Jan Slapeta

List of Publications by Year in descending order

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117625 133252 4,889 187 34 59 citations g-index h-index papers 190 190 190 4629 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	A photosynthetic alveolate closely related to apicomplexan parasites. Nature, 2008, 451, 959-963.	27.8	437
2	Global Dispersal and Ancient Cryptic Species in the Smallest Marine Eukaryotes. Molecular Biology and Evolution, 2006, 23, 23-29.	8.9	210
3	The extent of protist diversity: insights from molecular ecology of freshwater eukaryotes. Proceedings of the Royal Society B: Biological Sciences, 2005, 272, 2073-2081.	2.6	203
4	Australian dingoes are definitive hosts of Neospora caninum. International Journal for Parasitology, 2010, 40, 945-950.	3.1	188
5	A review of neosporosis and pathologic findings of Neospora caninum infection in wildlife. International Journal for Parasitology: Parasites and Wildlife, 2015, 4, 216-238.	1.5	148
6	Cryptosporidiosis and Cryptosporidium species in animals and humans: A thirty colour rainbow?. International Journal for Parasitology, 2013, 43, 957-970.	3.1	119
7	Molecular identification of Cryptosporidium spp. in animal and human hosts from the Czech Republic. Veterinary Parasitology, 2004, 122, 183-192.	1.8	102
8	The Phylogeny of Goussia and Choleoeimeria (Apicomplexa; Eimeriorina) and the Evolution of Excystation Structures in Coccidia. Protist, 2002, 153, 379-390.	1.5	87
9	Cryptosporidia: Epicellular parasites embraced by the host cell membrane. International Journal for Parasitology, 2008, 38, 913-922.	3.1	72
10	The â€~other' coral symbiont: <i>Ostreobium</i> diversity and distribution. ISME Journal, 2017, 11, 296-299.	9.8	72
11	A Genetically Tractable, Natural Mouse Model of Cryptosporidiosis Offers Insights into Host Protective Immunity. Cell Host and Microbe, 2019, 26, 135-146.e5.	11.0	72
12	High phylogenetic diversity of the cat flea (<i>Ctenocephalides felis</i>) at two mitochondrial <scp>DNA</scp> markers. Medical and Veterinary Entomology, 2014, 28, 330-336.	1.5	71
13	Cryptosporidium parvum Mitochondrial-Type HSP70 Targets Homologous and Heterologous Mitochondria. Eukaryotic Cell, 2004, 3, 483-494.	3.4	65
14	Parasite spread at the domestic animal - wildlife interface: anthropogenic habitat use, phylogeny and body mass drive risk of cat and dog flea (Ctenocephalides spp.) infestation in wild mammals. Parasites and Vectors, 2018, 11, 8.	2.5	64
15	Phylogenetic analysis of Sarcocystis spp. of mammals and reptiles supports the coevolution of Sarcocystis spp. with their final hosts1Note: The nucleotide sequences of Sarcocystis dispersa and Sarcocystis sp. have been deposited in the GenBankâ,, under the accession numbers AF120115 and AF120114, respectively.1. International Journal for Parasitology, 1999, 29, 795-798.	3.1	61
16	Tritrichomonas foetus from domestic cats and cattle are genetically distinct. Experimental Parasitology, 2010, 126, 209-213.	1.2	61
17	A curious coincidence: mosquito biodiversity and the limits of the Japanese encephalitis virus in Australasia. BMC Evolutionary Biology, 2007, 7, 100.	3.2	59
18	Differences in the faecal microbiome of non-diarrhoeic clinically healthy dogs and cats associated with Giardia duodenalis infection: impact of hookworms and coccidia. International Journal for Parasitology, 2015, 45, 585-594.	3.1	59

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19	Evolutionary relationships among cyst-forming coccidia Sarcocystis spp. (Alveolata: Apicomplexa:) Tj ETQq1 1 C	0.784314 rg 2.7	gBT /Overlock 57
20	Molecular Phylogenetics and Evolution, 2003, 27, 464-475. Integrated morphological and molecular identification of cat fleas (Ctenocephalides felis) and dog fleas (Ctenocephalides canis) vectoring Rickettsia felis in central Europe. Veterinary Parasitology, 2015, 210, 215-223.	1.8	55
