Stephen L. Cameron

List of Publications by Year
in descending order

126 papers	8,607 citations	53794 45 h-index	48315 88 g-index
127 all docs	127 docs citations	127 times ranked	6059 citing authors

#	Article	IF	Citations
1	Middle Jurassic origin in India: a new look at evolution of Vermileonidae and time-scaled relationships of lower brachyceran flies. Zoological Journal of the Linnean Society, 2022, 194, 938-959.	2.3	0
2	Gene arrangement, phylogeny and divergence time estimation of mitogenomes in Thrips. Molecular Biology Reports, 2022, 49, 6269-6283.	2.3	5
3	Independent evolution of highly variable, fragmented mitogenomes of parasitic lice. Communications Biology, 2022, 5, .	4.4	8
4	Structure, gene order, and nucleotide composition of mitochondrial genomes in parasitic lice from Amblycera. Gene, 2021, 768, 145312.	2.2	13
5	Polyzosteria cockroaches in Tasmania (Blattodea: Blattidae: Polyzosteriinae) represent a new, endemic species, with allopatric alpine and coastal sub-populations . Zootaxa, 2021, 4926, 384-400.	0.5	0
6	How are the mitochondrial genomes reorganized in Hexapoda? Differential evolution and the first report of convergences within Hexapoda. Gene, 2021, 791, 145719.	2.2	17
7	Development of internal COI primers to improve and extend barcoding of fruit flies (Diptera:) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf	3.0	11
8	A newly recorded Rickettsia of the Torix group is a recent intruder and an endosymbiont in the whitefly Bemisia tabaci. Environmental Microbiology, 2020, 22, 1207-1221.	3.8	20
9	Rearrangement and evolution of mitochondrial genomes in Thysanoptera (Insecta). Scientific Reports, 2020, 10, 695.	3.3	49
10	How well do multispecies coalescent methods perform with mitochondrial genomic data? A case study of butterflies and moths (Insecta: Lepidoptera). Systematic Entomology, 2020, 45, 857-873.	3.9	15
11	Mitochondrial genomes of<i>Columbicola</i>feather lice are highly fragmented, indicating repeated evolution of minicircle-type genomes in parasitic lice. PeerJ, 2020, 8, e8759.	2.0	21
12	Closeâ€distance courtship of laboratory reared <scp><i>Bactrocera tryoni</i></scp> (Diptera:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 30	1.4	5
13	<i>Bactrocera dorsalis</i> (Hendel) (Diptera: Tephritidae) is not invasive through Asia: It's been there all along. Journal of Applied Entomology, 2019, 143, 797-801.	1.8	28
14	In the footsteps of Wallace: population structure in the breadfruit fruit fly, <i>Bactrocera umbrosa</i> (F.) (Diptera: Tephritidae), suggests disjunction across the Indoâ€Australian Archipelago. Austral Entomology, 2019, 58, 602-613.	1.4	10
15	Insight into the microbial world of Bemisia tabaci cryptic species complex and its relationships with its host. Scientific Reports, 2019, 9, 6568.	3.3	23
16	The phylogeny and evolutionary timescale of stoneflies (Insecta: Plecoptera) inferred from mitochondrial genomes. Molecular Phylogenetics and Evolution, 2019, 135, 123-135.	2.7	35
17	Extensive host-switching of avian feather lice following the Cretaceous-Paleogene mass extinction event. Communications Biology, 2019, 2, 445.	4.4	20
18	Evidence from Australian mesic zone dung beetles supports their Gondwanan origin and Mesozoic diversification of the Scarabaeinae. Insect Systematics and Evolution, 2019, 50, 162-188.	0.7	12

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19	A transcriptomeâ€based analytical workflow for identifying loci for species diagnosis: a case study with <i>Bactrocera</i> fruit flies (Diptera: Tephritidae). Austral Entomology, 2019, 58, 395-408.	1.4	12
20	Discovery of a new species of Stromatium Audinetâ€Serville, 1834 (Coleoptera: Cerambycidae) native to Australia, based on morphology and DNA barcoding. Austral Entomology, 2019, 58, 137-147.	1.4	1
