John T Ellis

List of Publications by Year
in descending order

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210 papers	9,365 citations	30070 54 h-index	53230 85 g-index
212 all docs	212 docs citations	212 times ranked	6177 citing authors

#	Article	IF	Citations
1	Host transmission dynamics of first- and third-stage Angiostrongylus cantonensis larvae in Bullastra lessoni. Parasitology, 2022, 149, 1034-1044.	1.5	5
2	Compilation of parasitic immunogenic proteins from 30Âyears of published research using machine learning and natural language processing. Scientific Reports, 2022, 12, .	3.3	2
3	Recent trends in the use of social media in parasitology and the application of alternative metrics. Current Research in Parasitology and Vector-borne Diseases, 2021, 1, 100013.	1.9	4
4	Plasmodium falciparum Histidine-Rich Protein 2 and 3 Gene Deletions in Strains from Nigeria, Sudan, and South Sudan. Emerging Infectious Diseases, 2021, 27, 471-479.	4.3	23
5	Machine learning and applications in microbiology. FEMS Microbiology Reviews, 2021, 45, .	8.6	81
6	A new subspecies of Trypanosoma cyclops found in the Australian terrestrial leech Chtonobdella bilineata. Parasitology, 2021, 148, 1125-1136.	1.5	9
7	Applying Machine Learning to Predict the Exportome of Bovine and Canine Babesia Species That Cause Babesiosis. Pathogens, 2021, 10, 660.	2.8	7
8	Predicting Protein Therapeutic Candidates for Bovine Babesiosis Using Secondary Structure Properties and Machine Learning. Frontiers in Genetics, 2021, 12, 716132.	2.3	4
9	The controversies surrounding Giardia intestinalis assemblages A and B. Current Research in Parasitology and Vector-borne Diseases, 2021, 1, 100055.	1.9	11
10	Computational Antigen Discovery for Eukaryotic Pathogens Using Vacceed. Methods in Molecular Biology, 2021, 2183, 29-42.	0.9	1
11	Diversity profiling of xenic cultures of Dientamoeba fragilis following systematic antibiotic treatment and prospects for genome sequencing. Parasitology, 2020, 147, 29-38.	1.5	0
12	Contribution of introns to the species diversity associated with the apicomplexan parasite, Neospora caninum. Parasitology Research, 2020, 119, 431-445.	1.6	2
13	Detecting sequence variants in clinically important protozoan parasites. International Journal for Parasitology, 2020, 50, 1-18.	3.1	2
14	30 years of parasitology research analysed by text mining. Parasitology, 2020, 147, 1643-1657.	1.5	7
15	Research into Neospora caninumâ€"What Have We Learnt in the Last Thirty Years?. Pathogens, 2020, 9, 505.	2.8	18
16	Species diversity and genome evolution of the pathogenic protozoan parasite, Neospora caninum. Infection, Genetics and Evolution, 2020, 84, 104444.	2.3	8
17	Molecular Detection of Antimalarial Drug Resistance in Plasmodium vivax from Returned Travellers to NSW, Australia during 2008â€"2018. Pathogens, 2020, 9, 101.	2.8	8
18	A review of the systematics, species identification and diagnostics of the Trypanosomatidae using the maxicircle kinetoplast DNA: from past to present. International Journal for Parasitology, 2020, 50, 449-460.	3.1	5

