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	Insect Mitochondrial Genomics: Implications for Evolution and Phylogeny. Annual Review of Entomology, 2014, 59, 95-117.	11.8	1,012
2	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
3	A Genomic Perspective on the Shortcomings of Mitochondrial DNA for â€œBarcodingâ€•Identification. Journal of Heredity, 2006, 97, 581-594.	2.4	401
4	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
5	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
6	The Evolutionary History of Termites as Inferred from 66 Mitochondrial Genomes. Molecular Biology and Evolution, 2015, 32, 406-421.	8.9	268
7	Integrative taxonomy, or iterative taxonomy?. Systematic Entomology, 2011, 36, 209-217.	3.9	254
8	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
9	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
10	How to sequence and annotate insect mitochondrial genomes for systematic and comparative genomics research. Systematic Entomology, 2014, 39, 400-411.	3.9	206
11	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
12	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
13	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
14	Nonstationary Evolution and Compositional Heterogeneity in Beetle Mitochondrial Phylogenomics. Systematic Biology, 2009, 58, 381-394.	5.6	162
15	Beyond barcoding: A mitochondrial genomics approach to molecular phylogenetics and diagnostics of blowflies (Diptera: Calliphoridae). Gene, 2012, 511, 131-142.	2.2	142
16	Mitochondrial genomics and the new insect order Mantophasmatodea. Molecular Phylogenetics and Evolution, 2006, 38, 274-279.	2.7	138
17	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
18	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

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19	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
20	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
21	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
22	The complete mitochondrial genome sequence of the Mormon cricket (Anabrus simplex: Tettigoniidae:) Tj ETQq0 0 0 rgBT /Overlock 10	2.0	107
23	Are plant DNA barcodes a search for the Holy Grail?. Trends in Ecology and Evolution, 2006, 21, 1-2.	8.7	103
24	Mitochondrial genomic comparisons of the subterranean termites from the Genus Reticulitermes (Insecta: Isoptera: Rhinotermitidae). Genome, 2007, 50, 188-202.	2.0	93
25	Mitochondrial genome organization and phylogeny of two vespid wasps. Genome, 2008, 51, 800-808.	2.0	93
26	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
27	Mitochondrial genome deletions and minicircles are common in lice (Insecta: Phthiraptera). BMC Genomics, 2011, 12, 394.	2.8	90
28	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
29	Potential efficacy of mitochondrial genes for animal DNA barcoding: a case study using eutherian mammals. BMC Genomics, 2011, 12, 84.	2.8	83
30	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
31	A molecular phylogeny for the Tribe Dacini (Diptera: Tephritidae): Systematic and biogeographic implications. Molecular Phylogenetics and Evolution, 2012, 64, 513-523.	2.7	76
32	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
33	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
34	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
35	Transoceanic Dispersal and Plate Tectonics Shaped Global Cockroach Distributions: Evidence from Mitochondrial Phylogenomics. Molecular Biology and Evolution, 2018, 35, 970-983.	8.9	73
36	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

