Paul T Monis

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

 120
 6,240
 43
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 papers
 6,853
 5.9
 5.81

 ext. papers
 ext. citations
 avg, IF
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#	Paper	IF	Citations
120	Discrimination of all genotypes of Giardia duodenalis at the glutamate dehydrogenase locus using PCR-RFLP. <i>Infection, Genetics and Evolution</i> , 2004 , 4, 125-30	4.5	381
119	Comparison of SYTO9 and SYBR Green I for real-time polymerase chain reaction and investigation of the effect of dye concentration on amplification and DNA melting curve analysis. <i>Analytical Biochemistry</i> , 2005 , 340, 24-34	3.1	225
118	Molecular systematics of the parasitic protozoan Giardia intestinalis. <i>Molecular Biology and Evolution</i> , 1999 , 16, 1135-44	8.3	225
117	Epidemiological and molecular evidence supports the zoonotic transmission of Giardia among humans and dogs living in the same community. <i>Parasitology</i> , 2004 , 128, 253-62	2.7	214
116	Genetic diversity within the morphological species Giardia intestinalis and its relationship to host origin. <i>Infection, Genetics and Evolution</i> , 2003 , 3, 29-38	4.5	205
115	Variation in Giardia: implications for taxonomy and epidemiology. <i>Advances in Parasitology</i> , 2004 , 58, 69-137	3.2	193
114	Variation in Giardia: towards a taxonomic revision of the genus. <i>Trends in Parasitology</i> , 2009 , 25, 93-100	6.4	189
113	Demonstration of preferential binding of SYBR Green I to specific DNA fragments in real-time multiplex PCR. <i>Nucleic Acids Research</i> , 2003 , 31, e136	20.1	181
112	Molecular and phylogenetic characterisation of Cryptosporidium from birds. <i>International Journal for Parasitology</i> , 2001 , 31, 289-96	4.3	161
111	Enumeration of water-borne bacteria using viability assays and flow cytometry: a comparison to culture-based techniques. <i>Journal of Microbiological Methods</i> , 2003 , 55, 585-97	2.8	160
110	Cryptosporidium and Giardia-zoonoses: fact or fiction?. <i>Infection, Genetics and Evolution</i> , 2003 , 3, 233-44	1 4.5	144
109	Critical processes affecting Cryptosporidium oocyst survival in the environment. <i>Parasitology</i> , 2007 , 134, 309-23	2.7	129
108	Cryptosporidium suis n. sp. (Apicomplexa: Cryptosporidiidae) in pigs (Sus scrofa). <i>Journal of Parasitology</i> , 2004 , 90, 769-73	0.9	124
107	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, 1105-13	4.3	123
106	Profiling bacterial survival through a water treatment process and subsequent distribution system. Journal of Applied Microbiology, 2005 , 99, 175-86	4.7	109
105	Novel lineages of Giardia intestinalis identified by genetic analysis of organisms isolated from dogs in Australia. <i>Parasitology</i> , 1998 , 116 (Pt 1), 7-19	2.7	102
104	Biochemistry and genetics of taste- and odor-producing cyanobacteria. <i>Harmful Algae</i> , 2016 , 54, 112-12	7 5.3	101