21	Cryptosporidium species found in cattle: a proposal for a new species. Trends in Parasitology, 2006, 22, 469-474.	3.3	53
22	Out-of-Africa, human-mediated dispersal of the common cat flea, Ctenocephalides felis: The hitchhiker's guide to world domination. International Journal for Parasitology, 2019, 49, 321-336.	3.1	51
23	Comparative analysis of Tritrichomonas foetus (Riedmüller, 1928) cat genotype, T. foetus (Riedmüller,) Tj F	ETQq1 1 0. 3.1	784314 rgBT 49
24	A Suspected Parasite Spill-Back of Two Novel Myxidium spp. (Myxosporea) Causing Disease in Australian Endemic Frogs Found in the Invasive Cane Toad. PLoS ONE, 2011, 6, e18871.	2.5	49
25	New species of Cryptosporidium Tyzzer, 1907 (Apicomplexa) from amphibian host: morphology, biology and phylogeny. Folia Parasitologica, 2008, 55, 81-94.	1.3	49
26	The first report of ovine cerebral neosporosis and evaluation of Neospora caninum prevalence in sheep in New South Wales. Veterinary Parasitology, 2010, 170, 137-142.	1.8	48
27	Cryptosporidium from tortoises: Genetic characterisation, phylogeny and zoonotic implications. Molecular and Cellular Probes, 2008, 22, 122-128.	2.1	45
28	Oocysts and high seroprevalence of Neospora caninum in dogs living in remote Aboriginal communities and wild dogs in Australia. Veterinary Parasitology, 2012, 187, 85-92.	1.8	45
29	Conservation of proteins involved in oocyst wall formation in Eimeria maxima, Eimeria tenella and Eimeria acervulina. International Journal for Parasitology, 2009, 39, 1063-1070.	3.1	42
30	Implications of wild dog ecology on the sylvatic and domestic life cycle of Neospora caninum in Australia. Veterinary Journal, 2011, 188, 24-33.	1.7	42
31	Evidence for a specific host-endosymbiont relationship between †Rickettsia sp. genotype RF2125†and Ctenocephalides felis orientis infesting dogs in India. Parasites and Vectors, 2015, 8, 169.	2.5	40
32	The "tropical lineage―of the brown dog tick Rhipicephalus sanguineus sensu lato identified as Rhipicephalus linnaei (). International Journal for Parasitology, 2021, 51, 431-436.	3.1	40
33	Naturally occurring Tritrichomonas foetus infections in Australian cats: 38 cases. Journal of Feline Medicine and Surgery, 2010, 12, 889-898.	1.6	38
34	The cat flea (Ctenocephalides f. felis) is the dominant flea on domestic dogs and cats in Australian veterinary practices. Veterinary Parasitology, 2011, 180, 383-388.	1.8	37
35	Resolving genetic diversity in Australasian Culex mosquitoes: Incongruence between the mitochondrial cytochrome c oxidase I and nuclear acetylcholine esterase 2. Molecular Phylogenetics and Evolution, 2009, 50, 317-325.	2.7	34
36	Reptile-associated Borrelia species in the goanna tick (Bothriocroton undatum) from Sydney, Australia. Parasites and Vectors, 2017, 10, 616.	2.5	34

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37	Low diversity of Angiostrongylus cantonensis complete mitochondrial DNA sequences from Australia, Hawaii, French Polynesia and the Canary Islands revealed using whole genome next-generation sequencing. Parasites and Vectors, 2019, 12, 241.	2.5	34
38	Phylogenetic analysis of the light-harvesting system in Chromera velia. Photosynthesis Research, 2012, 111, 19-28.	2.9	32
39	Prolonged resilience of Tritrichomonas foetus in cat faeces at ambient temperature. Veterinary Parasitology, 2009, 166, 60-65.	1.8	31
40	Museum material reveals a frog parasite emergence after the invasion of the cane toad in Australia. Parasites and Vectors, 2010, 3, 50.	2.5	31
41	Peroxidase catalysed cross-linking of an intrinsically unstructured protein via dityrosine bonds in the oocyst wall of the apicomplexan parasite, Eimeria maxima. International Journal for Parasitology, 2011, 41, 1157-1164.	3.1	31
