

List of Publications by Year in descending order

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123
papers

3,939
citations

94381

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all docs

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docs citations

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times ranked

2550
citing authors

#	ARTICLE	IF	CITATIONS
1	<i>Chlamydia pecorum</i> “Induced Arthritis in Experimentally and Naturally Infected Sheep. <i>Veterinary Pathology</i> , 2021, 58, 346-360.	0.8	7
2	The trends of human dirofilariasis in Croatia: Yesterday “ Today “ Tomorrow. <i>One Health</i> , 2020, 10, 100153.	1.5	10
3	Recent history of psittacosis in Australia: expanding our understanding of the epidemiology of this important globally distributed zoonotic disease. <i>Internal Medicine Journal</i> , 2020, 50, 246-249.	0.5	14
4	New insights into chlamydial zoonoses. <i>Microbiology Australia</i> , 2020, 41, 14.	0.1	5
5	Detection of <i>Chlamydiaceae</i> in ocular swabs from Australian pre-export feedlot sheep. <i>Australian Veterinary Journal</i> , 2019, 97, 401-403.	0.5	14
6	Population-scale treatment informs solutions for control of environmentally transmitted wildlife disease. <i>Journal of Applied Ecology</i> , 2019, 56, 2363-2375.	1.9	22
7	Chlamydial infection and on-farm risk factors in dairy cattle herds in South East Queensland. <i>Australian Veterinary Journal</i> , 2019, 97, 505-508.	0.5	8
8	The impact of human activities on Australian wildlife. <i>PLoS ONE</i> , 2019, 14, e0206958.	1.1	61
9	<i>Chlamydia pecorum</i> prevalence in South Australian koala (<i>Phascolarctos cinereus</i>) populations: Identification and modelling of a population free from infection. <i>Scientific Reports</i> , 2019, 9, 6261.	1.6	23
10	Detection of a range of genetically diverse chlamydiae in Australian domesticated and wild ungulates. <i>Transboundary and Emerging Diseases</i> , 2019, 66, 1132-1137.	1.3	16
11	In vitro analysis of genetically distinct <i>Chlamydia pecorum</i> isolates reveals key growth differences in mammalian epithelial and immune cells. <i>Veterinary Microbiology</i> , 2019, 232, 22-29.	0.8	10
12	Identification of A Novel Picorna-Like Virus, Burpengary Virus, that is Negatively Associated with Chlamydial Disease in the Koala. <i>Viruses</i> , 2019, 11, 211.	1.5	6
13	Mini Review: Antimicrobial Control of Chlamydial Infections in Animals: Current Practices and Issues. <i>Frontiers in Microbiology</i> , 2019, 10, 113.	1.5	33
14	Seroprevalence of vector-borne pathogens in dogs from Croatia. <i>Parasitology Research</i> , 2019, 118, 347-352.	0.6	11
15	New evidence for domesticated animals as reservoirs of <i>Chlamydia</i> -associated community-acquired pneumonia. <i>Clinical Microbiology and Infection</i> , 2019, 25, 131-132.	2.8	10
16	Multilocus Sequence Typing (MLST) of Chlamydiales. <i>Methods in Molecular Biology</i> , 2019, 2042, 69-86.	0.4	8
17	The limitations of commercial serological assays for detection of chlamydial infections in Australian livestock. <i>Journal of Medical Microbiology</i> , 2019, 68, 627-632.	0.7	11
18	Prevalence and clinical significance of koala retrovirus in two South Australian koala (<i>Phascolarctos cinereus</i>) populations. <i>Journal of Medical Microbiology</i> , 2019, 68, 1072-1080.	0.7	20

