

# Una M Ryan

## List of Publications by Year in descending order

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252  
papers

11,972  
citations

34016

52  
h-index

37111

96  
g-index

262  
all docs

262  
docs citations

262  
times ranked

6009  
citing authors

#	ARTICLE	IF	CITATIONS
1	Cryptosporidium Taxonomy: Recent Advances and Implications for Public Health. <i>Clinical Microbiology Reviews</i> , 2004, 17, 72-97.	5.7	742
2	<i>Cryptosporidium</i> species in humans and animals: current understanding and research needs. <i>Parasitology</i> , 2014, 141, 1667-1685.	0.7	505
3	Zoonotic potential of Giardia. <i>International Journal for Parasitology</i> , 2013, 43, 943-956.	1.3	451
4	Molecular epidemiology of giardiasis. <i>Molecular and Biochemical Parasitology</i> , 2008, 160, 75-80.	0.5	403
5	Genetic Diversity and Population Structure of Cryptosporidium. <i>Trends in Parasitology</i> , 2018, 34, 997-1011.	1.5	365
6	Identification of Novel Cryptosporidium Genotypes from the Czech Republic. <i>Applied and Environmental Microbiology</i> , 2003, 69, 4302-4307.	1.4	311
7	Host adaptation and host-parasite co-evolution in Cryptosporidium: implications for taxonomy and public health. <i>International Journal for Parasitology</i> , 2002, 32, 1773-1785.	1.3	252
8	Cryptosporidiosis: an update in molecular epidemiology. <i>Current Opinion in Infectious Diseases</i> , 2004, 17, 483-490.	1.3	238
9	Cryptosporidium and Giardia in Africa: current and future challenges. <i>Parasites and Vectors</i> , 2017, 10, 195.	1.0	188
10	<i>Cryptosporidium</i> in humans and animals—a one health approach to prophylaxis. <i>Parasite Immunology</i> , 2016, 38, 535-547.	0.7	187
11	Sheep May Not Be an Important Zoonotic Reservoir for Cryptosporidium and Giardia Parasites. <i>Applied and Environmental Microbiology</i> , 2005, 71, 4992-4997.	1.4	183
12	Subtyping <i>Cryptosporidium ubiquitum</i> , a Zoonotic Pathogen Emerging in Humans. <i>Emerging Infectious Diseases</i> , 2014, 20, 217-224.	2.0	172
13	Identification of Novel Cryptosporidium Genotypes from Avian Hosts. <i>Applied and Environmental Microbiology</i> , 2006, 72, 7548-7553.	1.4	164
14	Comparison of next-generation droplet digital PCR (ddPCR) with quantitative PCR (qPCR) for enumeration of Cryptosporidium oocysts in faecal samples. <i>International Journal for Parasitology</i> , 2014, 44, 1105-1113.	1.3	152
15	Recent insights into the tick microbiome gained through next-generation sequencing. <i>Parasites and Vectors</i> , 2018, 11, 12.	1.0	146
16	Foodborne cryptosporidiosis. <i>International Journal for Parasitology</i> , 2018, 48, 1-12.	1.3	143
17	Public health significance of zoonotic Cryptosporidium species in wildlife: Critical insights into better drinking water management. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2016, 5, 88-109.	0.6	142
18	Giardia: an under-reported foodborne parasite. <i>International Journal for Parasitology</i> , 2019, 49, 1-11.	1.3	131

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19	Taxonomy and molecular epidemiology of <i>Cryptosporidium</i> and <i>Giardia</i> – a 50-year perspective (1971–2021). <i>International Journal for Parasitology</i> , 2021, 51, 1099-1119.	1.3	128
20	New developments in <i>Cryptosporidium</i> research. <i>International Journal for Parasitology</i> , 2015, 45, 367-373.	1.3	124
21	Specific and quantitative detection and identification of <i>Cryptosporidium hominis</i> and <i>C. parvum</i> in clinical and environmental samples. <i>Experimental Parasitology</i> , 2013, 135, 142-147.	0.5	123
22	<i>Cryptosporidium</i> Rabbit Genotype, a Newly Identified Human Pathogen. <i>Emerging Infectious Diseases</i> , 2009, 15, 829-830.	2.0	122
23	<i>Cryptosporidium</i> in birds, fish and amphibians. <i>Experimental Parasitology</i> , 2010, 124, 113-120.	0.5	118
24	Genetic Diversity of <i>Cryptosporidium</i> spp. in Captive Reptiles. <i>Applied and Environmental Microbiology</i> , 2004, 70, 891-899.	1.4	117
25	It's official – <i>Cryptosporidium</i> is a gregrarine: What are the implications for the water industry?. <i>Water Research</i> , 2016, 105, 305-313.	5.3	110
26	Blood, Bull Terriers and Babesiosis: further evidence for direct transmission of <i>Babesia gibsoni</i> in dogs. <i>Australian Veterinary Journal</i> , 2007, 85, 459-463.	0.5	107
27	Inhibition of the endosymbiont <i>Candidatus</i> <i>Mitochondria</i> during 16S rRNA gene profiling reveals potential pathogens in <i>Ixodes</i> ticks from Australia. <i>Parasites and Vectors</i> , 2015, 8, 345.	1.0	95
28	<i>Cryptosporidium</i> – An update with an emphasis on foodborne and waterborne transmission. <i>Research in Veterinary Science</i> , 2020, 132, 500-512.	0.9	95
29	Prevalence and molecular characterisation of <i>Cryptosporidium</i> and <i>Giardia</i> species in pre-weaned sheep in Australia. <i>Veterinary Parasitology</i> , 2009, 161, 19-24.	0.7	94
30	Molecular typing of <i>Giardia duodenalis</i> in humans in Queensland – first report of Assemblage E. <i>Parasitology</i> , 2017, 144, 1154-1161.	0.7	94
31	New Technologies for Detection of Enteric Parasites. <i>Trends in Parasitology</i> , 2017, 33, 532-546.	1.5	94
32	Genotypes of <i>Cryptosporidium</i> Species Infecting Fur-Bearing Mammals Differ from Those of Species Infecting Humans. <i>Applied and Environmental Microbiology</i> , 2004, 70, 7574-7577.	1.4	86
33	An Update on Zoonotic <i>Cryptosporidium</i> Species and Genotypes in Humans. <i>Animals</i> , 2021, 11, 3307.	1.0	84
34	<i>Cryptosporidium</i> GP60 genotypes from humans and domesticated animals in Australia, North America and Europe. <i>Experimental Parasitology</i> , 2008, 118, 118-121.	0.5	78
35	Development of an untargeted metabolomics method for the analysis of human faecal samples using <i>Cryptosporidium</i> -infected samples. <i>Molecular and Biochemical Parasitology</i> , 2012, 185, 145-150.	0.5	73
36	Zoonotic giardiasis: an update. <i>Parasitology Research</i> , 2021, 120, 4199-4218.	0.6	71