42	Sequence differences in the diagnostic region of the cysteine protease 8 gene of Tritrichomonas foetus parasites of cats and cattle. Veterinary Parasitology, 2012, 186, 445-449.	1.8	31
43	Combined Amplicon Pyrosequencing Assays Reveal Presence of the Apicomplexan "type-N―(cf.) Tj ETQq1 ☐ e76095.	l 0.784314 2.5	rgBT /Overlo
44	Comparative transcriptomics reveals striking similarities between the bovine and feline isolates of Tritrichomonas foetus: consequences for in silico drug-target identification. BMC Genomics, 2014, 15, 955.	2.8	31
45	Evaluation of the bacterial microbiome of two flea species using different DNA-isolation techniques provides insights into flea host ecology. FEMS Microbiology Ecology, 2015, 91, fiv134.	2.7	31
46	Multiple origin of the dihomoxenous life cycle in sarcosporidia. International Journal for Parasitology, 2001, 31, 413-417.	3.1	30
47	The flagellar contribution to the apical complex: a new tool for the eukaryotic Swiss Army knife?. Trends in Parasitology, 2014, 30, 58-64.	3.3	30
48	Evidence of Intraflagellar Transport and Apical Complex Formation in a Free-Living Relative of the Apicomplexa. Eukaryotic Cell, 2014, 13, 10-20.	3.4	30
49	The Occurrence and Apeâ€toâ€Ape Transmission of the Entodiniomorphid Ciliate <i>Troglodytella abrassarti</i> in Captive Gorillas. Journal of Eukaryotic Microbiology, 2009, 56, 83-87.	1.7	29
50	Cat fleas (Ctenocephalides felis) from cats and dogs in New Zealand: Molecular characterisation, presence of Rickettsia felis and Bartonella clarridgeiae and comparison with Australia. Veterinary Parasitology, 2017, 234, 25-30.	1.8	28
51	New species of Myxosporea from frogs and resurrection of the genus <i>Cystodiscus</i> Lutz, 1889 for species with myxospores in gallbladders of amphibians. Parasitology, 2012, 139, 478-496.	1.5	27
52	New insights on the epidemiology of Coxiella burnetii in pet dogs and cats from New South Wales, Australia. Acta Tropica, 2020, 205, 105416.	2.0	27
53	Nonreductive Iron Uptake Mechanism in the Marine Alveolate <i>Chromera velia</i> \hat{A} \hat{A} . Plant Physiology, 2010, 154, 991-1000.	4.8	26
54	Scrambled eggs: A highly sensitive molecular diagnostic workflow for Fasciola species specific detection from faecal samples. PLoS Neglected Tropical Diseases, 2017, 11, e0005931.	3.0	26

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55	Notes on coccidian phylogeny, based on the apicoplast small subunit ribosomal DNA. Parasitology Research, 2002, 88, 360-363.	1.6	25
56	Comparison of early detection of Fasciola hepatica in experimentally infected Merino sheep by real-time PCR, coproantigen ELISA and sedimentation. Veterinary Parasitology, 2018, 251, 85-89.	1.8	24
57	Phylogenetic analyses suggest lateral gene transfer from the mitochondrion to the apicoplast. Gene, 2002, 285, 109-118.	2.2	23
58	Phylogeny and sequence variability of the Sarcocystis singaporensis Zaman and Colley, (1975) 1976 ssrDNA. Parasitology Research, 2002, 88, 810-815.	1.6	23
59	Afternoon shedding of a new species of <i>Isospora </i> (Apicomplexa) in the endangered Regent Honeyeater (<i>Xanthomyza phrygia </i>). Parasitology, 2011, 138, 713-724.	1.5	23
60	Detection of <i>Cryptosporidium molnari</i> Oocysts from Fish by Fluorescent-Antibody Staining Assays for <i>Cryptosporidium</i> spp. Affecting Humans. Applied and Environmental Microbiology, 2011, 77, 1878-1880.	3.1	23
61	Multisystemic toxoplasmosis associated with a type II-like Toxoplasma gondii strain in a New Zealand fur seal (Arctocephalus forsteri) from New South Wales, Australia. Veterinary Parasitology, 2014, 205, 347-353.	1.8	23
62	Mosquito-borne heartworm Dirofilaria immitis in dogs from Australia. Parasites and Vectors, 2016, 9, 535.	2.5	23