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19	Koala immunology and infectious diseases: How much can the koala bear?. <i>Developmental and Comparative Immunology</i> , 2018, 82, 177-185.	1.0	12
20	Understanding the health and production impacts of endemic <i>Chlamydia pecorum</i> infections in lambs. <i>Veterinary Microbiology</i> , 2018, 217, 90-96.	0.8	16
21	Molecular evidence to suggest pigeon-type <i>Chlamydia psittaci</i> in association with an equine foal loss. <i>Transboundary and Emerging Diseases</i> , 2018, 65, 911-915.	1.3	22
22	Comparative diagnostics reveals PCR assays on skin scrapings is the most reliable method to detect <i>Sarcoptes scabiei</i> infestations. <i>Veterinary Parasitology</i> , 2018, 251, 119-124.	0.7	23
23	Characterization of the In Vitro <i>Chlamydia pecorum</i> Response to Gamma Interferon. <i>Infection and Immunity</i> , 2018, 86, .	1.0	11
24	A Review on Chlamydial Diseases in Animals: Still a Challenge for Pathologists?. <i>Veterinary Pathology</i> , 2018, 55, 374-390.	0.8	112
25	Transcriptome sequencing of the long-nosed bandicoot (<i>Perameles nasuta</i>) reveals conservation and innovation of immune genes in the marsupial order Peramelemorphia. <i>Immunogenetics</i> , 2018, 70, 327-336.	1.2	3
26	Characterisation of MHC class I genes in the koala. <i>Immunogenetics</i> , 2018, 70, 125-133.	1.2	15
27	<i>Chlamydia pecorum</i> gastrointestinal tract infection associations with urogenital tract infections in the koala (<i>Phascolarctos cinereus</i>). <i>PLoS ONE</i> , 2018, 13, e0206471.	1.1	18
28	Cloacal and Ocular Microbiota of the Endangered Australian Northern Quoll. <i>Microorganisms</i> , 2018, 6, 68.	1.6	5
29	Metagenomic Analysis of Fish-Associated <i>Ca. Parilichlamydiaceae</i> Reveals Striking Metabolic Similarities to the Terrestrial <i>Chlamydiaceae</i> . <i>Genome Biology and Evolution</i> , 2018, 10, 2587-2595.	1.1	5
30	Epitheliocystis in fish: An emerging aquaculture disease with a global impact. <i>Transboundary and Emerging Diseases</i> , 2018, 65, 1436-1446.	1.3	52
31	Adaptation and conservation insights from the koala genome. <i>Nature Genetics</i> , 2018, 50, 1102-1111.	9.4	163
32	An epizootic of <i>Chlamydia psittaci</i> equine reproductive loss associated with suspected spillover from native Australian parrots. <i>Emerging Microbes and Infections</i> , 2018, 7, 1-13.	3.0	48
33	<i>Chlamydia pecorum</i> in Joint Tissue and Synovial Fluid of a Koala (<i>Phascolarctos cinereus</i>) with Arthritis. <i>Journal of Wildlife Diseases</i> , 2018, 54, 646-649.	0.3	3
34	A <i>Sarcoptes scabiei</i> specific isothermal amplification assay for detection of this important ectoparasite of wombats and other animals. <i>PeerJ</i> , 2018, 6, e5291.	0.9	17
35	Culture-independent approaches to chlamydial genomics. <i>Microbial Genomics</i> , 2018, 4, .	1.0	22
36	Molecular and serological dynamics of <i>Chlamydia pecorum</i> infection in a longitudinal study of prime lamb production. <i>PeerJ</i> , 2018, 6, e4296.	0.9	13