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37	Molecular characterization of <i>Cryptosporidium</i> and <i>Giardia</i> in pre-weaned calves in Western Australia and New South Wales. <i>Veterinary Parasitology</i> , 2011, 176, 145-150.	0.7	67
38	Evidence of <i>Cryptosporidium</i> transmission between cattle and humans in northern New South Wales. <i>Experimental Parasitology</i> , 2012, 130, 437-441.	0.5	67
39	Molecular epidemiology of giardiasis from a veterinary perspective. <i>Advances in Parasitology</i> , 2019, 106, 209-254.	1.4	66
40	Identification of zoonotic <i>Cryptosporidium</i> and <i>Giardia</i> genotypes infecting animals in Sydney's water catchments. <i>Experimental Parasitology</i> , 2011, 128, 138-144.	0.5	65
41	<i>Cryptosporidium huwi</i> n. sp. (Apicomplexa: Eimeriidae) from the guppy ( <i>Poecilia reticulata</i> ). <i>Experimental Parasitology</i> , 2015, 150, 31-35.	0.5	64
42	Novel <i>Borrelia</i> species detected in echidna ticks, <i>Bothriocroton concolor</i> , in Australia. <i>Parasites and Vectors</i> , 2016, 9, 339.	1.0	63
43	<i>Trypanosoma irwini</i> n. sp (Sarcomastigophora: Trypanosomatidae) from the koala ( <i>Phascolarctos cinereus</i> ). <i>Parasitology</i> , 2009, 136, 875-885.	0.7	60
44	Development of a Multilocus Sequence Tool for Typing <i>Cryptosporidium muris</i> and <i>Cryptosporidium andersoni</i> . <i>Journal of Clinical Microbiology</i> , 2011, 49, 34-41.	1.8	60
45	Identification of rare and novel <i>Cryptosporidium</i> GP60 subtypes in human isolates from Jordan. <i>Experimental Parasitology</i> , 2010, 125, 161-164.	0.5	59
46	Longitudinal multi-locus molecular characterisation of sporadic Australian human clinical cases of cryptosporidiosis from 2005 to 2008. <i>Experimental Parasitology</i> , 2010, 125, 348-356.	0.5	59
47	Bacterial Profiling Reveals Novel <i>Neohhrlichia</i> , <i>Ehrlichia</i> , and <i>Anaplasma</i> Species in Australian Human-Biting Ticks. <i>PLoS ONE</i> , 2015, 10, e0145449.	1.1	58
48	Evidence supporting zoonotic transmission of <i>Cryptosporidium</i> in rural New South Wales. <i>Experimental Parasitology</i> , 2008, 119, 192-195.	0.5	57
49	<i>Giardia</i> genotypes in pigs in Western Australia: Prevalence and association with diarrhea. <i>Experimental Parasitology</i> , 2009, 121, 381-383.	0.5	57
50	<i>Cryptosporidium</i> species in Australian wildlife and domestic animals. <i>Parasitology</i> , 2012, 139, 1673-1688.	0.7	56
51	Wastewater-based epidemiology surveillance and early detection of waterborne pathogens with a focus on SARS-CoV-2, <i>Cryptosporidium</i> and <i>Giardia</i> . <i>Parasitology Research</i> , 2021, 120, 4167-4188.	0.6	55
52	<i>Cryptosporidium fayeri</i> n. sp. (Apicomplexa: Cryptosporidiidae) from the Red Kangaroo ( <i>Macropus rufus</i> ). <i>Journal of Eukaryotic Microbiology</i> , 2008, 55, 22-26.	0.8	54
53	Prevalence of and management factors contributing to <i>Cryptosporidium</i> sp. infection in pre-weaned and post-weaned calves in Johor, Malaysia. <i>Experimental Parasitology</i> , 2011, 127, 534-538.	0.5	54
54	Longitudinal prevalence, oocyst shedding and molecular characterisation of <i>Cryptosporidium</i> species in sheep across four states in Australia. <i>Veterinary Parasitology</i> , 2014, 200, 50-58.	0.7	54

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55	Cryptosporidium species in sheep and goats from Papua New Guinea. <i>Experimental Parasitology</i> , 2014, 141, 134-137.	0.5	54
56	Cryptosporidium species and subtypes in animals inhabiting drinking water catchments in three states across Australia. <i>Water Research</i> , 2018, 134, 327-340.	5.3	54
57	Longitudinal investigation of protozoan parasites in meat lamb farms in southern Western Australia. <i>Preventive Veterinary Medicine</i> , 2011, 101, 192-203.	0.7	53
58	Phylogenetic analysis of <i>Cystoisospora</i> species at the rRNA ITS1 locus and development of a PCR-RFLP assay. <i>Experimental Parasitology</i> , 2008, 118, 592-595.	0.5	52
59	Molecular characterization of <i>Cryptosporidium</i> spp. from wild rats and mice from rural communities in the Philippines. <i>Infection, Genetics and Evolution</i> , 2013, 16, 5-12.	1.0	52
60	Prevalence of <i>Cryptosporidium</i> and <i>Giardia</i> species in animals in irrigation catchments in the southwest of Australia. <i>Experimental Parasitology</i> , 2008, 118, 596-599.	0.5	50
61	Molecular characterization of <i>Cryptosporidium</i> and <i>Giardia</i> in farmers and their ruminant livestock from the Coastal Savannah zone of Ghana. <i>Infection, Genetics and Evolution</i> , 2017, 55, 236-243.	1.0	50
62	Endemic, exotic and novel apicomplexan parasites detected during a national study of ticks from companion animals in Australia. <i>Parasites and Vectors</i> , 2018, 11, 197.	1.0	49
63	Molecular characterisation of <i>Cryptosporidium</i> outbreaks in Western and South Australia. <i>Experimental Parasitology</i> , 2010, 125, 325-328.	0.5	48
64	Identification of novel and zoonotic <i>Cryptosporidium</i> species in marine fish. <i>Veterinary Parasitology</i> , 2010, 168, 190-195.	0.7	48
65	Genetic characterization of <i>Cryptosporidium</i> in animal and human isolates from Jordan. <i>Veterinary Parasitology</i> , 2016, 228, 116-120.	0.7	48
66	Zoonotic <i>Cryptosporidium</i> Species in Animals Inhabiting Sydney Water Catchments. <i>PLoS ONE</i> , 2016, 11, e0168169.	1.1	47
67	Novel trypanosome <i>Trypanosoma gilletti</i> sp. (Euglenozoa: Trypanosomatidae) and the extension of the host range of <i>Trypanosoma copemani</i> to include the koala ( <i>Phascolarctos</i> ) Tj ETQq1 1 0.784314 rgBT7/Overlock 10 Tf		
68	Multi-locus analysis of <i>Giardia duodenalis</i> intra-Assemblage B substitution patterns in cloned culture isolates suggests sub-Assemblage B analyses will require multi-locus genotyping with conserved and variable genes. <i>International Journal for Parasitology</i> , 2011, 41, 495-503.	1.3	46
69	Prevalence of <i>Cryptosporidium</i> genotypes in pre and post-weaned pigs in Australia. <i>Experimental Parasitology</i> , 2008, 119, 418-421.	0.5	45
70	Profiling the diversity of <i>Cryptosporidium</i> species and genotypes in wastewater treatment plants in Australia using next generation sequencing. <i>Science of the Total Environment</i> , 2018, 644, 635-648.	3.9	45
71	Morphological and molecular characterization of <i>Trypanosoma copemani</i> n. sp. (Trypanosomatidae) isolated from Gilbert's potoroo ( <i>Potorous gilbertii</i> ) and quokka ( <i>Setonix brachyurus</i> ). <i>Parasitology</i> , 2009, 136, 783-792.	0.7	44
72	Evaluation of 16S next-generation sequencing of hypervariable region 4 in wastewater samples: An unsuitable approach for bacterial enteric pathogen identification. <i>Science of the Total Environment</i> , 2019, 670, 1111-1124.	3.9	44

