

Daniel F.R. Cleary

List of Publications by Year in descending order

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139
papers

4,485
citations

94433

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138484

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docs citations

142
times ranked

5873
citing authors

#	ARTICLE	IF	CITATIONS
1	Microeukaryotic communities of golf-ball sponges inside and outside of marine lakes. <i>Journal of Sea Research</i> , 2022, 180, 102151.	1.6	1
2	Draft Genome Sequence of <i>Vibrio mediterranei</i> Strain CyArs1. <i>Microbiology Resource Announcements</i> , 2022, , e0015522.	0.6	0
3	The effect of natural disturbances on forest biodiversity: an ecological synthesis. <i>Biological Reviews</i> , 2022, 97, 1930-1947.	10.4	40
4	Characterization of putative circular plasmids in sponge-associated bacterial communities using a selective multiply-primed rolling circle amplification. <i>Molecular Ecology Resources</i> , 2021, 21, 110-121.	4.8	6
5	Composition and diversity of prokaryotic communities sampled from sponges and soft corals in Maldivian waters. <i>Marine Ecology</i> , 2021, 42, e12638.	1.1	5
6	Humic substances modulate fish bacterial communities in a marine recirculating aquaculture system. <i>Aquaculture</i> , 2021, 544, 737121.	3.5	11
7	Archaeal communities of low and high microbial abundance sponges inhabiting the remote western Indian Ocean island of Mayotte. <i>Antonie Van Leeuwenhoek</i> , 2021, 114, 95-112.	1.7	11
8	A global database for metacommunity ecology, integrating species, traits, environment and space. <i>Scientific Data</i> , 2020, 7, 6.	5.3	28
9	Bacterial composition of sponges, sediment and seawater in enclosed and open marine lakes in Ha Long Bay Vietnam. <i>Marine Biology Research</i> , 2020, 16, 18-31.	0.7	6
10	Compositional variation between high and low prokaryotic diversity coral reef biotopes translates to different predicted metagenomic gene content. <i>Antonie Van Leeuwenhoek</i> , 2020, 113, 563-587.	1.7	1
11	Marine lake populations of jellyfish, mussels and sponges host compositionally distinct prokaryotic communities. <i>Hydrobiologia</i> , 2020, 847, 3409-3425.	2.0	3
12	Geographical location and habitat predict variation in prokaryotic community composition of <i>Suberites diversicolor</i> . <i>Annals of Microbiology</i> , 2020, 70, .	2.6	1
13	Aquaponics using a fish farm effluent shifts bacterial communities profile in halophytes rhizosphere and endosphere. <i>Scientific Reports</i> , 2020, 10, 10023.	3.3	9
14	Prokaryote Communities Inhabiting Endemic and Newly Discovered Sponges and Octocorals from the Red Sea. <i>Microbial Ecology</i> , 2020, 80, 103-119.	2.8	14
15	A comparison of the prokaryotic communities associated with seven seaweed species, sediment, and seawater from the Penghu archipelago, Taiwan. <i>Marine Biology Research</i> , 2020, 16, 744-761.	0.7	3
16	Bacterial composition and putative functions associated with sponges, sediment and seawater from the Tioman coral reef system, Peninsular Malaysia. <i>Marine Biology Research</i> , 2020, 16, 729-743.	0.7	1
17	Archaeal communities in sponge, sediment and water from marine lakes and open water habitats. <i>Marine Biology Research</i> , 2019, 15, 259-274.	0.7	7
18	A comparison of prokaryote communities inhabiting sponges, bacterial mats, sediment and seawater in Southeast Asian coral reefs. <i>FEMS Microbiology Ecology</i> , 2019, 95, .	2.7	11