21	Transoceanic Dispersal and Plate Tectonics Shaped Global Cockroach Distributions: Evidence from Mitochondrial Phylogenomics. Molecular Biology and Evolution, 2018, 35, 970-983.	8.9	73
22	Plant-Mediated Female Transcriptomic Changes Post-Mating in a Tephritid Fruit Fly, Bactrocera tryoni. Genome Biology and Evolution, 2018, 10, 94-107.	2.5	12
23	Mitochondrial phylogenomics and genome rearrangements in the barklice (Insecta: Psocodea). Molecular Phylogenetics and Evolution, 2018, 119, 118-127.	2.7	44
24	Phylogenomics and the evolution of hemipteroid insects. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 12775-12780.	7.1	275
25	Population structure of a global agricultural invasive pest, <i>Bactrocera dorsalis</i> (Diptera:) Tj ETQq1 1 0.784314 rgBT /Overlock 1	3.1	40
26	Chromatin immunoprecipitation (ChIP) method for non-model fruit flies (Diptera: Tephritidae) and evidence of histone modifications. PLoS ONE, 2018, 13, e0194420.	2.5	4
27	The mating system of the true fruit fly <i>Bactrocera tryoni</i> and its sister species, <i>Bactrocera neohumeralis</i>. Insect Science, 2017, 24, 478-490.	3.0	12
28	Parallel evolution of mound-building and grass-feeding in Australian nasute termites. Biology Letters, 2017, 13, 20160665.	2.3	20
29	Signatures of invasion: using an integrative approach to infer the spread of melon fly, Zeugodacus cucurbitae (Diptera: Tephritidae), across Southeast Asia and the West Pacific. Biological Invasions, 2017, 19, 1597-1619.	2.4	13
30	Population structure in Zeugodacus cucurbitae (Diptera: Tephritidae) across Thailand and the Thaiâ€"Malay peninsula: natural barriers to a great disperser. Biological Journal of the Linnean Society, 2017, 121, 540-555.	1.6	10
31	A review of the status of Coptotermes (Isoptera : Rhinotermitidae) species in Australia with the description of two new small termite species from northern and eastern Australia. Invertebrate Systematics, 2017, 31, 180.	1.3	3
32	Ecological diversification of the Australian <i>Coptotermes</i> termites and the evolution of mound building. Journal of Biogeography, 2017, 44, 1405-1417.	3.0	12
33	Integrative taxonomy versus taxonomic authority without peer review: the case of the <scp>O</scp>riental fruit fly, <i>Bactrocera dorsalis</i> (<scp>T</scp>ephritidae). Systematic Entomology, 2017, 42, 609-620.	3.9	24
34	Effect of Body Size, Age, and Premating Experience on Male Mating Success in Bactrocera tryoni (Diptera: Tephritidae). Journal of Economic Entomology, 2017, 110, 2278-2281.	1.8	17
35	Tephritid Integrative Taxonomy: Where We Are Now, with a Focus on the Resolution of Three Tropical Fruit Fly Species Complexes. Annual Review of Entomology, 2017, 62, 147-164.	11.8	38
36	Mitochondrial Genomes Provide Insights into the Phylogeny of Lauxanioidea (Diptera: Cyclorrhapha). International Journal of Molecular Sciences, 2017, 18, 773.	4.1	17

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37	Transcriptomes of three species of Tipuloidea (Diptera, Tipulomorpha) and implications for phylogeny of Tipulomorpha. PLoS ONE, 2017, 12, e0173207.	2.5	14
38	If Dung Beetles (Scarabaeidae: Scarabaeinae) Arose in Association with Dinosaurs, Did They Also Suffer a Mass Co-Extinction at the K-Pg Boundary?. PLoS ONE, 2016, 11, e0153570.	2.5	74
39	Comparative Mt Genomics of the Tipuloidea (Diptera: Nematocera: Tipulomorpha) and Its Implications for the Phylogeny of the Tipulomorpha. PLoS ONE, 2016, 11, e0158167.	2.5	27
40	Molecular phylogenetics of <scp>A</scp>ustralian weevils (<scp>C</scp>oleoptera:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 627 Td (<scp independent analyses. Austral Entomology, 2016, 55, 217-233.	1.4	38
41	â€˜Ladd trapsâ€™ as a visual trap for male and female Queensland fruit fly, <i><scp>Bactrocera tryoni</scp></i> (<scp>Diptera: Tephritidae</scp>). Austral Entomology, 2016, 55, 324-329.	1.4	8