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73	Genotypes of <i>Cryptosporidium</i> from Sydney water catchment areas. <i>Journal of Applied Microbiology</i> , 2005, 98, 1221-1229.	1.4	43
74	<i>Cryptosporidium varanii</i> takes precedence over <i>C. saurophilum</i> . <i>Experimental Parasitology</i> , 2008, 118, 434-437.	0.5	43
75	Phylogenetic characterisation of two novel Anaplasmataceae from Australian <i>Ixodes holocyclus</i> ticks: 'Candidatus <i>Neoehrlichia australis</i> ' and 'Candidatus <i>Neoehrlichia arcana</i> '. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016, 66, 4256-4261.	0.8	43
76	Molecular characterisation of <i>Cryptosporidium</i> and <i>Giardia</i> in cats ( <i>Felis catus</i> ) in Western Australia. <i>Experimental Parasitology</i> , 2015, 155, 13-18.	0.5	42
77	A survey of ticks (Acari: Ixodidae) of companion animals in Australia. <i>Parasites and Vectors</i> , 2016, 9, 207.	1.0	42
78	Molecular and Biological Characterization of a <i>Cryptosporidium molnari</i> -Like Isolate from a Guppy ( <i>Poecilia reticulata</i> ). <i>Applied and Environmental Microbiology</i> , 2004, 70, 3761-3765.	1.4	41
79	<i>Babesia gibsoni</i> : Detection during experimental infections and after combined atovaquone and azithromycin therapy. <i>Experimental Parasitology</i> , 2007, 117, 115-123.	0.5	41
80	Increased genetic diversity and prevalence of co-infection with <i>Trypanosoma</i> spp. in koalas ( <i>Phascolarctos cinereus</i> ) and their ticks identified using next-generation sequencing (NGS). <i>PLoS ONE</i> , 2017, 12, e0181279.	1.1	41
81	Identification of eukaryotic microorganisms with 18S rRNA next-generation sequencing in wastewater treatment plants, with a more targeted NGS approach required for <i>Cryptosporidium</i> detection. <i>Water Research</i> , 2019, 158, 301-312.	5.3	41
82	A New Species of <i>Cryptosporidium</i> (Apicomplexa: Cryptosporidiidae) from Eastern Grey Kangaroos ( <i>Macropus giganteus</i> ). <i>Journal of Parasitology</i> , 2008, 94, 1114-1117.	0.3	40
83	High prevalence <i>Giardia duodenalis</i> assemblage B and potentially zoonotic subtypes in sporadic human cases in Western Australia. <i>International Journal for Parasitology</i> , 2010, 40, 293-297.	1.3	40
84	Molecular characterization of 'Candidatus <i>Borrelia tachyglossi</i> ' (family Spirochaetaceae) in echidna ticks, <i>Bothriocroton concolor</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 1075-1080.	0.8	39
85	Identification of novel <i>Babesia</i> and <i>Theileria</i> genotypes in the endangered marsupials, the woylie ( <i>Bettongia penicillata ogilbyi</i> ) and boodie ( <i>Bettongia lesueur</i> ). <i>Experimental Parasitology</i> , 2012, 131, 25-30.	0.5	38
86	Next Generation Sequencing uncovers within-host differences in the genetic diversity of <i>Cryptosporidium</i> gp60 subtypes. <i>International Journal for Parasitology</i> , 2017, 47, 601-607.	1.3	38
87	Identification of zoonotic <i>Giardia</i> genotypes in marsupials in Australia. <i>Experimental Parasitology</i> , 2008, 120, 88-93.	0.5	37
88	The potential impact of native Australian trypanosome infections on the health of koalas ( <i>Phascolarctos cinereus</i> ). <i>Parasitology</i> , 2011, 138, 873-883.	0.7	37
89	Complete development and multiplication of <i>Cryptosporidium hominis</i> in cell-free culture. <i>Veterinary Parasitology</i> , 2010, 169, 29-36.	0.7	36
90	Identification of novel trypanosome genotypes in native Australian marsupials. <i>Veterinary Parasitology</i> , 2011, 183, 21-30.	0.7	36