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19	Molecular detection of drug resistant malaria in Southern Thailand. Malaria Journal, 2019, 18, 275.	2.3	28
20	Semi-Quantitative, Duplexed qPCR Assay for the Detection of Leishmania spp. Using Bisulphite Conversion Technology. Tropical Medicine and Infectious Disease, 2019, 4, 135.	2.3	5
21	Evolutionary Insight into the Trypanosomatidae Using Alignment-Free Phylogenomics of the Kinetoplast. Pathogens, 2019, 8, 157.	2.8	8
22	Evaluation of the EasyScreen Protozoan Detection Kit for the diagnosis of Entamoeba histolytica. Pathology, 2019, 51, 426-428.	0.6	1
23	The complete coding region of the maxicircle as a superior phylogenetic marker for exploring evolutionary relationships between members of the Leishmaniinae. Infection, Genetics and Evolution, 2019, 70, 90-100.	2.3	18
24	Comparison and Recommendations for Use of Dientamoeba fragilis Real-Time PCR Assays. Journal of Clinical Microbiology, 2019, 57, .	3.9	13
25	Annotating the â€˜hypotheticalâ€™ in hypothetical proteins: In-silico analysis of uncharacterised proteins for the Apicomplexan parasite, Neospora caninum. Veterinary Parasitology, 2019, 265, 29-37.	1.8	4
26	Epidemiology and associated risk factors of giardiasis in a peri-urban setting in New South Wales Australia. Epidemiology and Infection, 2019, 147, e15.	2.1	7
27	Identification of Clinical Infections of Leishmania Imported into Australia: Revising Speciation with Polymerase Chain Reaction-RFLP of the Kinetoplast Maxicircle. American Journal of Tropical Medicine and Hygiene, 2019, 101, 590-601.	1.4	5
28	Evolutionary ARMS Race: Antimalarial Resistance Molecular Surveillance. Trends in Parasitology, 2018, 34, 322-334.	3.3	9
29	A Gene-Based Positive Selection Detection Approach to Identify Vaccine Candidates Using Toxoplasma gondii as a Test Case Protozoan Pathogen. Frontiers in Genetics, 2018, 9, 332.	2.3	17
30	Resistance screening and trend analysis of imported falciparum malaria in NSW, Australia (2010 to) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	2.5	13
31	Genome Wide Identification of Mutational Hotspots in the Apicomplexan Parasite Neospora caninum and the Implications for Virulence. Genome Biology and Evolution, 2018, 10, 2417-2431.	2.5	17
32	Geospatial Distribution of Giardiasis in NSW, Australia. ISEE Conference Abstracts, 2018, 2017, 206.	0.0	0
33	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
34	On the application of reverse vaccinology to parasitic diseases: a perspective on feature selection and ranking of vaccine candidates. International Journal for Parasitology, 2017, 47, 779-790.	3.1	13
35	Dynamic island model based on spectral clustering in genetic algorithm. , 2017, , .		10
36	Staged heterogeneity learning to identify conformational B-cell epitopes from antigen sequences. BMC Genomics, 2017, 18, 113.	2.8	5

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37	The evolution of trypanosomatid taxonomy. Parasites and Vectors, 2017, 10, 287.	2.5	123
38	Isolation of Novel Trypanosomatid, Zelonia australiensis sp. nov. (Kinetoplastida: Trypanosomatidae) Provides Support for a Gondwanan Origin of Dixenous Parasitism in the Leishmaniinae. PLoS Neglected Tropical Diseases, 2017, 11, e0005215.	3.0	55
39	Angiostrongylus cantonensis: a review of its distribution, molecular biology and clinical significance as a human pathogen. Parasitology, 2016, 143, 1087-1118.	1.5	162
40	Comparison of enteric protozoan infections in four Australian hospitals: variable tests and variable results. Parasitology Open, 2016, 2, .	0.9	1
41	Dientamoeba fragilis, the Neglected Trichomonad of the Human Bowel. Clinical Microbiology Reviews, 2016, 29, 553-580.	13.6	96
42	Detection of Dientamoeba fragilis in animal faeces using species specific real time PCR assay. Veterinary Parasitology, 2016, 227, 42-47.	1.8	22
43	Bulky Trichomonad Genomes: Encoding a Swiss Army Knife. Trends in Parasitology, 2016, 32, 783-797.	3.3	13
44	Positive-unlabeled learning for the prediction of conformational B-cell epitopes. BMC Bioinformatics, 2015, 16, S12.	2.6	22
45	A live vaccine against Neospora caninum abortions in cattle. Vaccine, 2015, 33, 1299-1301.	3.8	29
46	In Vitro Antimicrobial Susceptibility Patterns of Blastocystis. Antimicrobial Agents and Chemotherapy, 2015, 59, 4417-4423.	3.2	15
47	The Transcriptome Sequence of Dientamoeba fragilis Offers New Biological Insights on its Metabolism, Kinome, Degradome and Potential Mechanisms of Pathogenicity. Protist, 2015, 166, 389-408.	1.5	18
48	Improving the gene structure annotation of the apicomplexan parasite Neospora caninum fulfils a vital requirement towards an in silico-derived vaccine. International Journal for Parasitology, 2015, 45, 305-318.	3.1	11
49	Molecular Epidemiology of Imported Cases of Leishmaniasis in Australia from 2008 to 2014. PLoS ONE, 2015, 10, e0119212.	2.5	17
50	The Prevalence of Angiostrongylus cantonensis/mackerrasae Complex in Molluscs from the Sydney Region. PLoS ONE, 2015, 10, e0128128.	2.5	20
51	Recent Advances in Molecular Biology of Parasitic Viruses. Infectious Disorders - Drug Targets, 2015, 14, 155-167.	0.8	9
52	Descriptive epidemiology of infectious gastrointestinal illnesses in Sydney, Australia, 2007â€“2010. Western Pacific Surveillance and Response Journal: WPSAR, 2015, 6, 7-16.	0.6	15
53	Update on the Molecular Epidemiology and Diagnostic Tools for Blastocystis sp. , 2014, 03, .		0
54	Influenza A HA's conserved epitopes and broadly neutralizing antibodies: A prediction method. Journal of Bioinformatics and Computational Biology, 2014, 12, 1450023.	0.8	5