63	Fasciola Species Introgression: Just a Fluke or Something More?. Trends in Parasitology, 2021, 37, 25-34.	3.3	23
64	Detection of Dientamoeba fragilis in animal faeces using species specific real time PCR assay. Veterinary Parasitology, 2016, 227, 42-47.	1.8	22
65	Apicomplexa., 2017,, 567-624.		21
66	Molecular evidence confirms occurrence of Rhipicephalus microplus Clade A in Kenya and sub-Saharan Africa. Parasites and Vectors, 2020, 13, 432.	2.5	21
67	Pig-hunting dogs are an at-risk population for canine heartworm (Dirofilaria immitis) infection in eastern Australia. Parasites and Vectors, 2020, 13, 69.	2.5	21
68	AcanR3990 qPCR: A Novel, Highly Sensitive, Bioinformatically-Informed Assay to Detect <i>Angiostrongylus cantonensis</i> Infections. Clinical Infectious Diseases, 2021, 73, e1594-e1600.	5.8	21
69	Intercontinental distribution of a new trypanosome species from Australian endemic Regent Honeyeater (<i>Anthochaera phrygia</i>). Parasitology, 2016, 143, 1012-1025.	1.5	20
70	A real-time PCR tool for the surveillance of zoonotic Onchocerca lupi in dogs, cats and potential vectors. PLoS Neglected Tropical Diseases, 2018, 12, e0006402.	3.0	20
71	Global selective sweep of a highly inbred genome of the cattle parasite Neospora caninum. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 22764-22773.	7.1	20
72	Apicomplexa. , 2016, , 1-58.		20

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73	Occurrence, diagnosis and follow-up of canine strongyloidiosis in naturally infected shelter dogs. Parasitology, 2019, 146, 246-252.	1.5	19
74	Efficacy of ivermectin to control Strongyloides stercoralis infection in sheltered dogs. Acta Tropica, 2019, 190, 204-209.	2.0	19
75	Sarcocystis atheridis sp. nov., a new sarcosporidian coccidium from Nitsche's bush viper, Atheris nitschei Tornier, 1902, from Uganda. Parasitology Research, 1999, 85, 758-764.	1.6	18
76	Naming of Cryptosporidium pestis is in accordance with the ICZN Code and the name is available for this taxon previously recognized as C. parvum †bovine genotype'. Veterinary Parasitology, 2011, 177, 1-5.	1.8	18
77	Surface and Flagella Morphology of the Motile Form of Chromera velia Revealed by Field-Emission Scanning Electron Microscopy. Protist, 2011, 162, 142-153.	1.5	18
78	Extensive production of Neospora caninum tissue cysts in a carnivorous marsupial succumbing to experimental neosporosis. Veterinary Research, 2011, 42, 75.	3.0	18
79	Clinical and pathological features of toxoplasmosis in free-ranging common wombats (Vombatus) Tj ETQq1 1 0.7 International, 2015, 64, 148-153.	84314 rgE 1.3	3T /Overlock 18
80	Accurate identification of Australian mosquitoes using protein profiling. Parasitology, 2019, 146, 462-471.	1.5	18
81	Cat genotype Tritrichomonas foetus survives passage through the alimentary tract of two common slug species. Veterinary Parasitology, 2011, 177, 262-266.	1.8	17
82	Ten simple rules for describing a new (parasite) species. International Journal for Parasitology: Parasites and Wildlife, 2013, 2, 152-154.	1.5	17
83	Quantitative PCR detection of Theileria equi using laboratory workflows to detect asymptomatic persistently infected horses. Veterinary Parasitology, 2014, 206, 138-145.	1.8	17
84	Comparative RNA-seq analysis of the Tritrichomonas foetus PIG30/1 isolate from pigs reveals close association with Tritrichomonas foetus BP-4 isolate †bovine genotype'. Veterinary Parasitology, 2015, 212, 111-117.	1.8	17
85	The utility of diversity profiling using Illumina 18S rRNA gene amplicon deep sequencing to detect and discriminate Toxoplasma gondii among the cyst-forming coccidia. Veterinary Parasitology, 2016, 216, 38-45.	1.8	17