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91	<i>Cryptosporidium homai</i> n. sp. (Apicomplexa: Cryptosporididae) from the guinea pig ( <i>Cavia porcellus</i> ). <i>Veterinary Parasitology</i> , 2017, 245, 92-101.	0.7	36
92	Identification of novel <i>Cryptosporidium</i> species in aquarium fish. <i>Veterinary Parasitology</i> , 2010, 174, 43-48.	0.7	35
93	Molecular characterization of <i>Eimeria</i> species in macropods. <i>Experimental Parasitology</i> , 2012, 132, 216-221.	0.5	35
94	Diversity of <i>Bartonella</i> species detected in arthropod vectors from animals in Australia. <i>Comparative Immunology, Microbiology and Infectious Diseases</i> , 2011, 34, 411-417.	0.7	34
95	Vector of <i>Trypanosoma copemani</i> identified as <i>Ixodes</i> sp.. <i>Parasitology</i> , 2011, 138, 866-872.	0.7	34
96	Development of a quantitative PCR (qPCR) for <i>Giardia</i> and analysis of the prevalence, cyst shedding and genotypes of <i>Giardia</i> present in sheep across four states in Australia. <i>Experimental Parasitology</i> , 2014, 137, 46-52.	0.5	32
97	Identification of polymorphic genes for use in assemblage B genotyping assays through comparative genomics of multiple assemblage B <i>Giardia duodenalis</i> isolates. <i>Molecular and Biochemical Parasitology</i> , 2015, 201, 1-4.	0.5	32
98	Comparison of Sanger and next generation sequencing performance for genotyping <i>Cryptosporidium</i> isolates at the 18S rRNA and actin loci. <i>Experimental Parasitology</i> , 2015, 151-152, 21-27.	0.5	32
99	Detection and phylogenetic characterisation of novel <i>Anaplasma</i> and <i>Ehrlichia</i> species in <i>Amblyomma triguttatum</i> subsp. from four allopatric populations in Australia. <i>Ticks and Tick-borne Diseases</i> , 2017, 8, 749-756.	1.1	32
100	Are molecular tools clarifying or confusing our understanding of the public health threat from zoonotic enteric protozoa in wildlife?. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2019, 9, 323-341.	0.6	32
101	The first finding of a natural infection of <i>Cryptosporidium muris</i> in a cat. <i>Veterinary Parasitology</i> , 2007, 144, 349-352.	0.7	31
102	Multiple <i>Cryptosporidium</i> genotypes detected in wild black rats ( <i>Rattus rattus</i> ) from northern Australia. <i>Experimental Parasitology</i> , 2012, 131, 404-412.	0.5	31
103	Longitudinal prevalence and faecal shedding of <i>Chlamydia pecorum</i> in sheep. <i>Veterinary Journal</i> , 2014, 201, 322-326.	0.6	31
104	<i>Isoospora serinuse</i> n. sp. (Apicomplexa: Eimeriidae) from a domestic canary ( <i>Serinus canaria forma</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 59-66.	0.5	31
105	Limiting swimming pool outbreaks of cryptosporidiosis – the roles of regulations, staff, patrons and research. <i>Journal of Water and Health</i> , 2017, 15, 1-16.	1.1	31
106	Prevalence, Molecular Identification, and Risk Factors for <i>Cryptosporidium</i> Infection in Edible Marine Fish: A Survey Across Sea Areas Surrounding France. <i>Frontiers in Microbiology</i> , 2019, 10, 1037.	1.5	31
107	Novel <i>Cryptosporidium</i> Genotype in Wild Australian Mice ( <i>Mus domesticus</i> ). <i>Applied and Environmental Microbiology</i> , 2007, 73, 7693-7696.	1.4	30
108	Specific and genotypic identification of <i>Cryptosporidium</i> from a broad range of host species by nonisotopic SSCP analysis of nuclear ribosomal DNA. <i>Electrophoresis</i> , 2007, 28, 2818-2825.	1.3	29



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109	Identification of zoonotic Giardia genotypes in fish. International Journal for Parasitology, 2010, 40, 779-785.	1.3	29
110	Genetic diversity of Cryptosporidium in fish at the 18S and actin loci and high levels of mixed infections. Veterinary Parasitology, 2015, 214, 255-263.	0.7	29
111	Gastrointestinal helminths in farmers and their ruminant livestock from the Coastal Savannah zone of Ghana. Parasitology Research, 2018, 117, 3183-3194.	0.6	29
112	Genome-wide analysis of Borrelia turcica and Candidatus Borrelia taylori shows relapsing fever-like genomes with unique genomic links to Lyme disease Borrelia. Infection, Genetics and Evolution, 2018, 66, 72-81.	1.0	28
113	Retrospective analysis of Cryptosporidium species in Western Australian human populations (2015-2018), and emergence of the C. hominis fA12G1R5 subtype. Infection, Genetics and Evolution, 2019, 73, 306-313.	1.0	28
114	Small ruminants and zoonotic cryptosporidiosis. Parasitology Research, 2021, 120, 4189-4198.	0.6	28
115	Identification of Anisakis species (Nematoda: Anisakidae) in marine fish hosts from Papua New Guinea. Veterinary Parasitology, 2013, 193, 126-133.	0.7	27
116	Human Cryptosporidiosis Diagnosed in Western Australia: a Mixed Infection with Cryptosporidium meleagridis, the Cryptosporidium Mink Genotype, and an Unknown Cryptosporidium Species. Journal of Clinical Microbiology, 2013, 51, 2463-2465.	1.8	27
117	Polyphasic identification of cyanobacterial isolates from Australia. Water Research, 2014, 59, 248-261.	5.3	27
118	Cryptosporidium abrahamseni n. sp. (Apicomplexa: Cryptosporidiidae) from red-eye tetra (Moenkhausia tj ETQq0 0 0 r gBT /Overlock 10	0.5	27
119	Comparison of molecular and McMaster microscopy techniques to confirm the presence of naturally acquired strongylid nematode infections in sheep. Molecular and Biochemical Parasitology, 2011, 180, 62-67.	0.5	26
120	Candidatus Bartonella antechini: A novel Bartonella species detected in fleas and ticks from the yellow-footed antechinus (Antechinus flavipes), an Australian marsupial. Veterinary Microbiology, 2011, 149, 517-521.	0.8	26
121	Novel genotypes of Trypanosoma binneyi from wild platypuses (Ornithorhynchus anatinus) and identification of a leech as a potential vector. Experimental Parasitology, 2014, 145, 42-50.	0.5	26
122	Dynamics of Th17 associating cytokines in Cryptosporidium parvum-infected mice. Parasitology Research, 2016, 115, 879-887.	0.6	26
123	Prevalence and risk factors associated with gastrointestinal parasites in ruminant livestock in the Coastal Savannah zone of Ghana. Acta Tropica, 2019, 199, 105126.	0.9	26
124	Identification of novel Cryptosporidium genotypes in kangaroos from Western Australia. Veterinary Parasitology, 2011, 179, 22-27.	0.7	25
125	Novel Eimeria sp. isolated from a King's skink (Egernia kingii) in Western Australia. Experimental Parasitology, 2013, 133, 162-165.	0.5	25
126	Greater intensity and frequency of Cryptosporidium and Giardia oocyst shedding beyond the neonatal period is associated with reductions in growth, carcass weight and dressing efficiency in sheep. Veterinary Parasitology, 2016, 228, 42-51.	0.7	25