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37	Characterization of shifts of koala (<i>Phascolarctos cinereus</i>) intestinal microbial communities associated with antibiotic treatment. <i>PeerJ</i> , 2018, 6, e4452.	0.9	30
38	Culture-independent genomics of a novel chlamydial pathogen of fish provides new insight into host-specific adaptations utilized by these intracellular bacteria. <i>Environmental Microbiology</i> , 2017, 19, 1899-1913.	1.8	31
39	Multilocus sequence typing identifies an avian-like <i>Chlamydia psittaci</i> strain involved in equine placentitis and associated with subsequent human psittacosis. <i>Emerging Microbes and Infections</i> , 2017, 6, 1-3.	3.0	39
40	A new equine and zoonotic threat emerges from an old avian pathogen, <i>Chlamydia psittaci</i> . <i>Clinical Microbiology and Infection</i> , 2017, 23, 693-694.	2.8	26
41	Safety and immunogenicity of a prototype anti- <i>Chlamydia pecorum</i> recombinant protein vaccine in lambs and pregnant ewes. <i>Vaccine</i> , 2017, 35, 3461-3465.	1.7	10
42	Molecular evidence of <i>Chlamydia pecorum</i> and arthropod-associated Chlamydiae in an expanded range of marsupials. <i>Scientific Reports</i> , 2017, 7, 12844.	1.6	8
43	Molecular detection of <i>Anaplasma platys</i> , <i>Anaplasma phagocytophilum</i> and <i>Wolbachia</i> sp. but not <i>Ehrlichia canis</i> in Croatian dogs. <i>Parasitology Research</i> , 2017, 116, 3019-3026.	0.6	23
44	Culture-independent metagenomics supports discovery of uncultivable bacteria within the genus <i>Chlamydia</i> . <i>Scientific Reports</i> , 2017, 7, 10661.	1.6	49
45	New and emerging chlamydial infections of creatures great and small. <i>New Microbes and New Infections</i> , 2017, 18, 28-33.	0.8	40
46	Novel Chlamydiales genotypes identified in ticks from Australian wildlife. <i>Parasites and Vectors</i> , 2017, 10, 46.	1.0	21
47	Immunization of a wild koala population with a recombinant <i>Chlamydia pecorum</i> Major Outer Membrane Protein (MOMP) or Polymorphic Membrane Protein (PMP) based vaccine: New insights into immune response, protection and clearance. <i>PLoS ONE</i> , 2017, 12, e0178786.	1.1	24
48	Epidemiology of chlamydial infection and disease in a free-ranging koala (<i>Phascolarctos cinereus</i>) population. <i>PLoS ONE</i> , 2017, 12, e0190114.	1.1	39
49	Mitochondrial genome sequencing reveals potential origins of the scabies mite <i>Sarcoptes scabiei</i> infesting two iconic Australian marsupials. <i>BMC Evolutionary Biology</i> , 2017, 17, 233.	3.2	22
50	Asymptomatic infections with highly polymorphic <i>Chlamydia suis</i> are ubiquitous in pigs. <i>BMC Veterinary Research</i> , 2017, 13, 370.	0.7	31
51	From genomes to genotypes: molecular epidemiological analysis of <i>Chlamydia gallinacea</i> reveals a high level of genetic diversity for this newly emerging chlamydial pathogen. <i>BMC Genomics</i> , 2017, 18, 949.	1.2	15
52	<i>Chlamydia pecorum</i> : successful pathogen of koalas or Australian livestock?. <i>Microbiology Australia</i> , 2017, 38, 101.	0.1	1
53	Humoral immune response against two surface antigens of <i>Chlamydia pecorum</i> in vaccinated and naturally infected sheep. <i>PLoS ONE</i> , 2017, 12, e0188370.	1.1	5
54	Development and evaluation of rapid novel isothermal amplification assays for important veterinary pathogens: <i>Chlamydia psittaci</i> and <i>Chlamydia pecorum</i> . <i>PeerJ</i> , 2017, 5, e3799.	0.9	39