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19	Characterization of bacterioplankton communities from a hatchery recirculating aquaculture system (RAS) for juvenile sole (<i>Solea senegalensis</i>) production. PLoS ONE, 2019, 14, e0211209.	2.5	15
20	Indonesia: Threats to the Country's Biodiversity. , 2019, , 622-632.		0
21	Baseline information on prokaryotic and microeukaryotic plankton communities inside and outside of Indonesian marine lakes. Journal of Sea Research, 2019, 148-149, 23-32.	1.6	3
22	A comparison of microeukaryote communities inhabiting sponges and seawater in a Taiwanese coral reef system. Annals of Microbiology, 2019, 69, 861-866.	2.6	7
23	Microcosm evaluation of the impact of oil contamination and chemical dispersant addition on bacterial communities and sediment remediation of an estuarine port environment. Journal of Applied Microbiology, 2019, 127, 134-149.	3.1	9
24	The sponge microbiome within the greater coral reef microbial metacommunity. Nature Communications, 2019, 10, 1644.	12.8	86
25	Bacterial and microeukaryotic plankton communities in a semi-intensive aquaculture system of sea bass (<i>Dicentrarchus labrax</i>): A seasonal survey. Aquaculture, 2019, 503, 59-69.	3.5	29
26	Assessment of fish community structure along the Jakarta Bay-Pulau Seribu reef complex. Journal of the Marine Biological Association of the United Kingdom, 2019, 99, 503-516.	0.8	2
27	Comparison of bacterial communities associated with <i>Xestospongia testudinaria</i> , sediment and seawater in a Singaporean coral reef ecosystem. Journal of the Marine Biological Association of the United Kingdom, 2019, 99, 331-342.	0.8	3
28	Bacterial Communities Inhabiting the Sponge <i>Biemna fortis</i> , Sediment and Water in Marine Lakes and the Open Sea. Microbial Ecology, 2018, 76, 610-624.	2.8	23
29	Micro-eukaryotic plankton diversity in an intensive aquaculture system for production of <i>Scophthalmus maximus</i> and <i>Solea senegalensis</i> . Aquaculture, 2018, 490, 321-328.	3.5	10
30	Seasonal patterns of bacterioplankton composition in a semi-intensive European seabass (<i>Dicentrarchus labrax</i>) aquaculture system. Aquaculture, 2018, 490, 240-250.	3.5	17
31	Prokaryote composition and predicted metagenomic content of two <i>Cinachyrella</i> Morphospecies and water from West Papuan Marine Lakes. FEMS Microbiology Ecology, 2018, 94, .	2.7	32
32	Compositional analysis of bacterial communities in seawater, sediment, and sponges in the Misool coral reef system, Indonesia. Marine Biodiversity, 2018, 48, 1889-1901.	1.0	32
33	Sponge Prokaryote Communities in Taiwanese Coral Reef and Shallow Hydrothermal Vent Ecosystems. Microbial Ecology, 2018, 75, 239-254.	2.8	17
34	Assessing the bacterial communities of sponges inhabiting the remote western Indian Ocean island of Mayotte. Marine Ecology, 2018, 39, e12517.	1.1	18
35	Prokaryotic communities of Indo-Pacific giant barrel sponges are more strongly influenced by geography than host phylogeny. FEMS Microbiology Ecology, 2018, 94, .	2.7	26
36	Independent and interactive effects of reduced seawater pH and oil contamination on subsurface sediment bacterial communities. Environmental Science and Pollution Research, 2018, 25, 32756-32766.	5.3	6