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127	<i>Cryptosporidium bollandi</i> n. sp. (Apicomplexa: Cryptosporididae) from angelfish ( <i>Pterophyllum scalare</i> ) and Oscar fish ( <i>Astronotus ocellatus</i> ). <i>Experimental Parasitology</i> , 2020, 217, 107956.	0.5	25
128	Comparison of various staining methods for the detection of <i>Cryptosporidium</i> in cell-free culture. <i>Experimental Parasitology</i> , 2008, 120, 67-72.	0.5	24
129	The molecular characterization of an <i>Eimeria</i> and <i>Cryptosporidium</i> detected in Asian seabass ( <i>Lates</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 0.7 24	0.7	24
130	Impacts of naturally acquired protozoa and strongylid nematode infections on growth and faecal attributes in lambs. <i>Veterinary Parasitology</i> , 2012, 184, 298-308.	0.7	24
131	Metabolomic Profiling of Faecal Extracts from <i>Cryptosporidium parvum</i> Infection in Experimental Mouse Models. <i>PLoS ONE</i> , 2013, 8, e77803.	1.1	24
132	Investigation of a swimming pool-associated cryptosporidiosis outbreak in the Kimberley region of Western Australia. <i>Epidemiology and Infection</i> , 2015, 143, 1037-1041.	1.0	24
133	Taxonomy and Molecular Taxonomy. , 2014, , 3-41.		24
134	Emergence of zoonotic <i>Cryptosporidium parvum</i> in China. <i>Trends in Parasitology</i> , 2022, 38, 335-343.	1.5	24
135	<i>Trypanosoma teixeirae</i> : A new species belonging to the <i>T. cruzi</i> clade causing trypanosomosis in an Australian little red flying fox ( <i>Pteropus scapulatus</i> ). <i>Veterinary Parasitology</i> , 2016, 223, 214-221.	0.7	23
136	A novel <i>Ehrlichia</i> species in blood and <i>Ixodes ornithorhynchi</i> ticks from platypuses ( <i>Ornithorhynchus</i> ) Tj ETQq0 0 0 1.1 rgBT /Overlock 10 Tf 1.1 23	1.1	23
137	<i>Isospora anthochaerae</i> n. sp. (Apicomplexa: Eimeriidae) from a Red wattlebird ( <i>Anthochaera</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 0.5 22	0.5	22
138	Detection of <i>Dientamoeba fragilis</i> in animal faeces using species specific real time PCR assay. <i>Veterinary Parasitology</i> , 2016, 227, 42-47.	0.7	22
139	Organoids and Bioengineered Intestinal Models: Potential Solutions to the <i>Cryptosporidium</i> Culturing Dilemma. <i>Microorganisms</i> , 2020, 8, 715.	1.6	22
140	Transient transfection of <i>Cryptosporidium parvum</i> using green fluorescent protein (GFP) as a marker. <i>Molecular and Biochemical Parasitology</i> , 2009, 168, 143-148.	0.5	21
141	Differences in the occurrence and epidemiology of cryptosporidiosis in Aboriginal and non-Aboriginal people in Western Australia (2002 â 2012). <i>Infection, Genetics and Evolution</i> , 2017, 53, 100-106.	1.0	21
142	Prevalence of <i>Cryptosporidium</i> species and subtypes in paediatric oncology and non-oncology patients with diarrhoea in Jordan. <i>Infection, Genetics and Evolution</i> , 2017, 55, 127-130.	1.0	21
143	<i>Cryptosporidium</i> in fish: alternative sequencing approaches and analyses at multiple loci to resolve mixed infections. <i>Parasitology</i> , 2017, 144, 1811-1820.	0.7	21
144	Molecular typing of <i>Giardia duodenalis</i> in cattle, sheep and goats in an arid area of central Iran. <i>Infection, Genetics and Evolution</i> , 2019, 75, 104021.	1.0	21

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145	Zoonotic <i>Bartonella</i> Species in Fleas and Blood from Red Foxes in Australia. <i>Vector-Borne and Zoonotic Diseases</i> , 2011, 11, 1549-1553.	0.6	20
146	Effect of dung burial by the dung beetle <i>Bubas bison</i> on numbers and viability of <i>Cryptosporidium</i> oocysts in cattle dung. <i>Experimental Parasitology</i> , 2011, 129, 1-4.	0.5	20
147	Piroplasms of New Zealand seabirds. <i>Parasitology Research</i> , 2014, 113, 4407-4414.	0.6	20
148	Global selective sweep of a highly inbred genome of the cattle parasite <i>Neospora caninum</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 22764-22773.	3.3	20
149	Morphological and molecular characterisation of a mixed <i>Cryptosporidium muris</i> / <i>Cryptosporidium felis</i> infection in a cat. <i>Veterinary Parasitology</i> , 2011, 175, 160-164.	0.7	19
150	Molecular confirmation of the first autochthonous case of human babesiosis in Australia using a novel primer set for the beta-tubulin gene. <i>Experimental Parasitology</i> , 2014, 141, 93-97.	0.5	19
151	Morphological and molecular characterization of <i>Isospora manorinae</i> n. sp. in a yellow-throated miner ( <i>Manorina flavigula wayensis</i> ) (Gould, 1840). <i>Experimental Parasitology</i> , 2016, 163, 16-23.	0.5	19
152	Molecular characterisation of a disseminated <i>Cryptosporidium</i> infection in a Koi carp ( <i>Cyprinus</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 46	0.7	19
153	Sequence analyses at mitochondrial and nuclear loci reveal a novel <i>Theileria</i> sp. and aid in the phylogenetic resolution of piroplasms from Australian marsupials and ticks. <i>PLoS ONE</i> , 2019, 14, e0225822.	1.1	19
154	A review of the molecular epidemiology of <i>Cryptosporidium</i> spp. and <i>Giardia duodenalis</i> in the Middle East and North Africa (MENA) region. <i>Infection, Genetics and Evolution</i> , 2022, 98, 105212.	1.0	19
155	Prevalence of <i>Cryptosporidium</i> species in recreational versus non-recreational water sources. <i>Experimental Parasitology</i> , 2012, 131, 399-403.	0.5	18
156	A new <i>Caryospora</i> coccidian species (Apicomplexa: Eimeriidae) from the laughing kookaburra ( <i>Dacelo</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 T	0.5	18
157	Longitudinal prevalence, oocyst shedding and molecular characterisation of <i>Eimeria</i> species in sheep across four states in Australia. <i>Experimental Parasitology</i> , 2014, 145, 14-21.	0.5	18
158	First Molecular Characterization of <i>Theileria ornithorhynchi</i> Mackerras, 1959: yet Another Challenge to the Systematics of the Piroplasms. <i>Protist</i> , 2015, 166, 609-620.	0.6	18
159	Prevalence, genetic diversity and potential clinical impact of blood-borne and enteric protozoan parasites in native mammals from northern Australia. <i>Veterinary Parasitology</i> , 2017, 238, 94-105.	0.7	18
160	Response to the newly proposed species <i>Cryptosporidium pestis</i> . <i>Trends in Parasitology</i> , 2007, 23, 41-42.	1.5	17
161	Molecular Characterization of <i>Babesia kiwiensis</i> From the Brown Kiwi ( <i>Apteryx mantelli</i> ). <i>Journal of Parasitology</i> , 2008, 94, 557-560.	0.3	17
162	Genetic characterization of flea-derived <i>Bartonella</i> species from native animals in Australia suggests hostâ€“parasite co-evolution. <i>Infection, Genetics and Evolution</i> , 2011, 11, 1868-1872.	1.0	17

