Joanne Meers

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Retroviral invasion of the koala genome. Nature, 2006, 442, 79-81.	13.7	322
2	Real-time reverse transcriptase PCR for the endogenous koala retrovirus reveals an association between plasma viral load and neoplastic disease in koalas. Journal of General Virology, 2005, 86, 783-787.	1.3	139
3	Prevalence of koala retrovirus in geographically diverse populations in Australia. Australian Veterinary Journal, 2012, 90, 404-409.	0.5	107
4	Bats Without Borders: Long-Distance Movements and Implications for Disease Risk Management. EcoHealth, 2010, 7, 204-212.	0.9	95
5	Endogenous retroviruses. Cellular and Molecular Life Sciences, 2008, 65, 3413-3421.	2.4	77
6	Avian paramyxoviruses and influenza viruses isolated from mallard ducks (Anas platyrhynchos) in New Zealand. Archives of Virology, 2002, 147, 1287-1302.	0.9	72
7	Farm- and flock-level risk factors associated with Highly Pathogenic Avian Influenza outbreaks on small holder duck and chicken farms in the Mekong Delta of Viet Nam. Preventive Veterinary Medicine, 2009, 91, 179-188.	0.7	63
8	Equine respiratory viruses in foals in New Zealand. New Zealand Veterinary Journal, 2002, 50, 140-147.	0.4	61
9	Molecular confirmation of an adenovirus in brushtail possums (Trichosurus vulpecula). Virus Research, 2002, 83, 189-195.	1.1	56
10	Survival of rabbit haemorrhagic disease virus (RHDV) in the environment. Epidemiology and Infection, 2005, 133, 719-730.	1.0	55
11	Genetic analysis of canine parvovirus from dogs in Australia. Australian Veterinary Journal, 2007, 85, 392-396.	0.5	54
12	Evidence of Endemic Hendra Virus Infection in Flying-Foxes (Pteropus conspicillatus)—Implications for Disease Risk Management. PLoS ONE, 2011, 6, e28816.	1.1	53
13	The Distribution of Henipaviruses in Southeast Asia and Australasia: Is Wallace's Line a Barrier to Nipah Virus?. PLoS ONE, 2013, 8, e61316.	1.1	48
14	Extensive sequence variation of feline immunodeficiency virusenv genes in isolates from naturally infected cats. Archives of Virology, 1993, 133, 51-62.	0.9	47
15	Viruses associated with outbreaks of equine respiratory disease in New Zealand. New Zealand Veterinary Journal, 2002, 50, 132-139.	0.4	47
16	Scavenging Ducks and Transmission of Highly Pathogenic Avian Influenza, Java, Indonesia. Emerging Infectious Diseases, 2010, 16, 1244-1250.	2.0	45
17	Highly pathogenic avian influenza (H5N1) in ducks and in-contact chickens in backyard and smallholder commercial duck farms in Viet Nam. Preventive Veterinary Medicine, 2011, 101, 229-240.	0.7	45
18	The essential and non-essential genes of Bovine herpesvirus 1. Journal of General Virology, 2008, 89, 2851-2863.	1.3	44

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19	Coronavirus Infection and Diversity in Bats in the Australasian Region. EcoHealth, 2016, 13, 72-82.	0.9	41
20	Phylogenetic Diversity of Koala Retrovirus within a Wild Koala Population. Journal of Virology, 2017, 91, .	1.5	40
21	Degradation and remobilization of endogenous retroviruses by recombination during the earliest stages of a germ-line invasion. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 8609-8614.	3.3	40
22	Circovirus-like infection in a southern black-backed gull (Larus dominicanus). Avian Pathology, 1999, 28, 513-516.	0.8	38
23	Characteristics of Nipah virus and Hendra virus replication in different cell lines and their suitability for antiviral screening. Virus Research, 2009, 142, 92-99.	1.1	38
24	Genetic typing of pestiviruses from New Zealand. New Zealand Veterinary Journal, 1998, 46, 35-37.	0.4	36
25	Evaluation of immune effects of fowlpox vaccine strains and field isolates. Veterinary Microbiology, 2006, 116, 106-119.	0.8	36
26	Discovery of a Novel Retrovirus Sequence in an Australian Native Rodent (Melomys burtoni): A Putative Link between Gibbon Ape Leukemia Virus and Koala Retrovirus. PLoS ONE, 2014, 9, e106954.	1.1	36
27	Isolation of equine herpesvirus type 5 in New Zealand. New Zealand Veterinary Journal, 1999, 47, 44-46.	0.4	34
28	Nucleotide sequences of Australian isolates of the feline immunodeficiency virus: comparison with other feline lentiviruses. Archives of Virology, 1993, 132, 369-379.	0.9	33
29	Acute phase proteins in healthy and sick cats. Research in Veterinary Science, 2012, 93, 649-654.	0.9	31
30	Molecular subtyping of feline immunodeficiency virus from domestic cats in Australia. Australian Veterinary Journal, 2006, 84, 112-116.	0.5	29