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37	Who Will Actually Use DNA Barcoding and What Will It Cost?. Systematic Biology, 2006, 55, 844-847.	5.6	67
38	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
39	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
40	Save Isoptera: A comment on Inward <i>et al</i> .. Biology Letters, 2007, 3, 562-563.	2.3	65
41	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
42	Extraordinary number of gene rearrangements in the mitochondrial genomes of lice (Phthiraptera:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	2.0	64
43	Evolution of lure response in tephritid fruit flies: phytochemicals as drivers of sexual selection. Animal Behaviour, 2013, 85, 781-789.	1.9	58
44	A Preliminary Framework for DNA Barcoding, Incorporating the Multispecies Coalescent. Systematic Biology, 2014, 63, 639-644.	5.6	53
45	Towards a phylogeny of the Tenebrionoidea (Coleoptera). Molecular Phylogenetics and Evolution, 2014, 79, 305-312.	2.7	49
46	Rearrangement and evolution of mitochondrial genomes in Thysanoptera (Insecta). Scientific Reports, 2020, 10, 695.	3.3	49
47	The complete mitochondrial genome of Spilonota lechriaspis Meyrick (Lepidoptera: Tortricidae). Molecular Biology Reports, 2011, 38, 3757-3764.	2.3	46
48	Distribution and phylogenetic relationships of Australian glow-worms Arachnocampa (Diptera,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 30	2.7	44
49	Mitochondrial phylogenomics and genome rearrangements in the barklice (Insecta: Psocodea). Molecular Phylogenetics and Evolution, 2018, 119, 118-127.	2.7	44
50	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
51	Population structure of a global agricultural invasive pest, <i>Bactrocera dorsalis</i> (Diptera:) Tj ETQq1 1 0.784314 rgBT /Overlock 10	3.1	40
52	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
53	Molecular phylogenetics of <scp>A</scp>ustralian weevils (<scp>C</scp>oleoptera:) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 1 independent analyses. Austral Entomology, 2016, 55, 217-233.	1.4	38
54	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

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55	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
56	The phylogeny and evolutionary timescale of stoneflies (Insecta: Plecoptera) inferred from mitochondrial genomes. Molecular Phylogenetics and Evolution, 2019, 135, 123-135.	2.7	35
57	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
58	The evolution and biogeography of the austral horse fly tribe Scionini (Diptera: Tabanidae:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 627 Td Evolution, 2013, 68, 516-540.	2.7	31
59	The Phylogeny and Evolutionary Timescale of Muscoidea (Diptera: Brachycera: Calyptratae) Inferred from Mitochondrial Genomes. PLoS ONE, 2015, 10, e0134170.	2.5	31
60	The complete mitochondrial genome of the flesh fly,<i>Sarcophaga impatiens</i>Walker (Diptera:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	0.6	29
61	Comprehensive evaluation of DNA barcoding for the molecular species identification of forensically important Australian Sarcophagidae (Diptera). Invertebrate Systematics, 2012, 26, 515.	1.3	28
62	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
63	<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
64	Evidence for an Independent Radiation of Endosymbiotic Litostome Ciliates within Australian Marsupial Herbivores. Molecular Phylogenetics and Evolution, 2001, 20, 302-310.	2.7	27
65	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
66	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
67	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
68	Trapping to Monitor Tephritid Movement: Results, Best Practice, and Assessment of Alternatives. , 2014, , 175-217.		27
69	The origins and radiation of Australian Coptotermes termites: From rainforest to desert dwellers. Molecular Phylogenetics and Evolution, 2015, 82, 234-244.	2.7	25
70	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
71	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
72	Phylogeny and Biogeography of the â€œAustralianâ€•Trichostomes (Ciliophora: Litostomata). Protist, 2004, 155, 215-235.	1.5	23

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73	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
74	The ultrastructure of Macropodinium moiri and revised diagnosis of the Macropodiniidae (Litostomatea: Trichostomatia). European Journal of Protistology, 2002, 38, 179-194.	1.5	22
75	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
76	Trichostome ciliates from Australian marsupials. I. Bandia gen. nov. (Litostomatea: Amylovoracidae). European Journal of Protistology, 2002, 38, 405-429.	1.5	20
77	Parallel evolution of mound-building and grass-feeding in Australian nasute termites. Biology Letters, 2017, 13, 20160665.	2.3	20
78	Extensive host-switching of avian feather lice following the Cretaceous-Paleogene mass extinction event. Communications Biology, 2019, 2, 445.	4.4	20
79	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
80	Four New Species of Macropodinium (Ciliophora: Litostomatea) from Australian Wallabies and Pademelons. Journal of Eukaryotic Microbiology, 2001, 48, 542-555.	1.7	19
81	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
82	First record of Cycloposthium edentatum Strelkow, 1928 from the black-striped wallaby, Macropus dorsalis. Parasitology Research, 2000, 86, 158-162.	1.6	18
83	Trichostome ciliates from Australian marsupials. II. Polycosta gen. nov. (Litostomatea: Polycostidae) Tj ETQq1 1 0.784314 rgBT /Overl	1.5	17
84	The First Mitochondrial Genome of the Sepsid Fly Nemopoda mamaevi Ozerov, 1997 (Diptera:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 30 e0123594.	2.5	17
85	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
86	Mitochondrial Genomes Provide Insights into the Phylogeny of Lauxanioidea (Diptera: Cyclorrhapha). International Journal of Molecular Sciences, 2017, 18, 773.	4.1	17
87	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
88	Novel isotrichid ciliates endosymbiotic in Australian macropodid marsupials. Systematic Parasitology, 2000, 46, 45-57.	1.1	16
89	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
90	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