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163	Infection levels of gastrointestinal parasites in sheep and goats in Papua New Guinea. <i>Journal of Helminthology</i> , 2013, 87, 409-415.	0.4	17
164	Morphological and molecular characterization of an uninucleated cyst-producing <i>Entamoeba</i> spp. in captured Rangeland goats in Western Australia. <i>Veterinary Parasitology</i> , 2017, 235, 41-46.	0.7	17
165	<i>Theileria gilberti</i> n. sp. (Apicomplexa: Theileriidae) in the Gilbert's Potoroo ( <i>Potorous</i> ) Tj ETQq1 1 0.784314 rgBT /Oyerlock 10	0.8	16
166	Cryptosporidium infection is associated with reduced growth and diarrhoea in goats beyond weaning. <i>Veterinary Parasitology</i> , 2018, 260, 30-37.	0.7	16
167	Characterization of INS-15, A Metalloprotease Potentially Involved in the Invasion of <i>Cryptosporidium parvum</i> . <i>Microorganisms</i> , 2019, 7, 452.	1.6	16
168	Prevalence and genotyping identification of <i>Cryptosporidium</i> in adult ruminants in central Iran. <i>Parasites and Vectors</i> , 2019, 12, 510.	1.0	16
169	Prevalence of <i>Giardia</i> spp. infection in pre-weaned and weaned calves in relation to management factors. <i>Veterinary Journal</i> , 2012, 191, 135-137.	0.6	15
170	Longitudinal prevalence, faecal shedding and molecular characterisation of <i>Campylobacter</i> spp. and <i>Salmonella enterica</i> in sheep. <i>Veterinary Journal</i> , 2014, 202, 250-254.	0.6	15
171	Target validation of the inosine monophosphate dehydrogenase (IMPDH) gene in <i>Cryptosporidium</i> using Phylomer® peptides. <i>Experimental Parasitology</i> , 2015, 148, 40-48.	0.5	15
172	Molecular investigation into the presence of a <i>Coxiella</i> sp. in <i>Rhipicephalus sanguineus</i> ticks in Australia. <i>Veterinary Microbiology</i> , 2017, 201, 141-145.	0.8	15
173	Morphological and molecular characterization of three <i>Eimeria</i> species from captured rangeland goats in Western Australia. <i>Veterinary Parasitology: Regional Studies and Reports</i> , 2017, 9, 75-83.	0.3	15
174	Morphological and molecular characterization of <i>Eimeria paludosa</i> coccidian parasite (Apicomplexa:Eimeriidae) in a dusky moorhen ( <i>Gallinula tenebrosa</i> , Gould, 1846) in Australia. <i>Experimental Parasitology</i> , 2014, 147, 16-22.	0.5	14
175	High prevalence of <i>Trypanosoma vegrandis</i> in bats from Western Australia. <i>Veterinary Parasitology</i> , 2015, 214, 342-347.	0.7	14
176	Morphological and molecular characterization of <i>Eimeria labbeana</i> -like (Apicomplexa:Eimeriidae) in a domestic pigeon ( <i>Columba livia domestica</i> , Gmelin, 1789) in Australia. <i>Experimental Parasitology</i> , 2016, 166, 124-130.	0.5	14
177	Trypanosomiasis in an Australian little red flying fox ( <i>Pteropus scapulatus</i> ). <i>Australian Veterinary Journal</i> , 2017, 95, 259-261.	0.5	14
178	Development of a modified molecular diagnostic procedure for the identification and quantification of naturally occurring strongylid larvae on pastures. <i>Veterinary Parasitology</i> , 2012, 190, 467-481.	0.7	13
179	Phylogenetic study of <i>Baylisascaris schroederi</i> isolated from Qinling subspecies of giant panda in China based on combined nuclear 5.8S and the second internal transcribed spacer (ITS-2) ribosomal DNA sequences. <i>Parasitology International</i> , 2012, 61, 497-500.	0.6	13
180	Prevalence and pathogen load of <i>Campylobacter</i> spp., <i>Salmonella enterica</i> and <i>Escherichia coli</i> O157/O145 serogroup in sheep faeces collected at sale yards and in abattoir effluent in Western Australia. <i>Australian Veterinary Journal</i> , 2017, 95, 143-148.	0.5	13

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181	First report of <i>Cryptosporidium parvum</i> in a dromedary camel calf from Western Australia. <i>Acta Parasitologica</i> , 2018, 63, 422-427.	0.4	13
182	Comparison of three cryptosporidiosis outbreaks in Western Australia: 2003, 2007 and 2011. <i>Epidemiology and Infection</i> , 2018, 146, 1413-1424.	1.0	13
183	Bacterial community profiling highlights complex diversity and novel organisms in wildlife ticks. <i>Ticks and Tick-borne Diseases</i> , 2020, 11, 101407.	1.1	13
184	Advances in molecular epidemiology of cryptosporidiosis in dogs and cats. <i>International Journal for Parasitology</i> , 2021, 51, 787-795.	1.3	13
185	Further characterisation of two <i>Eimeria</i> species ( <i>Eimeria quokka</i> and <i>Eimeria setonicis</i> ) in quokkas ( <i>Setonix brachyurus</i> ). <i>Experimental Parasitology</i> , 2014, 138, 48-54.	0.5	12
186	Morphological and molecular characterization of <i>Eimeria purpureicephali</i> n. sp. (Apicomplexa:Eimeriidae) in a red-capped parrot ( <i>Purpureicephalus spurius</i> , Kuhl, 1820) in Western Australia. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2016, 5, 34-39.	0.6	12
187	Molecular surveillance of piroplasms in ticks from small and medium-sized urban and peri-urban mammals in Australia. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2018, 7, 197-203.	0.6	12
188	Molecular analysis of cryptosporidiosis cases in Western Australia in 2019 and 2020 supports the occurrence of two swimming pool associated outbreaks and reveals the emergence of a rare <i>C. hominis</i> lbA12G3 subtype. <i>Infection, Genetics and Evolution</i> , 2021, 92, 104859.	1.0	12
189	Pyrrhocoricin as a potential drug delivery vehicle for <i>Cryptosporidium parvum</i> . <i>Experimental Parasitology</i> , 2008, 119, 301-303.	0.5	11
190	Molecular identification of naturally acquired strongylid infections in lambs—An investigation into how lamb age influences diagnostic sensitivity. <i>Veterinary Parasitology</i> , 2012, 187, 227-236.	0.7	11
191	<i>Isoospora strepera</i> n. sp. (Apicomplexa: Eimeriidae) from a grey currawong ( <i>Strepera versicolour</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 11 49-55.	0.5	11
192	Validation of cell-free culture using scanning electron microscopy (SEM) and gene expression studies. <i>Experimental Parasitology</i> , 2015, 153, 55-62.	0.5	11
193	Prevalence and pathogen load of <i>Cryptosporidium</i> and <i>Giardia</i> in sheep faeces collected from saleyards and in abattoir effluent in Western Australia. <i>Small Ruminant Research</i> , 2015, 130, 216-220.	0.6	11
194	Morphological and molecular characterization of <i>Isoospora neochmia</i> n. sp. in a captive-bred red-browed finch ( <i>Neochmia temporalis</i> ) (Latham, 1802). <i>Experimental Parasitology</i> , 2016, 166, 181-188.	0.5	11
195	Molecular identification of the <i>Trypanosoma</i> ( <i>Herpetosoma</i> ) <i>lewis</i> clade in black rats ( <i>Rattus rattus</i> ) from Australia. <i>Parasitology Research</i> , 2020, 119, 1691-1696.	0.6	11
196	Identification and differentiation of <i>Cryptosporidium</i> species by capillary electrophoresis single-strand conformation polymorphism. <i>FEMS Microbiology Letters</i> , 2011, 314, 34-41.	0.7	10
197	Investigation of the morphological diversity of the potentially zoonotic <i>Trypanosoma copemani</i> in quokkas and Gilbert's potoroos. <i>Parasitology</i> , 2015, 142, 1443-1452.	0.7	10
198	First genetic characterisation of <i>Giardia</i> in human isolates from Jordan. <i>Parasitology Research</i> , 2016, 115, 3723-3729.	0.6	10