31	Village chicken production in Myanmar – purpose, magnitude and major constraints. World's Poultry Science Journal, 2007, 63, 308-322.	1.4	28
32	Genomic variability of equine herpesvirus-5. Archives of Virology, 2000, 145, 1359-1371.	0.9	27
33	Physiological stress and Hendra virus in flying-foxes (Pteropus spp.), Australia. PLoS ONE, 2017, 12, e0182171.	1.1	27
34	Evaluation of strategies to improve village chicken production-controlled field trials to assess effects of Newcastle disease vaccination and altered chick rearing in Myanmar. Preventive Veterinary Medicine, 2009, 90, 17-30.	0.7	26
35	The pathobiology of two Indonesian H5N1 avian influenza viruses representing different clade 2.1 sublineages in chickens and ducks. Comparative Immunology, Microbiology and Infectious Diseases, 2013, 36, 175-191.	0.7	26
36	Identification of stable reference genes for quantitative PCR in koalas. Scientific Reports, 2018, 8, 3364.	1.6	26

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37	Mortality rates adjusted for unobserved deaths and associations with Newcastle disease virus serology among unvaccinated village chickens in Myanmar. Preventive Veterinary Medicine, 2008, 85, 241-252.	0.7	25
38	Back to BAC: The Use of Infectious Clone Technologies for Viral Mutagenesis. Viruses, 2012, 4, 211-235.	1.5	24
39	Haematology and Plasma Biochemistry of Wild Black Flying-Foxes, (Pteropus alecto) in Queensland, Australia. PLoS ONE, 2015, 10, e0125741.	1.1	24
40	Characteristics of two duck farming systems in the Mekong Delta of Viet Nam: stationary flocks and moving flocks, and their potential relevance to the spread of highly pathogenic avian influenza. Tropical Animal Health and Production, 2013, 45, 837-848.	0.5	23
41	Hendra Virus Infection Dynamics in the Grey-Headed Flying Fox (Pteropus poliocephalus) at the Southern-Most Extent of Its Range: Further Evidence This Species Does Not Readily Transmit the Virus to Horses. PLoS ONE, 2016, 11, e0155252.	1.1	23
42	Koala retrovirus viral load and disease burden in distinct northern and southern koala populations. Scientific Reports, 2020, 10, 263.	1.6	22
43	Genome sequence of the thermostable Newcastle disease virus (strain I-2) reveals a possible phenotypic locus. Veterinary Microbiology, 2006, 114, 134-141.	0.8	21
44	A molecular and antigenic survey of H5N1 highly pathogenic avian influenza virus isolates from smallholder duck farms in Central Java, Indonesia during 2007-2008. Virology Journal, 2011, 8, 425.	1.4	21
45	Experimentally Infected Domestic Ducks Show Efficient Transmission of Indonesian H5N1 Highly Pathogenic Avian Influenza Virus, but Lack Persistent Viral Shedding. PLoS ONE, 2014, 9, e83417.	1.1	21
46	Feline immunodeficiency virus infection: plasma, but not peripheral blood mononuclear cell virus titer is influenced by zidovudine and cyclosporine. Archives of Virology, 1993, 132, 67-81.	0.9	20
47	Emergence of canine parvovirus subtype 2b (CPV-2b) infections in Australian dogs. Infection, Genetics and Evolution, 2018, 58, 50-55.	1.0	20
48	Analysis of canine parvoviruses circulating in Australia reveals predominance of variant 2b and identifies feline parvovirusâ€like mutations in the capsid proteins. Transboundary and Emerging Diseases, 2021, 68, 656-666.	1.3	20
49	Genetic diversity of Koala retrovirus env gene subtypes: insights into northern and southern koala populations. Journal of General Virology, 2019, 100, 1328-1339.	1.3	20
50	Feline immunodeficiency virus: quantification in peripheral blood mononuclear cells and isolation from plasma of infected cats. Archives of Virology, 1992, 127, 233-243.	0.9	19
51	Who Is Spreading Avian Influenza in the Moving Duck Flock Farming Network of Indonesia?. PLoS ONE, 2016, 11, e0152123.	1.1	18
52	Pathological Findings in Koala Retrovirus-positive Koalas (Phascolarctos cinereus) from Northern and Southern Australia. Journal of Comparative Pathology, 2020, 176, 50-66.	0.1	18
53	The detection and quantification of feline immunodeficiency provirus in peripheral blood mononuclear cells using the polymerase chain reaction. Veterinary Microbiology, 1993, 38, 11-21.	0.8	17
54	Validation of real-time polymerase chain reaction tests for diagnosing feline immunodeficiency virus infection in domestic cats using Bayesian latent class models. Preventive Veterinary Medicine, 2012, 104, 136-148.	0.7	17