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55	Culture-independent genomic characterisation of <i>Candidatus Chlamydia sanzinia</i> , a novel uncultivated bacterium infecting snakes. <i>BMC Genomics</i> , 2016, 17, 710.	1.2	65
56	A Prototype Recombinant-Protein Based <i>Chlamydia pecorum</i> Vaccine Results in Reduced Chlamydial Burden and Less Clinical Disease in Free-Ranging Koalas (<i>Phascolarctos cinereus</i>). <i>PLoS ONE</i> , 2016, 11, e0146934.	1.1	42
57	<i>Chlamydia</i> Serine Protease Inhibitor, targeting HtrA, as a New Treatment for Koala <i>Chlamydia</i> infection. <i>Scientific Reports</i> , 2016, 6, 31466.	1.6	27
58	Koala translocations and <i>Chlamydia</i> : Managing risk in the effort to conserve native species. <i>Biological Conservation</i> , 2016, 197, 247-253.	1.9	20
59	Treatment of <i>Chlamydia</i> -associated ocular disease via a recombinant protein based vaccine in the koala (<i>Phascolarctos cinereus</i>). <i>Biologicals</i> , 2016, 44, 588-590.	0.5	12
60	Characterisation of the immune compounds in koala milk using a combined transcriptomic and proteomic approach. <i>Scientific Reports</i> , 2016, 6, 35011.	1.6	25
61	Clinical, diagnostic and pathologic features of presumptive cases of <i>Chlamydia pecorum</i> -associated arthritis in Australian sheep flocks. <i>BMC Veterinary Research</i> , 2016, 12, 193.	0.7	27
62	<i>Chlamydial</i> infections in wildlife – conservation threats and/or reservoirs of “spill-over” infections?. <i>Veterinary Microbiology</i> , 2016, 196, 78-84.	0.8	43
63	Australian human and parrot <i>Chlamydia psittaci</i> strains cluster within the highly virulent 6BC clade of this important zoonotic pathogen. <i>Scientific Reports</i> , 2016, 6, 30019.	1.6	58
64	The emergence of sarcoptic mange in Australian wildlife: an unresolved debate. <i>Parasites and Vectors</i> , 2016, 9, 316.	1.0	45
65	Serum Antibody Response to Koala Retrovirus Antigens Varies in Free-Ranging Koalas (<i>Phascolarctos</i>) Tj ETQq1 1 0.784314 rsgBT /Over	0.3	15
66	PREVALENCE AND PATHOLOGIC FEATURES OF <i>CHLAMYDIA PECORUM</i> INFECTIONS IN SOUTH AUSTRALIAN KOALAS (<i>PHASCOLARCTOS CINEREUS</i>). <i>Journal of Wildlife Diseases</i> , 2016, 52, 301-306.	0.3	26
67	Humoral immune responses in koalas (<i>Phascolarctos cinereus</i>) either naturally infected with <i>Chlamydia pecorum</i> or following administration of a recombinant chlamydial major outer membrane protein vaccine. <i>Vaccine</i> , 2016, 34, 775-782.	1.7	21
68	HapFlow: visualizing haplotypes in sequencing data. <i>Bioinformatics</i> , 2016, 32, 441-443.	1.8	2
69	Identification of unusual <i>Chlamydia pecorum</i> genotypes in Victorian koalas (<i>Phascolarctos cinereus</i>) and clinical variables associated with infection. <i>Journal of Medical Microbiology</i> , 2016, 65, 420-428.	0.7	29
70	Antibody and Cytokine Responses of Koalas (<i>Phascolarctos cinereus</i>) Vaccinated with Recombinant Chlamydial Major Outer Membrane Protein (MOMP) with Two Different Adjuvants. <i>PLoS ONE</i> , 2016, 11, e0156094.	1.1	23
71	Identification, characterisation and expression analysis of natural killer receptor genes in <i>Chlamydia pecorum</i> infected koalas (<i>Phascolarctos cinereus</i>). <i>BMC Genomics</i> , 2015, 16, 796.	1.2	12
72	Genetic diversity in the plasticity zone and the presence of the chlamydial plasmid differentiates <i>Chlamydia pecorum</i> strains from pigs, sheep, cattle, and koalas. <i>BMC Genomics</i> , 2015, 16, 893.	1.2	40