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55	Tertiary structure-based prediction of conformational B-cell epitopes through B factors. Bioinformatics, 2014, 30, i264-i273.	4.1	33
56	Vacceed: a high-throughput in silico vaccine candidate discovery pipeline for eukaryotic pathogens based on reverse vaccinology. Bioinformatics, 2014, 30, 2381-2383.	4.1	60
57	Enhancing In Silico Protein-Based Vaccine Discovery for Eukaryotic Pathogens Using Predicted Peptide-MHC Binding and Peptide Conservation Scores. PLoS ONE, 2014, 9, e115745.	2.5	25
58	Epidemiology and geographical distribution of enteric protozoan infections in Sydney, Australia. Journal of Public Health Research, 2014, 3, 298.	1.2	28
59	Control options forNeospora caninumâ€“ is there anything new or are we going backwards?. Parasitology, 2014, 141, 1455-1470.	1.5	43
60	Treatment failure in patients with chronic Blastocystis infection. Journal of Medical Microbiology, 2014, 63, 252-257.	1.8	37
61	Description of Dientamoeba fragilis Cyst and Precystic Forms from Human Samples. Journal of Clinical Microbiology, 2014, 52, 2680-2683.	3.9	53
62	Discovering a vaccine against neosporosis using computers: is it feasible?. Trends in Parasitology, 2014, 30, 401-411.	3.3	28
63	Update on the pathogenic potential and treatment options for Blastocystis sp. Gut Pathogens, 2014, 6, 17.	3.4	121
64	Evaluation of the EasyScreenâ„¢ Enteric Parasite Detection Kit for the detection of Blastocystis spp., Cryptosporidium spp., Dientamoeba fragilis, Entamoeba complex, and Giardia intestinalis from clinical stool samples. Diagnostic Microbiology and Infectious Disease, 2014, 78, 149-152.	1.8	42
65	Activity of benzimidazoles againstDientamoeba fragilis(Trichomonadida,) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 347 Tc resistance. Parasite, 2014, 21, 41.	2.0	4
66	What is the global economic impact of Neospora caninum in cattle â€“ The billion dollar question. International Journal for Parasitology, 2013, 43, 133-142.	3.1	381
67	A review of the infection, genetics, and evolution of Neospora caninum: From the past to the present. Infection, Genetics and Evolution, 2013, 13, 133-150.	2.3	111
68	A novel strategy for classifying the output from an in silicovaccine discovery pipeline for eukaryotic pathogens using machine learning algorithms. BMC Bioinformatics, 2013, 14, 315.	2.6	38
69	Cyst formation and faecalâ€“oral transmission of Dientamoeba fragilis â€“ the missing link in the life cycle of an emerging pathogen. International Journal for Parasitology, 2013, 43, 879-883.	3.1	58
70	A guide to in silico vaccine discovery for eukaryotic pathogens. Briefings in Bioinformatics, 2013, 14, 753-774.	6.5	29
71	Subtype distribution of Blastocystis isolates from a variety of animals from New South Wales, Australia. Veterinary Parasitology, 2013, 196, 85-89.	1.8	79
72	Gastrointestinal pathogen distribution in symptomatic children in Sydney, Australia. Journal of Epidemiology and Global Health, 2013, 3, 11.	2.9	19