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55	Association between feline immunodeficiency virus (FIV) plasma viral RNA load, concentration of acute phase proteins and disease severity. Veterinary Journal, 2014, 201, 181-183.	0.6	17
56	A new papillomavirus of possums (Trichosurus vulpecula) associated with typical wart-like papillomas. Archives of Virology, 2000, 145, 1247-1255.	0.9	16
57	A neurological syndrome in a free-living population of possums (Trichosurus vulpecula). New Zealand Veterinary Journal, 2000, 48, 9-15.	0.4	16
58	Husbandry and trade of indigenous chickens in Myanmar—Results of a participatory rural appraisal in the Yangon and the Mandalay divisions. Tropical Animal Health and Production, 2006, 38, 611-618.	0.5	16
59	A comparative evaluation of feathers, oropharyngeal swabs, and cloacal swabs for the detection of H5N1 highly pathogenic avian influenza virus infection in experimentally infected chickens and ducks. Journal of Veterinary Diagnostic Investigation, 2015, 27, 704-715.	0.5	16
60	Temporal Variation in Physiological Biomarkers in Black Flying-Foxes (Pteropus alecto), Australia. EcoHealth, 2016, 13, 49-59.	0.9	15
61	Co-infection with different subtypes of feline immunodeficiency virus can complicate subtype assignment by phylogenetic analysis. Archives of Virology, 2007, 152, 1187-1193.	0.9	14
62	Highly Pathogenic Avian Influenza (H5N1) Virus in Feathers. Veterinary Pathology, 2017, 54, 226-233.	0.8	14
63	Routes of transmission of wobbly possum disease. New Zealand Veterinary Journal, 2000, 48, 3-8.	0.4	13
64	Comparison of serological assays for detecting antibodies in ducks exposed to H5 subtype avian influenza virus. BMC Veterinary Research, 2012, 8, 117.	0.7	12
65	Economic analysis of interventions to improve village chicken production in Myanmar. Preventive Veterinary Medicine, 2013, 110, 525-540.	0.7	12
66	Assessing the risk of Nipah virus establishment in Australian flying-foxes. Epidemiology and Infection, 2015, 143, 2213-2226.	1.0	11
67	Canine parvovirus is shed infrequently by cats without diarrhoea in multi-cat environments. Veterinary Microbiology, 2021, 261, 109204.	0.8	11
68	Influence of equine herpesvirus type 2 infection on monocyte chemoattractant protein 1 gene transcription in equine blood mononuclear cells. Research in Veterinary Science, 2001, 71, 111-113.	0.9	10
69	Determination of Organ Tropism of Newcastle Disease Virus (Strain I-2) by Virus Isolation and Reverse Transcription–Polymerase Chain Reaction. Veterinary Research Communications, 2006, 30, 697-706.	0.6	10
70	Survival of avirulent thermostable Newcastle disease virus (strain I-2) in raw, baked, oiled, and cooked white rice at ambient temperatures. Journal of Veterinary Science, 2007, 8, 303.	0.5	10
71	Incidence and risk factors for H5 highly pathogenic avian influenza infection in flocks of apparently clinically healthy ducks. Epidemiology and Infection, 2013, 141, 390-401.	1.0	10
72	A novel Australian flying-fox retrovirus shares an evolutionary ancestor with Koala, Gibbon and Melomys gamma-retroviruses. Virus Genes, 2019, 55, 421-424.	0.7	10

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73	Quantification of lymphadenopathy in experimentally induced feline immunodeficiency virus infection in domestic cats. Veterinary Immunology and Immunopathology, 1995, 46, 3-12.	0.5	9
74	A survey for paramyxoviruses in caged birds, wild birds, and poultry in New Zealand. New Zealand Veterinary Journal, 2001, 49, 18-23.	0.4	9
75	Exposure of rabbits to ultraviolet light-inactivated rabbit haemorrhagic disease virus (RHDV) and subsequent challenge with virulent virus. Epidemiology and Infection, 2005, 133, 731-735.	1.0	9
76	Phylogenetic analysis to define feline immunodeficiency virus subtypes in 31 domestic cats in South Africa. Journal of the South African Veterinary Association, 2006, 77, 108-113.	0.2	8
77	Genomic deletions and mutations resulting in the loss of eight genes reduce the in vivo replication capacity of Meleagrid herpesvirus 1. Virus Genes, 2015, 51, 85-95.	0.7	8
78	A survey for torovirus in New Zealand cats with protruding nictitating membranes. New Zealand Veterinary Journal, 1997, 45, 41-43.	0.4	7
79	Temporal dynamics of rabbit haemorrhagic disease virus infection in a low-density population of wild rabbits (Oryctolagus cuniculus) in New Zealand. Wildlife Research, 2006, 33, 293.	0.7	7