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73	Phylogenetic analysis of human <i>Chlamydia pneumoniae</i> strains reveals a distinct Australian indigenous clade that predates European exploration of the continent. <i>BMC Genomics</i> , 2015, 16, 1094.	1.2	5
74	SNP Marker Discovery in Koala TLR Genes. <i>PLoS ONE</i> , 2015, 10, e0121068.	1.1	7
75	Comparative genomic analysis of human <i>Chlamydia pneumoniae</i> isolates from respiratory, brain and cardiac tissues. <i>Genomics</i> , 2015, 106, 373-383.	1.3	23
76	Characterisation of <i>Chlamydia pneumoniae</i> and other novel chlamydial infections in captive snakes. <i>Veterinary Microbiology</i> , 2015, 178, 88-93.	0.8	26
77	Twenty years of research into <i>Chlamydia</i> -like organisms: a revolution in our understanding of the biology and pathogenicity of members of the phylum <i>Chlamydiae</i> . <i>Pathogens and Disease</i> , 2015, 73, 1-15.	0.8	112
78	Novel Sequence Types of <i>Chlamydia pecorum</i> Infect Free-Ranging Alpine Ibex (<i>Capra ibex</i>) and Red Deer (<i>Cervus elaphus</i>) in Switzerland. <i>Journal of Wildlife Diseases</i> , 2015, 51, 479.	0.3	12
79	Culture-Independent Genome Sequencing of Clinical Samples Reveals an Unexpected Heterogeneity of Infections by <i>Chlamydia pecorum</i> . <i>Journal of Clinical Microbiology</i> , 2015, 53, 1573-1581.	1.8	40
80	<i>Chlamydia pecorum</i> infections in sheep and cattle: A common and under-recognised infectious disease with significant impact on animal health. <i>Veterinary Journal</i> , 2015, 206, 252-260.	0.6	53
81	A transcriptome resource for the koala (<i>Phascolarctos cinereus</i>): insights into koala retrovirus transcription and sequence diversity. <i>BMC Genomics</i> , 2014, 15, 786.	1.2	49
82	Chlamydial infections of fish: Diverse pathogens and emerging causes of disease in aquaculture species. <i>Veterinary Microbiology</i> , 2014, 170, 19-27.	0.8	39
83	Evaluation of the relationship between <i>Chlamydia pecorum</i> sequence types and disease using a species-specific multi-locus sequence typing scheme (MLST). <i>Veterinary Microbiology</i> , 2014, 174, 214-222.	0.8	37
84	Comparative genomics of koala, cattle and sheep strains of <i>Chlamydia pecorum</i> . <i>BMC Genomics</i> , 2014, 15, 667.	1.2	33
85	Molecular and pathological insights into <i>Chlamydia pecorum</i> -associated sporadic bovine encephalomyelitis (SBE) in Western Australia. <i>BMC Veterinary Research</i> , 2014, 10, 121.	0.7	29
86	Vaccination of koalas (<i>Phascolarctos cinereus</i>) with a recombinant chlamydial major outer membrane protein adjuvanted with poly I:C, a host defense peptide and polyphosphazine, elicits strong and long lasting cellular and humoral immune responses. <i>Vaccine</i> , 2014, 32, 5781-5786.	1.7	44
87	Chlamydial infections of fish: Diverse pathogens and emerging causes of disease in aquaculture species. <i>Veterinary Microbiology</i> , 2014, 171, 258-266.	0.8	15
88	Interleukin 17A is an immune marker for chlamydial disease severity and pathogenesis in the koala (<i>Phascolarctos cinereus</i>). <i>Developmental and Comparative Immunology</i> , 2014, 46, 423-429.	1.0	26
89	<i>Chlamydia</i> genomics: providing novel insights into chlamydial biology. <i>Trends in Microbiology</i> , 2014, 22, 464-472.	3.5	83
90	The koala immunological toolkit: sequence identification and comparison of key markers of the koala (<i>Phascolarctos cinereus</i>) immune response. <i>Australian Journal of Zoology</i> , 2014, 62, 195.	0.6	15