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103	Genetic analysis of Giardia from hoofed farm animals reveals artiodactyl-specific and potentially zoonotic genotypes. <i>Journal of Eukaryotic Microbiology</i> , 1997 , 44, 626-35	3.6	97
102	A comparative study of carboxyfluorescein diacetate and carboxyfluorescein diacetate succinimidyl ester as indicators of bacterial activity. <i>Journal of Microbiological Methods</i> , 2003 , 52, 379-88	2.8	96
101	It's official - Cryptosporidium is a gregarine: What are the implications for the water industry?. Water Research, 2016 , 105, 305-313	12.5	92
100	Isolation and characterization of the gene associated with geosmin production in cyanobacteria. <i>Environmental Science & Environmental Science & Envir</i>	10.3	92
99	Nucleic acid amplification-based techniques for pathogen detection and identification. <i>Infection, Genetics and Evolution</i> , 2006 , 6, 2-12	4.5	92
98	Cryptosporidiumbiotechnological advances in the detection, diagnosis and analysis of genetic variation. <i>Biotechnology Advances</i> , 2008 , 26, 304-17	17.8	91
97	Complete development of Cryptosporidium parvum in host cell-free culture. <i>International Journal for Parasitology</i> , 2004 , 34, 769-77	4.3	91
96	Metabolic flux network and analysis of fermentative hydrogen production. <i>Biotechnology Advances</i> , 2011 , 29, 375-87	17.8	85
95	Biosynthesis of 2-methylisoborneol in cyanobacteria. <i>Environmental Science & Environmental Science & </i>	10.3	79
94	Development and field testing of a real-time PCR assay for cylindrospermopsin-producing cyanobacteria. <i>Journal of Applied Microbiology</i> , 2008 , 104, 1503-15	4.7	79
93	Rapid, sensitive, and discriminating identification of Naegleria spp. by real-time PCR and melting-curve analysis. <i>Applied and Environmental Microbiology</i> , 2006 , 72, 5857-63	4.8	78
92	Environmental temperature controls Cryptosporidium oocyst metabolic rate and associated retention of infectivity. <i>Applied and Environmental Microbiology</i> , 2005 , 71, 3848-57	4.8	78
91	Giardiafrom genome to proteome. Advances in Parasitology, 2012, 78, 57-95	3.2	77
90	Culture-independent techniques for rapid detection of bacteria associated with loss of chloramine residual in a drinking water system. <i>Applied and Environmental Microbiology</i> , 2005 , 71, 6479-88	4.8	74
89	Phylogenetic Relationships among Isolates of Cryptosporidium: Evidence for Several New Species. Journal of Parasitology, 1999 , 85, 1126	0.9	71
88	Metabolic flux analysis of hydrogen production network by Clostridium butyricum W5: Effect of pH and glucose concentrations. <i>International Journal of Hydrogen Energy</i> , 2010 , 35, 6681-6690	6.7	64
87	Solar UV reduces Cryptosporidium parvum oocyst infectivity in environmental waters. <i>Journal of Applied Microbiology</i> , 2008 , 104, 1311-23	4.7	62
86	Cell culture-Taqman PCR assay for evaluation of Cryptosporidium parvum disinfection. <i>Applied and Environmental Microbiology</i> , 2003 , 69, 2505-11	4.8	62

85	Benthic cyanobacteria: A source of cylindrospermopsin and microcystin in Australian drinking water reservoirs. <i>Water Research</i> , 2017 , 124, 454-464	12.5	60
84	Molecular and phylogenetic analysis of Cryptosporidium muris from various hosts. <i>Parasitology</i> , 2000 , 120 (Pt 5), 457-64	2.7	55
83	Pathogen and Particle Associations in Wastewater: Significance and Implications for Treatment and Disinfection Processes. <i>Advances in Applied Microbiology</i> , 2016 , 97, 63-119	4.9	54
82	A molecular phylogeny of nuclear and mitochondrial sequences in Hymenolepis nana (Cestoda) supports the existence of a cryptic species. <i>Parasitology</i> , 2002 , 125, 567-75	2.7	53
81	Biodegradation of multiple cyanobacterial metabolites in drinking water supplies. <i>Chemosphere</i> , 2012 , 87, 1149-54	8.4	48
80	Emerging technologies for the detection and genetic characterization of protozoan parasites. <i>Trends in Parasitology</i> , 2005 , 21, 340-6	6.4	48
79	Genetic manipulation of butyrate formation pathways in Clostridium butyricum. <i>Journal of Biotechnology</i> , 2011 , 155, 269-74	3.7	46
78	Cooperative biodegradation of geosmin by a consortium comprising three gram-negative bacteria isolated from the biofilm of a sand filter column. <i>Letters in Applied Microbiology</i> , 2006 , 43, 417-23	2.9	44
77	Multi-locus analysis of Giardia duodenalis 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	4.3	43
76	The isolation and microbial community analysis of hydrogen producing bacteria from activated sludge. <i>Journal of Applied Microbiology</i> , 2007 , 103, 1415-23	4.7	43
75	Using Amplicon Sequencing To Characterize and Monitor Bacterial Diversity in Drinking Water Distribution Systems. <i>Applied and Environmental Microbiology</i> , 2015 , 81, 6463-73	4.8	42
74	Comparison of drinking water treatment process streams for optimal bacteriological water quality. <i>Water Research</i> , 2012 , 46, 3934-42	12.5	42
73	Cryptosporidium spp. in domestic dogs: the "dog" genotype. <i>Applied and Environmental Microbiology</i> , 2000 , 66, 2220-3	4.8	42
72	Zoonotic Cryptosporidium Species in Animals Inhabiting Sydney Water Catchments. <i>PLoS ONE</i> , 2016 , 11, e0168169	3.7	42
71	Use of DNA melting simulation software for in silico diagnostic assay design: targeting regions with complex melting curves and confirmation by real-time PCR using intercalating dyes. <i>BMC Bioinformatics</i> , 2007 , 8, 107	3.6	40
70	Cyst morphology and sequence analysis of the small subunit rDNA and ef1 alpha identifies a novel Giardia genotype in a quenda (Isoodon obesulus) from Western Australia. <i>Infection, Genetics and Evolution</i> , 2004 , 4, 365-70	4.5	40
69	A genetic and metabolic approach to redirection of biochemical pathways of Clostridium butyricum for enhancing hydrogen production. <i>Biotechnology and Bioengineering</i> , 2013 , 110, 338-42	4.9	38
68	Multiplication of the waterborne pathogen Cryptosporidium parvum in an aquatic biofilm system. <i>Parasites and Vectors</i> , 2013 , 6, 270	4	38