42	The complete mitochondrial genome of the tarnished plant bug,Lygus lineolaris(Heteroptera: Miridae). Mitochondrial DNA, 2016, 27, 48-49.	0.6	5
43	Revisiting <i>Coptotermes</i> (Isoptera: Rhinotermitidae): a global taxonomic road map for species validity and distribution of an economically important subterranean termite genus. Systematic Entomology, 2016, 41, 299-306.	3.9	65
44	The First Mitochondrial Genome of the Sepsid Fly Nemopoda mamaevi Ozerov, 1997 (Diptera:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 46 e0123594.	2.5	17
45	The Phylogeny and Evolutionary Timescale of Muscoidea (Diptera: Brachycera: Calyptratae) Inferred from Mitochondrial Genomes. PLoS ONE, 2015, 10, e0134170.	2.5	31
46	The complete mitochondrial genome of a flea,<i>Jellisonia amadoi</i>(Siphonaptera: Ceratophyllidae). Mitochondrial DNA, 2015, 26, 289-290.	0.6	8
47	A mitochondrial genome phylogeny of owlet moths (Lepidoptera: Noctuoidea), and examination of the utility of mitochondrial genomes for lepidopteran phylogenetics. Molecular Phylogenetics and Evolution, 2015, 85, 230-237.	2.7	76
48	Synonymization of key pest species within the <i><scp>B</scp>actrocera dorsalis</i> species complex (<scp>D</scp>iptera: <scp>T</scp>ephritidae): taxonomic changes based on a review of 20â€‰years of integrative morphological, molecular, cytogenetic, behavioural and chemoecological data. Systematic Entomology, 2015, 40, 456-471.	3.9	175
49	One and the same: integrative taxonomic evidence that <i><scp>B</scp>actrocera invadens</i> (<scp>D</scp>iptera: <scp>T</scp>ephritidae) is the same species as the <scp>O</scp>riental fruit fly <i><scp>B</scp>actrocera dorsalis</i>. Systematic Entomology, 2015, 40, 472-486.	3.9	88
50	The Evolutionary History of Termites as Inferred from 66 Mitochondrial Genomes. Molecular Biology and Evolution, 2015, 32, 406-421.	8.9	268
51	The origins and radiation of Australian Coptotermes termites: From rainforest to desert dwellers. Molecular Phylogenetics and Evolution, 2015, 82, 234-244.	2.7	25
52	Effects of laboratory colonization on Bactrocera dorsalis (Diptera, Tephritidae) mating behaviour: â€˜what a difference a year makesâ€™. ZooKeys, 2015, 540, 369-383.	1.1	34
53	Beyond Moaâ€™s Ark and Wallaceâ€™s Line: extralimital distribution of new species of Austronothrus (Acari, Oribatida, Crotoniidae) and the endemicity of the New Zealand oribatid mite fauna. Zootaxa, 2014, 3780, 263-81.	0.5	14
54	Australian species of spore-feeding Thysanoptera in the genera Carientothrips and Nesothrips (Thysanoptera: Idolothripinae). Zootaxa, 2014, 3821, 193.	0.5	6

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55	Diverse urban plantings managed with sufficient resource availability can increase plant productivity and arthropod diversity. Frontiers in Plant Science, 2014, 5, 517.	3.6	5
56	Trans-Bass Strait speciation and trans-Pacific dispersal in theMyoporumthrips (Thysanoptera,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 702	1.4	4
57	How to sequence and annotate insect mitochondrial genomes for systematic and comparative genomics research. Systematic Entomology, 2014, 39, 400-411.	3.9	206
58	Insect Mitochondrial Genomics: Implications for Evolution and Phylogeny. Annual Review of Entomology, 2014, 59, 95-117.	11.8	1,012
59	Multiâ€gene phylogenetic analysis of southâ€east Asian pest members of the <i>Bactrocera dorsalis</i> species complex (Diptera: Tephritidae) does not support current taxonomy. Journal of Applied Entomology, 2014, 138, 235-253.	1.8	67
60	Sexual selection in true fruit flies (<scp>D</scp>iptera: <scp>T</scp>ephritidae): transcriptome and experimental evidences for phytochemicals increasing male competitive ability. Molecular Ecology, 2014, 23, 4645-4657.	3.9	35
61	A Preliminary Framework for DNA Barcoding, Incorporating the Multispecies Coalescent. Systematic Biology, 2014, 63, 639-644.	5.6	53