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91	Draft Genome and Plasmid Sequences of <i>Chlamydia pneumoniae</i> Strain B21 from an Australian Endangered Marsupial, the Western Barred Bandicoot. <i>Genome Announcements</i> , 2014, 2, .	0.8	7
92	Molecular characterisation and expression analysis of Interferon gamma in response to natural <i>Chlamydia</i> infection in the koala, <i>Phascolarctos cinereus</i> . <i>Gene</i> , 2013, 527, 570-577.	1.0	28
93	Molecular Characterization of <i>Candidatus Similichlamydia latridicola</i> gen. nov., sp. nov. (Chlamydiales: <i>Candidatus Parilichlamydiaceae</i>), a Novel <i>Chlamydia</i> -Like Epitheliocystis Agent in the Striped Trumpeter, <i>Latris lineata</i> (Forster). <i>Applied and Environmental Microbiology</i> , 2013, 79, 4914-4920.	1.4	37
94	Comparison of antigen detection and quantitative PCR in the detection of chlamydial infection in koalas (<i>Phascolarctos cinereus</i>). <i>Veterinary Journal</i> , 2013, 195, 391-393.	0.6	16
95	Antigenic specificity of a monovalent versus polyvalent MOMP based <i>Chlamydia pecorum</i> vaccine in koalas (<i>Phascolarctos cinereus</i>). <i>Vaccine</i> , 2013, 31, 1217-1223.	1.7	33
96	Recent advances in understanding the biology, epidemiology and control of chlamydial infections in koalas. <i>Veterinary Microbiology</i> , 2013, 165, 214-223.	0.8	173
97	Investigation of the koala (<i>Phascolarctos cinereus</i>) hindgut microbiome via 16S pyrosequencing. <i>Veterinary Microbiology</i> , 2013, 167, 554-564.	0.8	51
98	<i>Chlamydia pneumoniae</i> : modern insights into an ancient pathogen. <i>Trends in Microbiology</i> , 2013, 21, 120-128.	3.5	78
99	Genetic diversity of <i>Chlamydia pecorum</i> strains in wild koala locations across Australia and the implications for a recombinant <i>C. pecorum</i> major outer membrane protein based vaccine. <i>Veterinary Microbiology</i> , 2013, 167, 513-522.	0.8	43
100	Conjunctivitis Associated with <i>Chlamydia pecorum</i> in Three Koalas (<i>Phascolarctos</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 387 1066-1069.	0.3	8
101	Multilocus Sequence Analysis Provides Insights into Molecular Epidemiology of <i>Chlamydia pecorum</i> Infections in Australian Sheep, Cattle, and Koalas. <i>Journal of Clinical Microbiology</i> , 2013, 51, 2625-2632.	1.8	48
102	Molecular Characterization of <i>Candidatus Parilichlamydia carangidicola</i> , a Novel <i>Chlamydia</i> -Like Epitheliocystis Agent in Yellowtail Kingfish, <i>Seriola lalandi</i> (Valenciennes), and the Proposal of a New Family, <i>Candidatus Parilichlamydiaceae</i> fam. nov. (Order Chlamydiales). <i>Applied and Environmental Microbiology</i> , 2013, 79, 1590-1597.	1.4	65
103	Preliminary Characterisation of Tumor Necrosis Factor Alpha and Interleukin-10 Responses to <i>Chlamydia pecorum</i> Infection in the Koala (<i>Phascolarctos cinereus</i>). <i>PLoS ONE</i> , 2013, 8, e59958.	1.1	28
104	Vaccination of Koalas with a Recombinant <i>Chlamydia pecorum</i> Major Outer Membrane Protein Induces Antibodies of Different Specificity Compared to Those Following a Natural Live Infection. <i>PLoS ONE</i> , 2013, 8, e74808.	1.1	19
105	<i>Candidatus Similichlamydia laticola</i> , a Novel <i>Chlamydia</i> -like Agent of epitheliocystis in Seven Consecutive Cohorts of Farmed Australian Barramundi, <i>Lates calcarifer</i> (Bloch). <i>PLoS ONE</i> , 2013, 8, e82889.	1.1	25
106	Vaccination of healthy and diseased koalas (<i>Phascolarctos cinereus</i>) with a <i>Chlamydia pecorum</i> multi-subunit vaccine: Evaluation of immunity and pathology. <i>Vaccine</i> , 2012, 30, 1875-1885.	1.7	59
107	A natural freshwater origin for two chlamydial species, <i>Candidatus</i> <i>Piscichlamydia salmonis</i> and <i>Candidatus</i> <i>Clavochlamydia salmonicola</i> , causing mixed infections in wild brown trout (<i>Salmo trutta</i>). <i>Environmental Microbiology</i> , 2012, 14, 2048-2057.	1.8	39
108	Novel molecular markers of <i>Chlamydia pecorum</i> genetic diversity in the koala (<i>Phascolarctos</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 62 T	1.3	50