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199	Chronic <i>Cystoisospora belli</i> infection in an immunocompetent Myanmar refugee – microscopy is not sensitive enough. <i>BMC Infectious Diseases</i> , 2016, 16, 221.	1.3	10
200	First report of <i>Trypanosoma vegrandis</i> in koalas ( <i>Phascolarctos cinereus</i> ). <i>Parasitology International</i> , 2016, 65, 316-318.	0.6	10
201	Comparison of ELISA, nested PCR and sequencing and a novel qPCR for detection of <i>Giardia</i> isolates from Jordan. <i>Experimental Parasitology</i> , 2018, 185, 23-28.	0.5	10
202	An Australian dog diagnosed with an exotic tick-borne infection: should Australia still be considered free from <i>Hepatozoon canis</i> ?. <i>International Journal for Parasitology</i> , 2018, 48, 805-815.	1.3	10
203	First report of <i>Trypanosoma dionisii</i> ( <i>Trypanosomatidae</i> ) identified in Australia. <i>Parasitology</i> , 2020, 147, 1801-1809.	0.7	10
204	<i>Cryptosporidium</i> and <i>Giardia</i> in dam water on sheep farms – An important source of transmission?. <i>Veterinary Parasitology</i> , 2020, 288, 109281.	0.7	10
205	Molecular Characterization of Novel <i>Cryptosporidium</i> Fish Genotypes in Edible Marine Fish. <i>Microorganisms</i> , 2020, 8, 2014.	1.6	10
206	Molecular identification and prevalence of <i>Isoospora</i> sp. in pigs in Western Australia using a PCR-RFLP assay. <i>Experimental Parasitology</i> , 2008, 120, 191-193.	0.5	9
207	A high prevalence of <i>Theileria penicillata</i> in woylies ( <i>Bettongia penicillata</i> ). <i>Experimental Parasitology</i> , 2012, 131, 157-161.	0.5	9
208	First report of <i>Cryptosporidium</i> species in farmed and wild buffalo from the Northern Territory, Australia. <i>Parasitology Research</i> , 2016, 115, 1349-1353.	0.6	9
209	Zoonotic <i>Cryptosporidium</i> and <i>Giardia</i> shedding by captured rangeland goats. <i>Veterinary Parasitology: Regional Studies and Reports</i> , 2017, 7, 32-35.	0.3	9
210	Illuminating the bacterial microbiome of Australian ticks with 16S and <i>Rickettsia</i> -specific next-generation sequencing. <i>Current Research in Parasitology and Vector-borne Diseases</i> , 2021, 1, 100037.	0.7	9
211	<i>Cryptosporidium tyzzeri</i> and <i>Cryptosporidium pestis</i> : Which name is valid?. <i>Experimental Parasitology</i> , 2012, 130, 308-309.	0.5	8
212	Characterization of two complete <i>Isoospora</i> mitochondrial genomes from passerine birds: <i>Isoospora serinuse</i> in a domestic canary and <i>Isoospora manorinae</i> in a yellow-throated miner. <i>Veterinary Parasitology</i> , 2017, 237, 137-142.	0.7	8
213	Novel Primer Sets for Next Generation Sequencing-Based Analyses of Water Quality. <i>PLoS ONE</i> , 2017, 12, e0170008.	1.1	8
214	<i>Salmonella enterica</i> isolates from Western Australian rangeland goats remain susceptible to critically important antimicrobials. <i>Scientific Reports</i> , 2018, 8, 15326.	1.6	8
215	Blood Parasites in Endangered Wildlife-Trypanosomes Discovered during a Survey of Haemoprotozoa from the Tasmanian Devil. <i>Pathogens</i> , 2020, 9, 873.	1.2	8
216	Haemoprotozoan surveillance in peri-urban native and introduced wildlife from Australia. <i>Current Research in Parasitology and Vector-borne Diseases</i> , 2021, 1, 100052.	0.7	8