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67	Molecular epidemiology: a multidisciplinary approach to understanding parasitic zoonoses. <i>International Journal for Parasitology</i> , 2005 , 35, 1295-307	4.3	38	
66	Cryptosporidium species and subtypes in animals inhabiting drinking water catchments in three states across Australia. <i>Water Research</i> , 2018 , 134, 327-340	12.5	37	
65	Development of a nested-PCR assay for the detection of Cryptosporidium parvum in finished water. <i>Water Research</i> , 2001 , 35, 1641-8	12.5	37	
64	Isolates of £Candidatus Nostocoida limicolaSBlackall et al. 2000 should be described as three novel species of the genus Tetrasphaera, as Tetrasphaera jenkinsii sp. nov., Tetrasphaera vanveenii sp. nov. and Tetrasphaera veronensis sp. nov. <i>International Journal of Systematic and Evolutionary</i>	2.2	36	
63	Monitoring of geosmin producing Anabaena circinalis using quantitative PCR. <i>Water Research</i> , 2014 , 49, 416-25	12.5	34	
62	Biodegradation of geosmin by a novel Gram-negative bacterium; isolation, phylogenetic characterisation and degradation rate determination. <i>Water Research</i> , 2009 , 43, 2927-35	12.5	34	
61	Molecular characterisation of Cryptosporidium and Giardia in cats (Felis catus) in Western Australia. <i>Experimental Parasitology</i> , 2015 , 155, 13-8	2.1	31	
60	Complete development and multiplication of Cryptosporidium hominis in cell-free culture. <i>Veterinary Parasitology</i> , 2010 , 169, 29-36	2.8	30	
59	Assessing the impact of water treatment on bacterial biofilms in drinking water distribution systems using high-throughput DNA sequencing. <i>Chemosphere</i> , 2014 , 117, 185-92	8.4	29	
58	Molecular epidemiology: assumptions and limitations of commonly applied methods. <i>International Journal for Parasitology</i> , 1998 , 28, 981-7	4.3	28	
57	Profiling the diversity of Cryptosporidium species and genotypes in wastewater treatment plants in Australia using next generation sequencing. <i>Science of the Total Environment</i> , 2018 , 644, 635-648	10.2	27	
56	Extracellular excystation and development of Cryptosporidium: tracing the fate of oocysts within Pseudomonas aquatic biofilm systems. <i>BMC Microbiology</i> , 2014 , 14, 281	4.5	27	
55	Biochemical kinetics of fermentative hydrogen production by Clostridium butyricum W5. <i>International Journal of Hydrogen Energy</i> , 2009 , 34, 791-798	6.7	27	
54	Legionella confirmation using real-time PCR and SYTO9 is an alternative to current methodology. <i>Applied and Environmental Microbiology</i> , 2005 , 71, 8944-8	4.8	27	
53	Comparison of the levels of intra-specific genetic variation within Giardia muris and Giardia intestinalis. <i>International Journal for Parasitology</i> , 1998 , 28, 1179-85	4.3	26	
52	Molecular biology techniques in parasite ecology. International Journal for Parasitology, 2002, 32, 551-0	52 4.3	26	
51	Investigating source water Cryptosporidium concentration, species and infectivity rates during rainfall-runoff in a multi-use catchment. <i>Water Research</i> , 2014 , 67, 310-20	12.5	25	
50	Identification of polymorphic genes for use in assemblage B genotyping assays through comparative genomics of multiple assemblage B Giardia duodenalis isolates. <i>Molecular and Biochemical Parasitology</i> , 2015 , 201, 1-4	1.9	24	