62	Towards a phylogeny of the Tenebrionoidea (Coleoptera). Molecular Phylogenetics and Evolution, 2014, 79, 305-312.	2.7	49
63	The Complexities of Knowing What It Is You Are Trapping. , 2014, , 611-632.		9
64	Trapping to Monitor Tephritid Movement: Results, Best Practice, and Assessment of Alternatives. , 2014, , 175-217.		27
65	Utility of COI, CAD and morphological data for resolving relationships within the genus Sarcophaga (sensu lato) (Diptera: Sarcophagidae): A preliminary study. Molecular Phylogenetics and Evolution, 2013, 69, 133-141.	2.7	28
66	The evolution and biogeography of the austral horse fly tribe Scionini (Diptera: Tabanidae:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 307 Td Evolution, 2013, 68, 516-540.	2.7	31
67	Piecing together an integrative taxonomic puzzle: microsatellite, wing shape and aedeagus length analyses of <i>Bactrocera dorsalis s.l.</i> (Diptera: Tephritidae) find no evidence of multiple lineages in a proposed contact zone along the Thai/Malay Peninsula. Systematic Entomology, 2013, 38, 2-13.	3.9	70
68	Evolution of lure response in tephritid fruit flies: phytochemicals as drivers of sexual selection. Animal Behaviour, 2013, 85, 781-789.	1.9	58
69	A molecular phylogeny of the checkered beetles and a description of <scp>E</scp>piclininae a new subfamily (<scp>C</scp>oleoptera: <scp>C</scp>leroidea: <scp>C</scp>leridae). Systematic Entomology, 2013, 38, 626-636.	3.9	40
70	A phylogenetic analysis and taxonomic revision of the oribatid mite family Malaconothridae (Acari:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 from Australia. Zootaxa, 2013, 3681, 301-46.	0.5	14
71	Comprehensive evaluation of DNA barcoding for the molecular species identification of forensically important Australian Sarcophagidae (Diptera). Invertebrate Systematics, 2012, 26, 515.	1.3	28
72	Systematics and biology of the iconic Australian scribbly gum moths Ogmograptis Meyrick (Lepidoptera : Bucculatricidae) and their unique insectâ€“plant interaction. Invertebrate Systematics, 2012, 26, 357.	1.3	13

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73	A molecular phylogeny for the Tribe Dacini (Diptera: Tephritidae): Systematic and biogeographic implications. Molecular Phylogenetics and Evolution, 2012, 64, 513-523.	2.7	76
74	Population structure of Bactrocera dorsalis s.s., B. papayae and B. philippinensis (Diptera: Tephritidae) in southeast Asia: evidence for a single species hypothesis using mitochondrial DNA and wing-shape data. BMC Evolutionary Biology, 2012, 12, 130.	3.2	75
75	A mitochondrial genome phylogeny of termites (Blattodea: Termitoidae): Robust support for interfamilial relationships and molecular synapomorphies define major clades. Molecular Phylogenetics and Evolution, 2012, 65, 163-173.	2.7	127
76	Beyond barcoding: A mitochondrial genomics approach to molecular phylogenetics and diagnostics of blowflies (Diptera: Calliphoridae). Gene, 2012, 511, 131-142.	2.2	142
77	The complete mitochondrial genome of the flesh fly,<i>Sarcophaga impatiens</i>Walker (Diptera:) Tj ETQq1 1 0.784314 rgBT /Overlock	0.6	29
78	A view from the edge of the forest: recent progress in understanding the relationships of the insect orders. Australian Journal of Entomology, 2012, 51, 79-87.	1.1	19
79	Integrative taxonomy, or iterative taxonomy?. Systematic Entomology, 2011, 36, 209-217.	3.9	254
80	The complete mitochondrial genome of Spilonota lechriaspis Meyrick (Lepidoptera: Tortricidae). Molecular Biology Reports, 2011, 38, 3757-3764.	2.3	46
81	Mitochondrial genome deletions and minicircles are common in lice (Insecta: Phthiraptera). BMC Genomics, 2011, 12, 394.	2.8	90
82	Potential efficacy of mitochondrial genes for animal DNA barcoding: a case study using eutherian mammals. BMC Genomics, 2011, 12, 84.	2.8	83
83	The complete mitochondrial genome of the gall-forming fly,Fergusonina tayloriNelson and Yeates (Diptera: Fergusoninidae). Mitochondrial DNA, 2011, 22, 197-199.	0.6	10