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217	The bacterial biome of ticks and their wildlife hosts at the urban-wildland interface. <i>Microbial Genomics</i> , 2021, 7, .	1.0	8
218	Longitudinal analysis of <i>Giardia duodenalis</i> assemblages in animals inhabiting drinking water catchments in New South Wales and Queensland Australia (2013-2015). <i>Science of the Total Environment</i> , 2020, 718, 137433.	3.9	7
219	Zoonotic infection by <i>Cryptosporidium fayeri</i> IVgA10G1T1R1 in a Western Australian human. <i>Zoonoses and Public Health</i> , 2021, 68, 358-360.	0.9	7
220	Review of generic screening level assumptions for quantitative microbial risk assessment (QMRA) for estimating public health risks from Australian drinking water sources contaminated with <i>Cryptosporidium</i> by recreational activities. <i>Water Research</i> , 2022, 220, 118659.	5.3	7
221	In vitro analysis of the TAT protein transduction domain as a drug delivery vehicle in protozoan parasites. <i>Experimental Parasitology</i> , 2008, 118, 303-307.	0.5	6
222	<i>Eimeria tiliquae</i> n. sp. (Apicomplexa: Eimeriidae) from the shingleback skink ( <i>Tiliqua rugosa rugosa</i> ). <i>Experimental Parasitology</i> , 2013, 133, 144-149.	0.5	6
223	<i>Eimeria collieie</i> n. sp. (Apicomplexa:Eimeriidae) from the western long-necked turtle ( <i>Chelodina</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	0.5	6
224	Morphological and molecular characterization of <i>Eimeria haematodi</i> , coccidian parasite (Apicomplexa: Eimeriidae) in a rainbow lorikeet ( <i>Trichoglossus haematodus</i> ). <i>Experimental Parasitology</i> , 2015, 153, 123-128.	0.5	6
225	Morphological and molecular characterization of <i>Choleoimeria pogonae</i> n. sp. coccidian parasite (Apicomplexa: Eimeriidae, 1989, Paperna and Landsberg) in a western bearded dragon ( <i>Pogona minor</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	0.5	6
226	Identification of <i>Theileria fuliginosa</i> -like species in <i>Ixodes australiensis</i> ticks from western grey kangaroos ( <i>Macropus fuliginosus</i> ) in Western Australia. <i>Ticks and Tick-borne Diseases</i> , 2018, 9, 632-637.	1.1	6
227	Morphological and molecular characterisation of <i>Isospora butcheriae</i> n. sp. in a silvereye ( <i>Zosterops</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	0.6	6
228	<i>Isospora coronoideae</i> n. sp. (Apicomplexa: Eimeriidae) from the Australian raven ( <i>Corvus coronoides</i> ) (Passeriformes: Corvidae) (Linnaeus, 1758) in Western Australia. <i>Parasitology Research</i> , 2019, 118, 2399-2408.	0.6	6
229	Response to the Letter to the Editor by Harris. <i>Parasites and Vectors</i> , 2019, 12, 178.	1.0	6
230	Association of Common Zoonotic Pathogens With Concentrated Animal Feeding Operations. <i>Frontiers in Microbiology</i> , 2021, 12, 810142.	1.5	6
231	Further characterisation of <i>Leucocytozoon podargii</i> in wild tawny frogmouths ( <i>Podargus</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	0.6	5
232	Genetic characterizations of <i>Cryptosporidium</i> spp. from pet rodents indicate high zoonotic potential of pathogens from chinchillas. <i>One Health</i> , 2021, 13, 100269.	1.5	5
233	<i>Molecular Epidemiology</i> , 2007, , 119-171.		4
234	The innate resistance of <i>Trypanosoma copemani</i> to human serum. <i>Experimental Parasitology</i> , 2015, 153, 105-110.	0.5	4



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235	Prevalence, faecal shedding and genetic characterisation of <i>Yersinia</i> spp. in sheep across four states of Australia. <i>Australian Veterinary Journal</i> , 2016, 94, 129-137.	0.5	4
236	Molecular characterization of native Australian trypanosomes in quokka ( <i>Setonix brachyurus</i> ) populations from Western Australia. <i>Parasitology International</i> , 2016, 65, 205-208.	0.6	4
237	Detection of <i>Chlamydia pecorum</i> in joints trimmed from ovine carcasses with arthritis at an abattoir in southern Australia. <i>Small Ruminant Research</i> , 2017, 150, 80-86.	0.6	4
238	Molecular characterisation of <i>Salmonella enterica</i> serovar Typhimurium and <i>Campylobacter jejuni</i> faecal carriage by captured rangeland goats. <i>Small Ruminant Research</i> , 2018, 158, 48-53.	0.6	4
239	Faecal shedding of pathogenic <i>Yersinia enterocolitica</i> determined by qPCR for <i>yst</i> virulence gene is associated with reduced live weight but not diarrhoea in prime lambs. <i>Preventive Veterinary Medicine</i> , 2018, 152, 56-64.	0.7	4
240	Morphological and molecular characterisation of <i>Eimeria</i> vison-like oocysts (Apicomplexa:Eimeriidae) in farmed mink ( <i>Neovison vison</i> ) in Denmark. <i>Parasitology Research</i> , 2018, 117, 2933-2939.	0.6	4
241	Molecular and morphological analysis of a <i>Caryospora</i> -like isolate (Apicomplexa: Eimeriidae) from the magpie-lark ( <i>Grallina cyanoleuca</i> ) (Latham, 1801) in Western Australia. <i>Parasitology Research</i> , 2020, 119, 611-621.	0.6	4
242	Further characterisation of <i>Haemocystidium chelodinae</i> -like Haemoproteidae isolated from the Bellinger River snapping turtle ( <i>Myuchelys georgesi</i> ). <i>Parasitology Research</i> , 2020, 119, 601-609.	0.6	4
243	Morphological and molecular description of a new species of <i>Isospora</i> (Apicomplexa) from a New Holland honeyeater ( <i>Phylidonyris novaehollandiae</i> ). <i>Parasitology International</i> , 2021, 83, 102348.	0.6	4
244	Morphological and genetic characterization of the first <i>Isospora</i> species ( <i>I. lugensae</i> n. sp.) from a Kerguelen petrel ( <i>Lugensa brevirostris</i> ). <i>Parasitology Research</i> , 2021, 120, 1037-1047.	0.6	3
245	Chemical removal in waste stabilisation pond systems of varying configuration. <i>Environmental Science: Water Research and Technology</i> , 2021, 7, 1587-1599.	1.2	3
246	Morphological and genetic characterization of <i>Eimeria chalcopterae</i> n. sp. (Apicomplexa: Eimeriidae) in a common bronzewing pigeon ( <i>Phaps chalcoptera</i> ) (Latham, 1790) in Western Australia. <i>Parasitology Research</i> , 2020, 119, 3729-3737.	0.6	1
247	Mass mortality of the black sea hare <i>Aplysia gigantea</i> (Gastropoda: Aplysiidae) at Augusta, Western Australia in the austral summer of 2021. <i>Molluscan Research</i> , 2021, 41, 269-273.	0.2	1
248	Knowledge, Attitude and Practices Towards <i>Cryptosporidium</i> Among Public Swimming Pool Patrons and Staff in Western Australia. <i>Acta Parasitologica</i> , 2021, , 1.	0.4	1
249	Molecular characterization of and species in stool samples collected from Jordanian patients suffering from gastroenteritis. <i>Tropical Parasitology</i> , 2021, 11, 122-125.	0.2	1
250	Editorial. <i>Parasitology Research</i> , 2018, 117, 1997-1997.	0.6	0
251	A simple method to test the reproducibility of the phylogenetic reconstructions: the molecular systematics of cyanobacteria as a case study. <i>Fottea</i> , 2016, 16, 209-217.	0.4	0
252	<i>Cryptosporidium</i> . , 2018, , 551-563.		0