49	Novel toxic effects associated with a tropical Limnothrix/Geitlerinema-like cyanobacterium. <i>Environmental Toxicology</i> , 2011 , 26, 260-70	4.2	24
48	Polyphasic identification of cyanobacterial isolates from Australia. <i>Water Research</i> , 2014 , 59, 248-61	12.5	22
47	Humans, dogs and parasitic zoonosesunravelling the relationships in a remote endemic community in northeast India using molecular tools. <i>Parasitology Research</i> , 2003 , 90 Suppl 3, S156-7	2.4	22
46	Understanding human infectious Cryptosporidium risk in drinking water supply catchments. <i>Water Research</i> , 2018 , 138, 282-292	12.5	21
45	Comparison of various staining methods for the detection of Cryptosporidium in cell-free culture. <i>Experimental Parasitology</i> , 2008 , 120, 67-72	2.1	21
44	The importance of systematics in parasitological research. <i>International Journal for Parasitology</i> , 1999 , 29, 381-8	4.3	20
43	Cryptosporidium cell culture infectivity assay design. <i>Parasitology</i> , 2011 , 138, 671-81	2.7	19
42	Wastewater-based epidemiology-surveillance and early detection of waterborne pathogens with a focus on SARS-CoV-2, Cryptosporidium and Giardia. <i>Parasitology Research</i> , 2021 , 120, 4167-4188	2.4	18
41	Cryptosporidium Attenuation across the Wastewater Treatment Train: Recycled Water Fit for Purpose. <i>Applied and Environmental Microbiology</i> , 2017 , 83,	4.8	17
40	EXPRESSION OF THE GEOSMIN SYNTHASE GENE IN THE CYANOBACTERIUM ANABAENA CIRCINALIS AWQC318(1). <i>Journal of Phycology</i> , 2011 , 47, 1338-43	3	17
39	Development and Evaluation of Three Real-Time PCR Assays for Genotyping and Source Tracking Cryptosporidium spp. in Water. <i>Applied and Environmental Microbiology</i> , 2015 , 81, 5845-54	4.8	15
38	Is nitrification the only cause of microbiologically induced chloramine decay?. <i>Water Research</i> , 2016 , 88, 904-911	12.5	15
37	Dissection of the hierarchy and synergism of the bile derived signal on Cryptosporidium parvum excystation and infectivity. <i>Parasitology</i> , 2012 , 139, 1533-46	2.7	15
36	Effect of water treatment processes on Cryptosporidium infectivity. Water Research, 2008, 42, 1805-11	12.5	14
35	Integrated cryptosporidium assay to determine oocyst density, infectivity, and genotype for risk assessment of source and reuse water. <i>Applied and Environmental Microbiology</i> , 2015 , 81, 3471-81	4.8	13
34	Virus removal by ultrafiltration: Understanding long-term performance change by application of Bayesian analysis. <i>Water Research</i> , 2017 , 122, 269-279	12.5	13
33	Organoids and Bioengineered Intestinal Models: Potential Solutions to the Culturing Dilemma. <i>Microorganisms</i> , 2020 , 8,	4.9	12
32	Toolbox for the sampling and monitoring of benthic cyanobacteria. <i>Water Research</i> , 2020 , 169, 115222	12.5	12