84	The complete mitochondrial genome of the yellow coaster, AcraeaÂissoria (Lepidoptera: Nymphalidae:) Tj ETQq0 0 0 rgBT /Overlock 10 T Molecular Biology Reports, 2010, 37, 3431-3438.	2.3	79
85	When phylogenetic assumptions are violated: base compositional heterogeneity and among-site rate variation in beetle mitochondrial phylogenomics. Systematic Entomology, 2010, 35, 429-448.	3.9	121
86	The mitochondrial genome of the onychophoran Opisthopatus cinctipes (Peripatopsidae) reflects the ancestral mitochondrial gene arrangement of Panarthropoda and Ecdysozoa. Molecular Phylogenetics and Evolution, 2010, 57, 285-292.	2.7	38
87	Extensive duplication events account for multiple control regions and pseudo-genes in the mitochondrial genome of the velvet worm Metaperipatus inae (Onychophora, Peripatopsidae). Molecular Phylogenetics and Evolution, 2010, 57, 293-300.	2.7	24
88	Genome sequences of the human body louse and its primary endosymbiont provide insights into the permanent parasitic lifestyle. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 12168-12173.	7.1	482
89	Revision of the oribatid mite genus Austronothrus Hammer (Acari:Oribatida):sexual dimorphism and a re-evaluation of the phylogenetic relationships of the family Crotoniidae. Invertebrate Systematics, 2009, 23, 87.	1.3	10
90	Nonstationary Evolution and Compositional Heterogeneity in Beetle Mitochondrial Phylogenomics. Systematic Biology, 2009, 58, 381-394.	5.6	162

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91	Characterization of 67 Mitochondrial tRNA Gene Rearrangements in the Hymenoptera Suggests That Mitochondrial tRNA Gene Position Is Selectively Neutral. Molecular Biology and Evolution, 2009, 26, 1607-1617.	8.9	176
92	A mitochondrial genome phylogeny of the Neuropterida (laceâ€wings, alderflies and snakeflies) and their relationship to the other holometabolous insect orders. Zoologica Scripta, 2009, 38, 575-590.	1.7	122
93	Phylogenetic approaches for the analysis of mitochondrial genome sequence data in the Hymenoptera â€“ A lineage with both rapidly and slowly evolving mitochondrial genomes. Molecular Phylogenetics and Evolution, 2009, 52, 512-519.	2.7	90
94	Distribution and phylogenetic relationships of Australian glow-worms Arachnocampa (Diptera,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 62	2.7	44
95	A preliminary mitochondrial genome phylogeny of Orthoptera (Insecta) and approaches to maximizing phylogenetic signal found within mitochondrial genome data. Molecular Phylogenetics and Evolution, 2008, 49, 59-68.	2.7	174
96	Converse Bergmann cline in a <i>Eucalyptus</i> herbivore, <i>Paropsis atomaria</i> Olivier (Coleoptera: Chrysomelidae): phenotypic plasticity or local adaptation?. Global Ecology and Biogeography, 2008, 17, 424-431.	5.8	27
97	The complete mitochondrial genome of the sexual oribatid mite Steganacarus magnus: genome rearrangements and loss of tRNAs. BMC Genomics, 2008, 9, 532.	2.8	67
98	The complete mitochondrial genome of the tobacco hornworm, Manduca sexta, (Insecta: Lepidoptera:) Tj ETQq0 0 0 rgBT /Overlock 10 2008, 408, 112-123.	2.2	335
99	A Comparative Analysis of Mitochondrial Genomes in Coleoptera (Arthropoda: Insecta) and Genome Descriptions of Six New Beetles. Molecular Biology and Evolution, 2008, 25, 2499-2509.	8.9	211
100	Mitochondrial genome organization and phylogeny of two vespid wasps. Genome, 2008, 51, 800-808.	2.0	93
101	Mitochondrial genomic comparisons of the subterranean termites from the Genus Reticulitermes (Insecta: Isoptera: Rhinotermitidae). Genome, 2007, 50, 188-202.	2.0	93
102	Save Isoptera: A comment on Inward <i>et al</i> .. Biology Letters, 2007, 3, 562-563.	2.3	65
103	The complete mitochondrial genome sequence of the Mormon cricket (Anabrus simplex: Tettigoniidae:) Tj ETQq1 1 0.784314 rgBT /Ov	2.0	107