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73	Subtype distribution of Blastocystis isolates identified in a Sydney population and pathogenic potential of Blastocystis. European Journal of Clinical Microbiology and Infectious Diseases, 2013, 32, 335-343.	2.9	58
74	On the Efficacy and Safety of Vaccination with Live Tachyzoites of Neospora caninum for Prevention of Neospora-Associated Fetal Loss in Cattle. Vaccine Journal, 2013, 20, 99-105.	3.1	46
75	Prevalence of Gastrointestinal Pathogens in Developed and Developing Countries: Systematic Review and Meta-Analysis. Journal of Public Health Research, 2013, 2, jphr.2013.e9.	1.2	111
76	Detection and Transmission of Dientamoeba fragilis from Environmental and Household Samples. American Journal of Tropical Medicine and Hygiene, 2012, 86, 233-236.	1.4	17
77	In Vitro Susceptibility Testing of Dientamoeba fragilis. Antimicrobial Agents and Chemotherapy, 2012, 56, 487-494.	3.2	18
78	The Core Mouse Response to Infection by Neospora Caninum Defined by Gene Set Enrichment Analyses. Bioinformatics and Biology Insights, 2012, 6, BBI.S9954.	2.0	3
79	New advances in thein-vitroculture ofDientamoeba fragilis. Parasitology, 2012, 139, 864-869.	1.5	9
80	Enteric Protozoa in the Developed World: a Public Health Perspective. Clinical Microbiology Reviews, 2012, 25, 420-449.	13.6	329
81	Current treatment options for Dientamoeba fragilis infections. International Journal for Parasitology: Drugs and Drug Resistance, 2012, 2, 204-215.	3.4	30
82	Evaluating High-Throughput Ab Initio Gene Finders to Discover Proteins Encoded in Eukaryotic Pathogen Genomes Missed by Laboratory Techniques. PLoS ONE, 2012, 7, e50609.	2.5	33
83	A microscopic description and ultrastructural characterisation of Dientamoeba fragilis: An emerging cause of human enteric disease. International Journal for Parasitology, 2012, 42, 139-153.	3.1	18
84	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
85	Comparison of Microscopy, Culture, and Conventional Polymerase Chain Reaction for Detection of Blastocystis sp. in Clinical Stool Samples. American Journal of Tropical Medicine and Hygiene, 2011, 84, 308-312.	1.4	102
86	Prevalence of gastrointestinal pathogens in Sub-Saharan Africa: systematic review and meta-analysis. Journal of Public Health in Africa, 2011, 2, 30.	0.4	42
87	The ambiguous life ofDientamoeba fragilis: the need to investigate current hypotheses on transmission. Parasitology, 2011, 138, 557-572.	1.5	38
88	A case-controlled study of Dientamoeba fragilis infections in children. Parasitology, 2011, 138, 819-823.	1.5	33
89	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
90	Extensive production of Neospora caninum tissue cysts in a carnivorous marsupial succumbing to experimental neosporosis. Veterinary Research, 2011, 42, 75.	3.0	18