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109	Chlamydia abortus YhbZ, a truncated Obg family GTPase, associates with the Escherichia coli large ribosomal subunit. Microbial Pathogenesis, 2011, 50, 200-206.	1.3	12
110	Using quantitative polymerase chain reaction to correlate Chlamydia pecorum infectious load with ocular, urinary and reproductive tract disease in the koala (Phascolarctos cinereus). Australian Veterinary Journal, 2011, 89, 409-412.	0.5	69
111	Novel Chlamydiales associated with epitheliocystis in a leopard shark Triakis semifasciata. Diseases of Aquatic Organisms, 2010, 91, 75-81.	0.5	34
112	The Waddlia Genome: A Window into Chlamydial Biology. PLoS ONE, 2010, 5, e10890.	1.1	104
113	Molecular evidence for chlamydial infections in the eyes of sheep. Veterinary Microbiology, 2009, 135, 142-146.	0.8	49
114	Predator or prey? Chlamydophila abortus infections of a free-living amoebae, Acanthamoeba castellanii 9GU. Microbes and Infection, 2008, 10, 591-597.	1.0	8
115	Chlamydophila pneumoniae HflX belongs to an uncharacterized family of conserved GTPases and associates with the Escherichia coli 50S large ribosomal subunit. Microbiology (United Kingdom), 2008, 154, 3537-3546.	0.7	26
116	Prevalence of chlamydiae in semen and genital tracts of bulls, rams and bucks. Theriogenology, 2007, 67, 303-310.	0.9	48
117	Ovine Enzootic Abortion (OEA): a comparison of antibody responses in vaccinated and naturally-infected swiss sheep over a two year period. BMC Veterinary Research, 2007, 3, 24.	0.7	18
118	Differential expression of chlamydial signal transduction genes in normal and interferon gamma-induced persistent Chlamydophila pneumoniae infections. Microbes and Infection, 2006, 8, 61-72.	1.0	27
119	Detection of chlamydiae in boar semen and genital tracts. Veterinary Microbiology, 2006, 116, 149-157.	0.8	32
120	Detection of Mycobacteria and Chlamydiae in Granulomatous Inflammation of Reptiles: A Retrospective Study. Veterinary Pathology, 2004, 41, 388-397.	0.8	98
121	Detection of novel chlamydiae in cats with ocular disease. American Journal of Veterinary Research, 2003, 64, 1421-1428.	0.3	67
122	Molecular Evidence for Novel Chlamydial Infections in the Koala (Phascolarctos cinereus). Systematic and Applied Microbiology, 2003, 26, 245-253.	1.2	50
123	The Koala Genome Consortium. Technical Reports of the Australian Museum Online, 0, 24, 91-92.	0.0	7