86	Uncinaria sanguinis sp. n. (Nematoda: Ancylostomatidae) from the endangered Australian sea lion, Neophoca cinerea (Carnivora: Otariidae). Folia Parasitologica, 2014, 61, 255-265.	1.3	17
87	High prevalence of Tritrichomonas foetus †bovine genotype' in faecal samples from domestic pigs at a farm where bovine trichomonosis has not been reported for over 30 years. Veterinary Parasitology, 2015, 212, 105-110.	1.8	16
88	Comparison of genotypes of Toxoplasma gondii in domestic cats from Australia with latent infection or clinical toxoplasmosis. Veterinary Parasitology, 2016, 228, 13-16.	1.8	16
89	Comparative proteomic analysis of two pathogenic Tritrichomonas foetus genotypes: there is more to the proteome than meets the eye. International Journal for Parasitology, 2017, 47, 203-213.	3.1	16
90	Cat fleas (Ctenocephalides felis) carrying Rickettsia felis and Bartonella species in Hong Kong. Parasitology International, 2018, 67, 209-212.	1.3	16

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91	Climate change models predict southerly shift of the cat flea (Ctenocephalides felis) distribution in Australia. Parasites and Vectors, 2019, 12, 137.	2.5	16
92	Eimeria telekii n.sp. (Apicomplexa: Coccidia) from Lemniscomys striatus (Rodentia: Muridae): morphology, pathology and phylogeny. Parasitology, 2001, 122, 133-43.	1.5	15
93	The Glycosylation Pathway of <i>Eimeria tenella</i> Is Upregulated during Gametocyte Development and May Play a Role in Oocyst Wall Formation. Eukaryotic Cell, 2010, 9, 127-135.	3.4	15
94	The mitochondrial genome of <i>Angiostrongylus mackerrasae</i> is distinct from <ia. cantonensis<="" i=""> and <ia. i="" malaysiensis<="">. Parasitology, 2020, 147, 681-688.</ia.></ia.>	1.5	15
95	Which species is in the faeces at a time of global livestock movements: single nucleotide polymorphism genotyping assays for the differentiation of Fasciola spp International Journal for Parasitology, 2020, 50, 91-101.	3.1	15
96	Unexpected absence of genetic separation of a highly diverse population of hookworms from geographically isolated hosts. Infection, Genetics and Evolution, 2014, 28, 192-200.	2.3	14
97	Molecular identity of cat fleas (Ctenocephalides felis) from cats in Georgia, USA carrying Bartonella clarridgeiae, Bartonella henselae and Rickettsia sp. RF2125. Veterinary Parasitology: Regional Studies and Reports, 2016, 3-4, 36-40.	0.5	14
98	High seroprevalance of Neospora caninum in dogs in Victoria, Australia, compared to 20Âyears ago. Parasites and Vectors, 2017, 10, 503.	2.5	14
99	Wild horse populations in south-east Australia have a high prevalence of Strongylus vulgaris and may act as a reservoir of infection for domestic horses. International Journal for Parasitology: Parasites and Wildlife, 2019, 8, 156-163.	1.5	14
100	Meta-transcriptomic identification of Trypanosoma spp. in native wildlife species from Australia. Parasites and Vectors, 2020, 13, 447.	2.5	14
101	Rhipicephalus sanguineus sensu lato from dogs and dromedary camels in Riyadh, Saudi Arabia: low prevalence of vector-borne pathogens in dogs detected using multiplexed tandem PCR panel. Folia Parasitologica, 2019, 66, .	1.3	14
102	A new diagnostic approach to fast-track and increase the accessibility of gastrointestinal nematode identification from faeces: FECPAKG2 egg nemabiome metabarcoding. International Journal for Parasitology, 2022, 52, 331-342.	3.1	14
103	Looks can deceive: Molecular identity of an intraerythrocytic apicomplexan parasite in Australian gliders. Veterinary Parasitology, 2009, 159, 105-111.	1.8	13
104	Phylogenetic relationships of Habronema microstoma and Habronema muscae (Spirurida:) Tj ETQq0 0 0 rgBT /Ov subunit 1 (cox1) gene analysis. Parasitology Research, 2009, 104, 979-984.	erlock 10 1.6	Tf 50 227 Td 12