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31	Evaluation of chromogenic technologies for use in Australian potable water. <i>Journal of Applied Microbiology</i> , 2008 , 105, 1138-49	4.7	11	
30	Selection of surrogate pathogens and process indicator organisms for pasteurisation of municipal wastewater a survey of literature data on heat inactivation of pathogens. <i>Chemical Engineering Research and Design</i> , 2020 , 133, 301-314	5.5	11	
29	Virus removal of new and aged UF membranes at full-scale in a wastewater reclamation plant. <i>Environmental Science: Water Research and Technology</i> , 2016 , 2, 1014-1021	4.2	11	
28	DNA extraction from benthic Cyanobacteria: comparative assessment and optimization. <i>Journal of Applied Microbiology</i> , 2017 , 122, 294-304	4.7	10	
27	Do skin peptide profiles reflect speciation in the Australian treefrog Litoria caerulea (Anura : Hylidae)?. <i>Australian Journal of Zoology</i> , 2000 , 48, 33	0.5	10	
26	Epidemiological evaluation of sewage surveillance as a tool to detect the presence of COVID-19 cases in a low case load setting. <i>Science of the Total Environment</i> , 2021 , 786, 147469	10.2	10	
25	Risk-based management of drinking water safety in Australia: Implementation of health based targets to determine water treatment requirements and identification of pathogen surrogates for validation of conventional filtration. <i>Food and Waterborne Parasitology</i> , 2017 , 8-9, 64-74	6	9	
24	Detection and significance of the potentially pathogenic amoeboflagellate Naegleria italica in Australia. <i>Parasitology International</i> , 2004 , 53, 23-7	2.1	9	
23	Solar radiation induces non-nuclear perturbations and a false start to regulated exocytosis in Cryptosporidium parvum. <i>PLoS ONE</i> , 2010 , 5, e11773	3.7	9	
22	Independent validation and regulatory agency approval for high rate algal ponds to treat wastewater from rural communities. <i>Environmental Science: Water Research and Technology</i> , 2018 , 4, 195-205	4.2	8	
21	Flow cytometric assessment of distinct physiological stages within Cryptosporidium parvum sporozoites post-excystation. <i>Parasitology</i> , 2009 , 136, 953-66	2.7	7	
20	Novel Primer Sets for Next Generation Sequencing-Based Analyses of Water Quality. <i>PLoS ONE</i> , 2017 , 12, e0170008	3.7	6	
19	Disaggregation of colonies of Microcystis (Cyanobacteria): efficiency of two techniques assessed using an image analysis system. <i>Journal of Applied Phycology</i> , 2004 , 16, 117-125	3.2	5	
18	Inactivation, removal, and regrowth potential of opportunistic pathogens and antimicrobial resistance genes in recycled water systems. <i>Water Research</i> , 2021 , 201, 117324	12.5	5	
17	Taxonomy of Giardia Species 2011 , 3-15		4	
16	Disposable microfluidic micromixers for effective capture of oocysts from water samples. <i>Journal of Biological Engineering</i> , 2018 , 12, 4	6.3	3	
15	PCR slippage across the ML-2 microsatellite of the Cryptosporidium MIC1 locus enables development of a PCR assay capable of distinguishing the zoonotic Cryptosporidium parvum from other human infectious Cryptosporidium species. <i>Zoonoses and Public Health</i> , 2014 , 61, 324-37	2.9	3	
14	Validation of activated sludge plant performance for virus and protozoan reduction. <i>Journal of Water Reuse and Desalination</i> , 2013 , 3, 140-147	2.6	3	

13	Field based pilot-scale drinking water distribution system: Simulation of long hydraulic retention times and microbiological mediated monochloramine decay. <i>MethodsX</i> , 2018 , 5, 684-696	1.9	2
12	Evaluation of heterotrophic plate and chromogenic agar colony counting in water quality laboratories. <i>MethodsX</i> , 2015 , 2, 415-22	1.9	2
11	Evaluating membrane performance in recycled water treatment plants for assets replacement strategy. <i>Water Science and Technology</i> , 2017 , 76, 2941-2948	2.2	1
10	Removal of pathogens by functionalised self-assembled monolayers 2008 , 57, 93-100		1
9	Stormwater monitoring using on-line UV-Vis spectroscopy. <i>Environmental Science and Pollution Research</i> , 2021 , 1	5.1	1
8	Wastewater monitoring for SARS-CoV-2. <i>Microbiology Australia</i> , 2021 , 42, 18	0.8	1
7	Removal and Inactivation of Cryptosporidium from Water 2014 , 515-552		O
6	Sanitation: Thwart fatal infant gut parasite. <i>Nature</i> , 2014 , 507, 431	50.4	
5	Cryptosporidium and the Environment Dverview and Summary 2003, 387-392		
4	The Use of Cell Culture and Real-time PCR to Assess Disinfection of Cryptosporidium Parvum 2003 , 25	7-260	
3	Evaluation of Oocyst DNA Extraction Methods Using Real-time PCR 2003 , 177-180		
2	A Novel and Rapid Legionella Detection System for Water Analysis453-455		
1	Effectiveness and Energy Requirements of Pasteurisation for the Treatment of Unfiltered Secondary Effluent from a Municipal Wastewater Treatment Plant. <i>Water (Switzerland)</i> , 2020 , 12, 2100	3	