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91	A review ofDientamoeba fragiliscarriage in humans: Several reasons why this organism should be considered in the diagnosis of gastrointestinal illness. Gut Microbes, 2011, 2, 3-12.	9.8	116
92	Evaluation of Multiplex Tandem Real-Time PCR for Detection of Cryptosporidium spp., Dientamoeba fragilis , Entamoeba histolytica , and Giardia intestinalis in Clinical Stool Samples. Journal of Clinical Microbiology, 2011, 49, 257-262.	3.9	114
93	Comparison of microscopy, two xenic culture techniques, conventional and real-time PCR for the detection of Dientamoeba fragilis in clinical stool samples. European Journal of Clinical Microbiology and Infectious Diseases, 2010, 29, 411-416.	2.9	70
94	Microarray analyses of mouse responses to infection by Neospora caninum identifies disease associated cellular pathways in the host response. Molecular and Biochemical Parasitology, 2010, 174, 117-127.	1.1	15
95	Australian dingoes are definitive hosts of Neospora caninum. International Journal for Parasitology, 2010, 40, 945-950.	3.1	188
96	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
97	On the Biological and Genetic Diversity in Neospora caninum. Diversity, 2010, 2, 411-438.	1.7	24
98	Importance of Nonenteric Protozoan Infections in Immunocompromised People. Clinical Microbiology Reviews, 2010, 23, 795-836.	13.6	89
99	Newly defined conditions for thein vitrocultivation and cryopreservation ofDientamoeba fragilis: new techniques set to fast track molecular studies on this organism. Parasitology, 2010, 137, 1867-1878.	1.5	25
100	A Review of the Clinical Presentation of Dientamoebiasis. American Journal of Tropical Medicine and Hygiene, 2010, 82, 614-619.	1.4	109
101	A second generation multiplex PCR for typing strains of Neospora caninum using six DNA targets. Molecular and Cellular Probes, 2010, 24, 20-26.	2.1	23
102	Repeated Dientamoeba fragilis Infections: A Case Report of Two Families from Sydney, Australia. Gastroenterology Insights, 2009, 1, e4.	1.2	8
103	Repeated Dientamoeba fragilis infections: a case report of two families from Sydney, Australia. Gastroenterology Insights, 2009, 1, e4.	1.2	7
104	Limited genetic diversity among genotypes of Enterocytozoon bieneusi strains isolated from HIV-infected patients from Sydney, Australia. Journal of Medical Microbiology, 2009, 58, 355-357.	1.8	35
105	A unique thioredoxin of the parasitic nematode Haemonchus contortus with glutaredoxin activity. Free Radical Biology and Medicine, 2009, 46, 579-585.	2.9	17
106	Neospora caninum â€“ How close are we to development of an efficacious vaccine that prevents abortion in cattle?. International Journal for Parasitology, 2009, 39, 1173-1187.	3.1	84
107	Isolation of Toxoplasma gondii from the brain of a dog in Australia and its biological and molecular characterization. Veterinary Parasitology, 2009, 164, 335-339.	1.8	25
108	Genetic diversity amongst isolates of Neospora caninum, and the development of a multiplex assay for the detection of distinct strains. Molecular and Cellular Probes, 2009, 23, 132-139.	2.1	36

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109	Clinical Significance of Enteric Protozoa in the Immunosuppressed Human Population. Clinical Microbiology Reviews, 2009, 22, 634-650.	13.6	187
110	Protozoal Hepatitis Associated with Immunosuppressive Therapy in a Dog. Journal of Veterinary Internal Medicine, 2009, 23, 366-368.	1.6	16
111	Entamoeba moshkovskii infections in Sydney, Australia. European Journal of Clinical Microbiology and Infectious Diseases, 2008, 27, 133-137.	2.9	61
112	Gorillas are a host for Dientamoeba fragilis: An update on the life cycle and host distribution. Veterinary Parasitology, 2008, 151, 21-26.	1.8	36
113	Re-evaluating the economics of neosporosis control. Veterinary Parasitology, 2008, 156, 361-362.	1.8	17
114	Thioredoxins of a parasitic nematode: Comparison of the 16- and 12-kDA thioredoxins from Haemonchus contortus. Free Radical Biology and Medicine, 2008, 44, 2026-2033.	2.9	13
115	The development and evaluation of a nested PCR assay for detection of Neospora caninum and Hammondia heydorni in feral mouse tissues. Molecular and Cellular Probes, 2008, 22, 228-233.	2.1	32
116	Evaluation of recombinant proteins of Neospora caninum as vaccine candidates (in a mouse model). Vaccine, 2008, 26, 5989-5996.	3.8	41
117	Comparison of Stool Antigen Detection Kits to PCR for Diagnosis of Amebiasis. Journal of Clinical Microbiology, 2008, 46, 1678-1681.	3.9	71
118	PCR Detection of Entamoeba histolytica , Entamoeba dispar , and Entamoeba moshkovskii in Stool Samples from Sydney, Australia. Journal of Clinical Microbiology, 2007, 45, 1035-1037.	3.9	109
119	Laboratory Diagnostic Techniques for Entamoeba Species. Clinical Microbiology Reviews, 2007, 20, 511-532.	13.6	382
120	Immunization of Cattle with Live Tachyzoites of Neospora caninum Confers Protection against Fetal Death. Infection and Immunity, 2007, 75, 1343-1348.	2.2	109
121	Amoebiasis: current status in Australia. Medical Journal of Australia, 2007, 186, 412-416.	1.7	66
122	Dientamoeba fragilisas a Cause of Travelersâ€™ Diarrhea: Report of Seven Cases: Table 1. Journal of Travel Medicine, 2007, 14, 72-73.	3.0	12
123	Neosporosis and hammondiosis in dogs. Journal of Small Animal Practice, 2007, 48, 308-312.	1.2	70
124	Irritable bowel syndrome: A review on the role of intestinal protozoa and the importance of their detection and diagnosis. International Journal for Parasitology, 2007, 37, 11-20.	3.1	152
125	PREVALENCE OF ENTERIC PROTOZOA IN HUMAN IMMUNODEFICIENCY VIRUS (HIV)â€“POSITIVE AND HIV-NEGATIVE MEN WHO HAVE SEX WITH MEN FROM SYDNEY, AUSTRALIA. American Journal of Tropical Medicine and Hygiene, 2007, 76, 549-552.	1.4	77
126	Prevalence of enteric protozoa in human immunodeficiency virus (HIV)-positive and HIV-negative men who have sex with men from Sydney, Australia. American Journal of Tropical Medicine and Hygiene, 2007, 76, 549-52.	1.4	33