105	Effect of Nutrient Concentration and Salinity on Immotile–Motile Transformation of <i>Chromera velia</i> . Journal of Eukaryotic Microbiology, 2010, 57, 444-446.	1.7	12
106	New species of Cryptosporidium Tyzzer, 1907 (Apicomplexa) from amphibian host: morphology, biology and phylogeny. Folia Parasitologica, 2008, 55, 81-94.	1.3	12
107	ANarf-like gene fromCryptosporidium parvumresembles homologues observed in aerobic protists and higher eukaryotes. FEMS Microbiology Letters, 2003, 229, 91-96.	1.8	11
108	Differential Gamma Interferon- and Tumor Necrosis Factor Alpha-Driven Cytokine Response Distinguishes Acute Infection of a Metatherian Host with Toxoplasma gondii and Neospora caninum. Infection and Immunity, 2017, 85, .	2.2	11

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109	Invasive Colonic Entamoebiasis in Wild Cane Toads, Australia. Emerging Infectious Diseases, 2018, 24, 1541-1543.	4.3	11
110	Fleas from domestic dogs and rodents in Rwanda carry <i>Rickettsia asembonensis</i> and <i>Bartonella tribocorum</i> . Medical and Veterinary Entomology, 2019, 33, 177-184.	1.5	11
111	The brown dog tick Rhipicephalus sanguineus sensu Roberts, 1965 across Australia: Morphological and molecular identification of R. sanguineus s.l. tropical lineage. Ticks and Tick-borne Diseases, 2020, 11, 101305.	2.7	11
112	The molecular identity of fleas (Siphonaptera) carrying Rickettsia felis, Bartonella clarridgeiae and Bartonella rochalimae from dogs and cats in Northern Laos. Heliyon, 2020, 6, e04385.	3.2	11
113	Unbiased Characterization of the Microbiome and Virome of Questing Ticks. Frontiers in Microbiology, 2021, 12, 627327.	3.5	11
114	Canine parvovirus is shed infrequently by cats without diarrhoea in multi-cat environments. Veterinary Microbiology, 2021, 261, 109204.	1.9	11
115	Centenary of the genus <i>Cryptosporidium</i> : from morphological to molecular species identification, 2009, , 31-50.		11
116	Description of Eimeria motelo sp. n. (Apicomplexa: Eimeriidae) from the yellow footed tortoise, Geochelone denticulata (Chelonia: Testudinidae), and replacement of Eimeria carinii Lainson, Costa & Shaw, 1990 by Eimeria lainsoni nom. nov Memorias Do Instituto Oswaldo Cruz, 2000, 95, 829-832.	1.6	10
117	Present Status of the Molecular Ecology of Kathablepharids. Protist, 2006, 157, 7-11.	1.5	10
118	Neosporosis in an Aborted Southern White Rhinoceros (Ceratotherium simum simum) Fetus. Journal of Zoo and Wildlife Medicine, 2010, 41, 725-728.	0.6	10
119	Gastric cryptosporidiosis in farmed Australian Murray cod, Maccullochella peelii peelii. Aquaculture, 2011, 314, 1-6.	3.5	10
120	Myxosporean parasites in Australian frogs: Importance, implications and future directions. International Journal for Parasitology: Parasites and Wildlife, 2013, 2, 62-68.	1.5	10
121	Increased growth and pigment content of <i>Chromera velia </i> in mixotrophic culture. FEMS Microbiology Ecology, 2014, 88, 121-128.	2.7	10
122	MT-PCR panel detection of canine parvovirus (CPV-2): Vaccine and wild-type CPV-2 can be difficult to differentiate in canine diagnostic fecal samples. Molecular and Cellular Probes, 2017, 33, 20-23.	2.1	10
123	<i>Cryptosporidium</i> : Identification and Genetic Typing. Current Protocols in Microbiology, 2017, 44, 20B.1.1-20B.1.17.	6.5	10
124	Comparison of multiplexed-tandem real-time PCR panel with reference real-time PCR molecular diagnostic assays for detection of Giardia intestinalis and Tritrichomonas foetus in cats. Veterinary Parasitology, 2019, 266, 12-17.	1.8	10
125	Further studies of neuroangiostrongyliasis (rat lungworm disease) in Australian dogs: 92 new cases (2010–2020) and results for a novel, highly sensitive qPCR assay. Parasitology, 2021, 148, 178-186.	1.5	10
126	Characterization of S-adenosylmethionine synthetase in Cryptosporidium parvum (Apicomplexa). FEMS Microbiology Letters, 2003, 225, 271-277.	1.8	9