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91	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
92	Transcriptomes of three species of Tipuloidea (Diptera, Tipulomorpha) and implications for phylogeny of Tipulomorpha. PLoS ONE, 2017, 12, e0173207.	2.5	14
93	Stomatogenesis in the ciliate genus MacropodiniumDehority, 1996 (Litostomatea: Macropodiniidae). European Journal of Protistology, 2001, 37, 199-206.	1.5	13
94	Trichostome ciliates from Australian marsupials. III. Megavestibulum gen. nov. (Litostomatea:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 622	1.5	13
95	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
96	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
97	Structure, gene order, and nucleotide composition of mitochondrial genomes in parasitic lice from Amblycera. Gene, 2021, 768, 145312.	2.2	13
98	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
99	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
100	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
101	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
102	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
103	Trichostome ciliates from Australian marsupials. IV. Distribution of the ciliate fauna. European Journal of Protistology, 2003, 39, 139-147.	1.5	11
104	Development of internal COI primers to improve and extend barcoding of fruit flies (Diptera:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 222	3.0	11
105	Taxonomy and phylogeny of endosymbiotic ciliates (Ciliophora: Litostomatea) associated with Australian herbivorous marsupials. International Journal for Parasitology, 2003, 33, 347-355.	3.1	10
106	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
107	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
108	The complete mitochondrial genome of the gall-forming fly,Fergusonina tayloriNelson and Yeates (Diptera: Fergusoninidae). Mitochondrial DNA, 2011, 22, 197-199.	0.6	10

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109	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
110	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
111	The Complexities of Knowing What It Is You Are Trapping. , 2014, , 611-632.		9
112	The complete mitochondrial genome of a flea,<i>Jellisonia amadoi</i>(Siphonaptera: Ceratophyllidae). Mitochondrial DNA, 2015, 26, 289-290.	0.6	8
113	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
114	Independent evolution of highly variable, fragmented mitogenomes of parasitic lice. Communications Biology, 2022, 5, .	4.4	8
115	Australian species of spore-feeding Thysanoptera in the genera Carientothrips and Nesothrips (Thysanoptera: Idolothripinae). Zootaxa, 2014, 3821, 193.	0.5	6
116	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
117	The complete mitochondrial genome of the tarnished plant bug,Lygus lineolaris(Heteroptera: Miridae). Mitochondrial DNA, 2016, 27, 48-49.	0.6	5
118	Closeâ€distance courtship of laboratory reared <scp><i>Bactrocera tryoni</i></scp> (Diptera:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 38	1.4	5
119	Gene arrangement, phylogeny and divergence time estimation of mitogenomes in Thrips. Molecular Biology Reports, 2022, 49, 6269-6283.	2.3	5
120	Trans-Bass Strait speciation and trans-Pacific dispersal in theMyoporumthrips (Thysanoptera,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 302	1.4	4
121	Chromatin immunoprecipitation (ChIP) method for non-model fruit flies (Diptera: Tephritidae) and evidence of histone modifications. PLoS ONE, 2018, 13, e0194420.	2.5	4
122	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
123	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
124	On Micromastigotes and the evolution of the hypermastigont condition in the Parabasalida. Journal of Eukaryotic Microbiology, 2005, 52, 7S-27S.	1.7	0
125	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
126	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