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127	Locally acquired infection with Entamoeba histolytica in men who have sex with men in Australia. Medical Journal of Australia, 2006, 185, 417-417.	1.7	11
128	Hammondia isolated from dogs and foxes are genetically distinct. Parasitology, 2006, 132, 187.	1.5	19
129	Performance characteristics and optimisation of cut-off values of two enzyme-linked immunosorbent assays for the detection of antibodies to Neospora caninum in the serum of cattle. Veterinary Parasitology, 2006, 140, 61-68.	1.8	5
130	Prevalence of Neospora caninum infection in Australian (NSW) dairy cattle estimated by a newly validated ELISA for milk. Veterinary Parasitology, 2006, 142, 173-178.	1.8	25
131	If control of Neospora caninum infection is technically feasible does it make economic sense?. Veterinary Parasitology, 2006, 142, 23-34.	1.8	70
132	Dientamoebiasis: clinical importance and recent advances. Trends in Parasitology, 2006, 22, 92-96.	3.3	78
133	Evaluation of Three Diagnostic Methods, Including Real-Time PCR, for Detection of Dientamoeba fragilis in Stool Specimens. Journal of Clinical Microbiology, 2006, 44, 232-235.	3.9	56
134	Attachment and invasion ofToxoplasma gondiiandNeospora caninumto epithelial and fibroblast cell linesin vitro. Parasitology, 2005, 131, 583-590.	1.5	18
135	Subcellular fractionation and molecular characterization of the pellicle and plasmalemma of Neospora caninum. Parasitology, 2005, 131, 467.	1.5	6
136	Culture of Neospora caninum in the presence of a Mycoplasma Removal Agent results in the selection of a mutant population of tachyzoites. Parasitology, 2005, 130, 607-610.	1.5	2
137	Neospora abortions in dairy cattle: diagnosis, mode of transmission and control. Veterinary Parasitology, 2005, 128, 231-241.	1.8	113
138	Detection of Dientamoeba fragilis in fresh stool specimens using PCR. International Journal for Parasitology, 2005, 35, 57-62.	3.1	75
139	Application of bioinformatics to parasitology. International Journal for Parasitology, 2005, 35, 463-464.	3.1	1
140	Reduction in transplacental transmission of Neospora caninum in outbred mice by vaccination. International Journal for Parasitology, 2005, 35, 821-828.	3.1	45
141	Prospective Study of the Prevalence, Genotyping, and Clinical Relevance of Dientamoeba fragilis Infections in an Australian Population. Journal of Clinical Microbiology, 2005, 43, 2718-2723.	3.9	84
142	AUTOFLUORESCENCE OF TOXOPLASMA GONDII AND NEOSPORA CANINUM CYSTS IN VITRO. Journal of Parasitology, 2005, 91, 17-23.	0.7	7
143	In Vitro Induction of Neospora caninum Bradyzoites in Vero Cells Reveals Differential Antigen Expression, Localization, and Host-Cell Recognition of Tachyzoites and Bradyzoites. Infection and Immunity, 2004, 72, 576-583.	2.2	70
144	An outbreak of abortion in a dairy herd associated with Neospora caninum and bovine pestivirus infection. Australian Veterinary Journal, 2004, 82, 99-101.	1.1	20

