704.	1.7	22
124	Detection of a Novel Gammaherpesvirus in Koalas (Phascolarctos cinereus). Journal of Wildlife Diseases, 2011, 47, 787-791.	0.3	22
125	Deep sequencing of the uterine immune response to bacteria during the equine oestrous cycle. BMC Genomics, 2015, 16, 934.	1.2	22
126	Genomic comparisons of Escherichia coli ST131 from Australia. Microbial Genomics, 2021, 7, .	1.0	22

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127	Mutation of chicken anemia virus VP2 differentially affects serine/threonine and tyrosine protein phosphatase activities. Journal of General Virology, 2005, 86, 623-630.	1.3	21
128	Rapid differentiation of current infectious bronchitis virus vaccine strains and field isolates in Australia. Australian Veterinary Journal, 2006, 84, 59-62.	0.5	21
129	Viral load in 1-day-old and 6-week-old chickens infected with chicken anaemia virus by the intraocular route. Avian Pathology, 2006, 35, 471-474.	0.8	21
130	Membrane proteins of Mycoplasma bovis and their role in pathogenesis. Research in Veterinary Science, 2013, 95, 321-325.	0.9	21
131	Two subspecies of bent-winged bats (Miniopterus orianae bassanii and oceanensis) in southern Australia have diverse fungal skin flora but not Pseudogymnoascus destructans. PLoS ONE, 2018, 13, e0204282.	1.1	21
132	Antimicrobial prescribing guidelines for poultry. Australian Veterinary Journal, 2021, 99, 181-235.	0.5	21
133	Indirect Enzyme-Linked Immunosorbent Assay for Detection of Immunoglobulin G Reactive with a Recombinant Protein Expressed from the Gene Encoding the 116-Kilodalton Protein of <i>Mycoplasma pneumoniae</i>). Journal of Clinical Microbiology, 1999, 37, 1024-1029.	1.8	21
134	Mycoplasma synoviae surface protein MSPB as a recombinant antigen in an indirect ELISA. Microbiology (United Kingdom), 1999, 145, 2087-2094.	0.7	20
135	Is <i>Chlamydophila felis</i> a significant zoonotic pathogen?. Australian Veterinary Journal, 2004, 82, 695-696.	0.5	20
136	Naturally occurring recombination between distant strains of infectious bronchitis virus. Archives of Virology, 2010, 155, 1581-1586.	0.9	20
137	<i>Mannheimia</i> Species Associated with Ovine Mastitis. Journal of Clinical Microbiology, 2010, 48, 3419-3422.	1.8	20
138	Infectious bronchitis viruses with naturally occurring genomic rearrangement and gene deletion. Archives of Virology, 2011, 156, 245-252.	0.9	20
139	Low genetic diversity among historical and contemporary clinical isolates of felid herpesvirus 1. BMC Genomics, 2016, 17, 704.	1.2	20
140	Lipoprotein Multigene Families in Mycoplasma pneumoniae. Journal of Bacteriology, 2006, 188, 5393-5399.	1.0	19
141	TonB is essential for virulence in avian pathogenic Escherichia coli. Comparative Immunology, Microbiology and Infectious Diseases, 2012, 35, 129-138.	0.7	19
142	Equine herpesvirus genomes: Heterogeneity of naturally occurring type 4 isolates and of a type 1 isolate after heterologous cell passage. Archives of Virology, 1986, 91, 375-381.	0.9	18
143	Application of highâ€resolution melt curve analysis for classification of infectious bronchitis viruses in field specimens. Australian Veterinary Journal, 2010, 88, 408-413.	0.5	18
144	Detection of a Second Novel Gammaherpesvirus in a Free-ranging Koala (Phascolarctos cinereus). Journal of Wildlife Diseases, 2012, 48, 226-229.	0.3	18

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145	Review: Water medication of growing pigs: sources of between-animal variability in systemic exposure to antimicrobials. Animal, 2019, 13, 3031-3040.	1.3	18
146	$\mbox{\sc (i)}$ Salmonella $\mbox{\sc / i)}$ Genomic Island 1B Variant Found in a Sequence Type 117 Avian Pathogenic Escherichia coli Isolate. MSphere, 2019, 4, .	1.3	18
147	Typing infectious bronchitis virus strains using reverse transcription-polymerase chain reaction and restriction fragment length polymorphism analysis to compare the 3′ 7.5 kb of their genomes. Avian Pathology, 2006, 35, 63-69.	0.8	17
148	ISOLATION AND CHARACTERIZATION OF A NOVEL HERPESVIRUS FROM A FREE-RANGING EASTERN GREY KANGAROO (MACROPUS GIGANTEUS). Journal of Wildlife Diseases, 2013, 49, 143-151.	0.3	17
149	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
150	Development of a <i>Mycoplasma gallisepticum</i> infection model in turkeys. Avian Pathology, 2015, 44, 35-42.	0.8	17
151	A Novel Glaesserella sp. Isolated from Pigs with Severe Respiratory Infections Has a Mosaic Genome with Virulence Factors Putatively Acquired by Horizontal Transfer. Applied and Environmental Microbiology, 2018, 84, .	1.4	17
152	Targeted mutagenesis of Mycoplasma gallisepticum using its endogenous CRISPR/Cas system. Veterinary Microbiology, 2020, 250, 108868.	0.8	17
153	The Effect of an Alternate Start Codon on Heterologous Expression of a PhoA Fusion Protein in Mycoplasma gallisepticum. PLoS ONE, 2015, 10, e0127911.	1.1	16
154	Impacts of poultry vaccination on viruses of wild bird. Current Opinion in Virology, 2016, 19, 23-29.	2.6	16
155	Genetic Diversity of Infectious Laryngotracheitis Virus during In Vivo Coinfection Parallels Viral Replication and Arises from Recombination Hot Spots within the Genome. Applied and Environmental Microbiology, 2017, 83, .	1.4	16
156	Development and application of a TaqMan single nucleotide polymorphism genotyping assay to study infectious laryngotracheitis virus recombination in the natural host. PLoS ONE, 2017, 12, e0174590.	1.1	16
157	Detection of Antibodies to Mycoplasma gallisepticum Vaccine ts-11 by an Autologous pMGA Enzyme-Linked Immunosorbent Assay. Avian Diseases, 2002, 46, 405-411.	0.4	15
158	Development and immunogenicity of recombinant GapA+ Mycoplasma gallisepticum vaccine strain ts-11 expressing infectious bronchitis virus-S1 glycoprotein and chicken interleukin-6. Vaccine, 2011, 29, 3197-3205.	1.7	15
159	Oestrous cycle-dependent equine uterine immune response to induced infectious endometritis. Veterinary Research, 2016, 47, 110.	1.1	15
160	Exploration of antibiotic resistance risks in a veterinary teaching hospital with Oxford Nanopore long read sequencing. PLoS ONE, 2019, 14, e0217600.	1.1	15
161	A combined metabolomic and bioinformatic approach to investigate the function of transport proteins of the important pathogen Mycoplasma bovis. Veterinary Microbiology, 2019, 234, 8-16.	0.8	15
162	Homotypic and heterotypic serum and milk antibody to rotavirus in normal, infected and vaccinated horses. Veterinary Microbiology, 1991, 27, 231-244.	0.8	14

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163	IL-18 Expression in Pigs Following Infection with Mycoplasma hyopneumoniae. Journal of Interferon and Cytokine Research, 2006, 26, 637-644.	0.5	14
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