80	Seropositivity to rabbit haemorrhagic disease virus in non-target mammals during periods of viral activity in a population of wild rabbits in New Zealand. Wildlife Research, 2006, 33, 305.	0.7	7
81	Prevalence and incidence of Newcastle disease and prevalence of Avian Influenza infection of scavenging village chickens in Timor-Lesté. Preventive Veterinary Medicine, 2012, 104, 301-308.	0.7	7
82	Genetic analysis of porcine circovirus type 2 (PCV2) in Queensland, Australia. Australian Veterinary Journal, 2020, 98, 388-395.	0.5	7
83	Pathogens associated with pleuritic pig lungs at an abattoir in Queensland Australia. Australian Veterinary Journal, 2021, 99, 163-171.	0.5	7
84	The Epidemiology of Koala Retrovirus. Journal of Veterinary Epidemiology, 2011, 15, 1-9.	0.2	7
85	Feline immunodeficiency virus subtypes in domestic cats in New Zealand. New Zealand Veterinary Journal, 2007, 55, 358-360.	0.4	6
86	Mortality patterns over 3Âyears in a sparse population of wild rabbits (Oryctolagus cuniculus) in New Zealand, with an emphasis on rabbit haemorrhagic disease (RHD). European Journal of Wildlife Research, 2008, 54, 619-626.	0.7	6
87	Interdisciplinary communication of infectious disease research – translating complex epidemiological findings into understandable messages for village chicken farmers in Myanmar. SpringerPlus, 2014, 3, 726.	1.2	6
88	Shelter-housed cats show no evidence of faecal shedding of canine parvovirus DNA. Veterinary Journal, 2018, 239, 54-58.	0.6	6
89	Transcriptomic and genomic variants between koala populations reveals underlying genetic components to disorders in a bottlenecked population. Conservation Genetics, 2021, 22, 329-340.	0.8	6
90	Novel insights into viral infection and oncogenesis from koala retrovirus (KoRV) infection of HEK293T cells. Gene, 2020, 733, 144366.	1.0	5

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91	An unprecedented cluster of Australian bat lyssavirus in <i>Pteropus conspicillatus</i> indicates preâ€flight flying fox pups are at risk of mass infection. Zoonoses and Public Health, 2020, 67, 435-442.	0.9	4
92	Koala retrovirus in free-ranging populations—prevalence. Technical Reports of the Australian Museum Online, 0, 24, 15-17.	0.0	4
93	Differential and defective transcription of koala retrovirus indicates the complexity of host and virus evolution. Journal of General Virology, 2022, 103, .	1.3	4
94	Thermostability profile of Newcastle disease virus (strain I–2) following serial passages without heat selection. Tropical Animal Health and Production, 2006, 38, 527-531.	0.5	3
95	Development of a Cell Culture Method for Quantal Assay of Strain I-2 of Newcastle Disease Virus. Veterinary Research Communications, 2006, 30, 689-696.	0.6	3
96	Deduced Amino Acid Sequences Surrounding the Fusion Glycoprotein Cleavage Site and of the Carboxyl-terminus of Haemagglutinin–Neuraminidase Protein of the Avirulent Thermostable Vaccine Strain I-2 of Newcastle disease virus. Veterinary Research Communications, 2007, 31, 105-112.	0.6	3
97	Training Veterinary Personnel for Effective Identification and Diagnosis of Exotic Animal Diseases. Journal of Veterinary Medical Education, 2008, 35, 255-261.	0.4	3
98	The management of smallholder duck flocks in Central Java, Indonesia, and potential hazards promoting the spread of Highly Pathogenic Avian Influenza virus. World's Poultry Science Journal, 2012, 68, 513-528.	1.4	3
99	Characterisation of New Zealand isolates of infectious bursal disease virus. Archives of Virology, 2001, 146, 1571-1580.	0.9	2
100	The Meleagrid herpesvirus 1 Genome Is Partially Resistant to Transposition. Avian Diseases, 2013, 57, 380-386.	0.4	2
101	Identification of non-essential loci within the Meleagrid herpesvirus 1 genome. Virology Journal, 2015, 12, 130.	1.4	2
102	Evaluation of serological, histological and immunocytochemical methods for the detection of infectious bursal disease virus infection in broiler flocks in New Zealand. New Zealand Veterinary Journal, 1999, 47, 175-179.	0.4	1
103	Development of a rapid biological assay for determination of potency of Newcastle disease vaccine (strain I-2). Tropical Animal Health and Production, 2006, 38, 463-466.	0.5	0
104	Transport of Moving Duck Flocks in Indonesia and Vietnam: Management Practices That Potentially Impact Avian Pathogen Dissemination. Frontiers in Veterinary Science, 2021, 8, 673624.	0.9	0