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145	Microarrays and stage conversion in Toxoplasma gondii. Trends in Parasitology, 2004, 20, 288-295.	3.3	9
146	The cell-mediated immune response to Neospora caninum during pregnancy in the mouse is associated with a bias towards production of interleukin-4. International Journal for Parasitology, 2004, 34, 723-732.	3.1	48
147	Genomics and Its Impact on Parasitology and the Potential for Development of New Parasite Control Methods. DNA and Cell Biology, 2003, 22, 395-403.	1.9	18
148	The Design and Analysis of Microarray Experiments: Applications in Parasitology. DNA and Cell Biology, 2003, 22, 357-394.	1.9	16
149	Hammondia heydornioocysts in the faeces of a greyhound in New Zealand. New Zealand Veterinary Journal, 2003, 51, 38-39.	0.9	5
150	HAMMONDIA HEYDORNI FROM THE ARABIAN MOUNTAIN GAZELLE AND RED FOX IN SAUDI ARABIA. Journal of Parasitology, 2003, 89, 535-539.	0.7	26
151	Control options forNeospora caninuminfections in cattle â€" current state of knowledge. New Zealand Veterinary Journal, 2002, 50, 86-92.	0.9	31
152	A previously unclassified trypanosomatid responsible for human cutaneous lesions in Martinique (French West Indies) is the most divergent member of the genus Leishmania ss. Parasitology, 2002, 124, 17-24.	1.5	58
153	CHARACTERIZATION OF AN OUTBRED PREGNANT MOUSE MODEL OF NEOSPORA CANINUM INFECTION. Journal of Parasitology, 2002, 88, 691-696.	0.7	36
154	Redescription of Neospora caninum and its differentiation from related coccidia. International Journal for Parasitology, 2002, 32, 929-946.	3.1	185
155	Neospora caninum: a cause of immune-mediated failure of pregnancy?. Trends in Parasitology, 2002, 18, 391-394.	3.3	95
156	Characterisation of the first Australian isolate of Neospora caninum from cattle. Australian Veterinary Journal, 2002, 80, 620-625.	1.1	66
157	Ribosomal DNA spacer genotypes of the Anopheles bancroftii group (Diptera: Culicidae) from Australia and Papua New Guinea. Insect Molecular Biology, 2001, 10, 407-413.	2.0	38
158	Isolation of Neospora caninum genes detected during a chronic murine infection. International Journal for Parasitology, 2001, 31, 67-71.	3.1	19
159	Isolation, characterization and expression of a GRA2 homologue from Neospora caninum. Parasitology, 2000, 120, 383-390.	1.5	29
160	A Phylogenetic Study of the Anopheles punctulatus Group of Malaria Vectors Comparing rDNA Sequence Alignments Derived from the Mitochondrial and Nuclear Small Ribosomal Subunits. Molecular Phylogenetics and Evolution, 2000, 17, 430-436.	2.7	37
161	Populations of the south-west Pacific malaria vector Anopheles farauti s.s. revealed by ribosomal DNA transcribed spacer polymorphisms. Heredity, 2000, 84, 244-253.	2.6	48
162	Subset partitioning of the ribosomal DNA small subunit and its effects on the phylogeny of the Anopheles punctulatus group. Insect Molecular Biology, 2000, 9, 515-520.	2.0	22

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163	Differential ecology of Anopheles punctulatus and three members of the Anopheles farauti complex of mosquitoes on Guadalcanal, Solomon Islands, identified by PCR-RFLP analysis. Medical and Veterinary Entomology, 2000, 14, 308-312.	1.5	36
164	Seroprevalence of Neospora caninum infection following an abortion outbreak in a dairy cattle herd. Australian Veterinary Journal, 2000, 78, 262-266.	1.1	53
165	The Leishmania major RNA Polymerase II Largest Subunit Lacks a Carboxy-Terminus Heptad Repeat Structure and its Encoding Gene is Linked with the Calreticulin Gene. Protist, 2000, 151, 57-68.	1.5	4
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182	Monophyletic origin of the genus Sauroleishmania. Archiv FÃ¼r Protistenkunde, 1997, 148, 269-275.	0.8	3
183	Evolution of the genus Leishmania revealed by comparison of DNA and RNA polymerase gene sequences1Note: Nucleotide sequence data reported in this paper have been submitted to the GenBankâ„¢ data base with the accession numbers: POLA/RPOIILS (AF009134/AF009153, Leishmania adleri); (AF009135/NS, Leishmania aethiopica); (AF009136/AF009154, Leishmania amazonensis); (AF009137/NS,) Tj ETQq1 1 0.784314 rgBT /O		

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