104	A mitochondrial genome phylogeny of Diptera: whole genome sequence data accurately resolve relationships over broad timescales with high precision. Systematic Entomology, 2007, 32, 40-59.	3.9	231
105	The Mitochondrial Genome of the Screamer Louse Bothriometopus (Phthiraptera: Ischnocera): Effects of Extensive Gene Rearrangements on the Evolution of the Genome. Journal of Molecular Evolution, 2007, 65, 589-604.	1.8	122
106	A Genomic Perspective on the Shortcomings of Mitochondrial DNA for â€œBarcodingâ€• Identification. Journal of Heredity, 2006, 97, 581-594.	2.4	401
107	Who Will Actually Use DNA Barcoding and What Will It Cost?. Systematic Biology, 2006, 55, 844-847.	5.6	67
108	Are plant DNA barcodes a search for the Holy Grail?. Trends in Ecology and Evolution, 2006, 21, 1-2.	8.7	103

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109	Species status and population structure of the Australian Eucalyptus pest Paropsis atomaria Olivier (Coleoptera: Chrysomelidae). Agricultural and Forest Entomology, 2006, 8, 323-332.	1.3	10
110	Extraordinary number of gene rearrangements in the mitochondrial genomes of lice (Phthiraptera:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	2.0	64
111	Mitochondrial genomics and the new insect order Mantophasmatodea. Molecular Phylogenetics and Evolution, 2006, 38, 274-279.	2.7	138
112	On Micromastigotes and the evolution of the hypermastigont condition in the Parabasalida. Journal of Eukaryotic Microbiology, 2005, 52, 7S-27S.	1.7	0
113	Mitochondrial genome data alone are not enough to unambiguously resolve the relationships of Entognatha, Insecta and Crustacea sensu lato (Arthropoda). Cladistics, 2004, 20, 534-557.	3.3	122
114	Phylogeny and Biogeography of the â€œAustralianâ€•Trichostomes (Ciliophora: Litostomata). Protist, 2004, 155, 215-235.	1.5	23
115	Trichostome ciliates from Australian marsupials. II. Polycosta gen. nov. (Litostomatea: Polycostidae) Tj ETQq1 1 0.784314 rgBT /Overl	1.5	17
116	Trichostome ciliates from Australian marsupials. III. Megavestibulum gen. nov. (Litostomatea:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 462	1.5	13
117	Trichostome ciliates from Australian marsupials. IV. Distribution of the ciliate fauna. European Journal of Protistology, 2003, 39, 139-147.	1.5	11
118	Taxonomy and phylogeny of endosymbiotic ciliates (Ciliophora: Litostomatea) associated with Australian herbivorous marsupials. International Journal for Parasitology, 2003, 33, 347-355.	3.1	10
119	The ultrastructure of Amylovorax dehorityi comb. nov. and erection of the Amylovoracidae fam. nov. (Ciliophora: Trichostomatia). European Journal of Protistology, 2002, 38, 29-44.	1.5	27
120	The ultrastructure of Macropodinium moiri and revised diagnosis of the Macropodiniidae (Litostomatea: Trichostomatia). European Journal of Protistology, 2002, 38, 179-194.	1.5	22
121	Trichostome ciliates from Australian marsupials. I. Bandia gen. nov. (Litostomatea: Amylovoracidae). European Journal of Protistology, 2002, 38, 405-429.	1.5	20
122	Evidence for an Independent Radiation of Endosymbiotic Litostome Ciliates within Australian Marsupial Herbivores. Molecular Phylogenetics and Evolution, 2001, 20, 302-310.	2.7	27
123	Four New Species of Macropodinium (Ciliophora: Litostomatea) from Australian Wallabies and Pademelons. Journal of Eukaryotic Microbiology, 2001, 48, 542-555.	1.7	19
124	Stomatogenesis in the ciliate genus MacropodiniumDehority, 1996 (Litostomatea: Macropodiniidae). European Journal of Protistology, 2001, 37, 199-206.	1.5	13
125	Novel isotrichid ciliates endosymbiotic in Australian macropodid marsupials. Systematic Parasitology, 2000, 46, 45-57.	1.1	16
126	First record of Cycloposthium edentatum Strelkow, 1928 from the black-striped wallaby, Macropus dorsalis. Parasitology Research, 2000, 86, 158-162.	1.6	18