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127	Emerging myxosporean parasites of Australian frogs take a ride with fresh fruit transport. Parasites and Vectors, 2012, 5, 208.	2.5	9
128	Comparative Pathology and Ecological Implications of Two Myxosporean Parasites in Native Australian Frogs and the Invasive Cane Toad. PLoS ONE, 2012, 7, e43780.	2.5	9
129	Neonatal neosporosis in a 2-week-old Bernese mountain dog infected with multiple Neospora caninum strains based on MS10 microsatellite analysis. Veterinary Parasitology, 2016, 221, 134-138.	1.8	9
130	Biotic Factors Influence Microbiota of Nymph Ticks from Vegetation in Sydney, Australia. Pathogens, 2020, 9, 566.	2.8	9
131	Monophyly of marsupial intraerythrocytic apicomplexan parasites from South America and Australia. Parasitology, 2010, 137, 37-43.	1.5	8
132	Severe Amoebic Placentitis in a Horse Caused by an Acanthamoeba hatchetti Isolate Identified Using Next-Generation Sequencing. Journal of Clinical Microbiology, 2014, 52, 3101-3104.	3.9	8
133	Host origin determines pH tolerance of Tritrichomonas foetus isolates from the feline gastrointestinal and bovine urogenital tracts. Experimental Parasitology, 2015, 157, 68-77.	1.2	8
134	Novel genotype of <i>Tritrichomonas foetus </i> from cattle in Southern Africa. Parasitology, 2016, 143, 1954-1959.	1.5	8
135	Deep-sequencing to resolve complex diversity of apicomplexan parasites in platypuses and echidnas: Proof of principle for wildlife disease investigation. Infection, Genetics and Evolution, 2017, 55, 218-227.	2.3	8
136	Response to Xiao et al.: Further debate on the description of Cryptosporidium pestis. Trends in Parasitology, 2007, 23, 42-43.	3.3	7
137	DNA barcoding ofCryptosporidium. Parasitology, 2018, 145, 574-584.	1.5	7
138	Multiple diagnostic tests demonstrate an increased risk of canine heartworm disease in northern Queensland, Australia. Parasites and Vectors, 2021, 14, 393.	2.5	7
139	Toxoplasmosis in a Pet Peach-Faced Lovebird(<i>Agapornis roseicollis</i>). Korean Journal of Parasitology, 2015, 53, 749-753.	1.3	7
140	Year-Long Presence of Eimeria echidnae and Absence of Eimeria tachyglossi In Captive Short-Beaked Echidnas (Tachyglossus aculeatus). Journal of Parasitology, 2012, 98, 543.	0.7	6
141	Myxozoan Parasite in Brain of Critically Endangered Frog. Emerging Infectious Diseases, 2012, 18, 693-695.	4.3	6
142	Unusual presentation of neosporosis in a neonatal puppy from a litter of bulldogs. Australian Veterinary Journal, 2016, 94, 411-414.	1.1	6
143	Shelter-housed cats show no evidence of faecal shedding of canine parvovirus DNA. Veterinary Journal, 2018, 239, 54-58.	1.7	6
144	Ctenocephalides felis (cat flea) infestation in neonatal dairy calves managed with deltamethrin pour-on in Australia. Veterinary Parasitology, 2020, 279, 109039.	1.8	6

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145	Which larvae are they? Use of single larva for the molecular confirmation of Cooperia pectinata and Cooperia punctata in Australian cattle. Veterinary Parasitology, 2020, 278, 109033.	1.8	6
146	Adaptation of gltA and ssrA assays for diversity profiling by Illumina sequencing to identify Bartonella henselae, B. clarridgeiae and B. koehlerae. Journal of Medical Microbiology, 2021, 70, .	1.8	6
147	Seroprevalence of Neospora caninum in dogs from greater Sydney, Australia unchanged from 1997 to 2019 and worldwide review of adult-onset of canine neosporosis. Current Research in Parasitology and Vector-borne Diseases, 2021, 1, 100005.	1.9	6
148	Identification of Chromera velia by fluorescence in situ hybridization. FEMS Microbiology Letters, 2012, 328, 144-149.	1.8	5
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