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37	Variation in the composition and diversity of ground-layer herbs and shrubs in unburnt and burnt landscapes. <i>Journal of Tropical Ecology</i> , 2018, 34, 243-256.	1.1	0
38	Compositional analysis of archaeal communities in high and low microbial abundance sponges in the Misool coral reef system, Indonesia. <i>Marine Biology Research</i> , 2018, 14, 537-550.	0.7	10
39	Bacterial and archaeal communities inhabiting mussels, sediment and water in Indonesian anchialine lakes. <i>Antonie Van Leeuwenhoek</i> , 2018, 111, 237-257.	1.7	14
40	“Blue Carbon” and Nutrient Stocks of Salt Marshes at a Temperate Coastal Lagoon (Ria de Aveiro,) <i>Tj ETQq0 0 0 rgBT /Overlock 10 T</i>	3.3	45
41	Sediment depth and habitat as predictors of the diversity and composition of sediment bacterial communities in an intertidal estuarine environment. <i>Marine Ecology</i> , 2017, 38, e12411.	1.1	25
42	The database of the <sc>PREDICTS</sc> (Projecting Responses of Ecological Diversity In Changing) <i>Tj ETQq0 0 0 rgBT /Overlock 10 T</i>	1.9	186
43	Impact of logging on tree, liana and herb assemblages in a Bornean forest. <i>Journal of Sustainable Forestry</i> , 2017, 36, 806-817.	1.4	7
44	Effect of spatio-temporal shifts in salinity combined with other environmental variables on the ecological processes provided by <i>Zostera noltei</i> meadows. <i>Scientific Reports</i> , 2017, 7, 1336.	3.3	15
45	Globally intertwined evolutionary history of giant barrel sponges. <i>Coral Reefs</i> , 2017, 36, 933-945.	2.2	24
46	Linking fish species traits to environmental conditions in the Jakarta Bay-Pulau Seribu coral reef system. <i>Marine Pollution Bulletin</i> , 2017, 122, 259-262.	5.0	9
47	Archaeal and bacterial communities of <i>Xestospongia testudinaria</i> and sediment differ in diversity, composition and predicted function in an Indonesian coral reef environment. <i>Journal of Sea Research</i> , 2017, 119, 37-53.	1.6	17
48	Environmental controls on estuarine nitrifying communities along a salinity gradient. <i>Aquatic Microbial Ecology</i> , 2017, 80, 167-180.	1.8	8
49	Temporal dynamics of sediment bacterial communities in monospecific stands of <i>Juncus maritimus</i> and <i>Spartina maritima</i> . <i>Plant Biology</i> , 2016, 18, 824-834.	3.8	13
50	Diversity and composition of plants, butterflies and odonates in an <i>Imperata cylindrica</i> grassland landscape in East Kalimantan, Indonesia. <i>Journal of Tropical Ecology</i> , 2016, 32, 555-560.	1.1	3
51	Biodiversity pattern of subtidal sponges (Porifera: Demospongiae) in the Penghu Archipelago (Pescadores), Taiwan. <i>Journal of the Marine Biological Association of the United Kingdom</i> , 2016, 96, 417-427.	0.8	12
52	Jellyfish-associated bacterial communities and bacterioplankton in Indonesian Marine lakes. <i>FEMS Microbiology Ecology</i> , 2016, 92, fiw064.	2.7	32
53	Variation in the composition of corals, fishes, sponges, echinoderms, ascidians, molluscs, foraminifera and macroalgae across a pronounced in-to-offshore environmental gradient in the Jakarta Bay “Thousand Islands coral reef complex. <i>Marine Pollution Bulletin</i> , 2016, 110, 701-717.	5.0	59
54	Multitaxon activity profiling reveals differential microbial response to reduced seawater pH and oil pollution. <i>Molecular Ecology</i> , 2016, 25, 4645-4659.	3.9	20

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55	Integrated analysis of bacterial and microeukaryotic communities from differentially active mud volcanoes in the Gulf of Cadiz. <i>Scientific Reports</i> , 2016, 6, 35272.	3.3	23
56	Comparison of archaeal and bacterial communities in two sponge species and seawater from an Indonesian coral reef environment. <i>Marine Genomics</i> , 2016, 29, 69-80.	1.1	20
57	Composition and predicted functional ecology of mussel-associated bacteria in Indonesian marine lakes. <i>Antonie Van Leeuwenhoek</i> , 2015, 107, 821-834.	1.7	53
58	Polycyclic aromatic hydrocarbons in deep sea sediments: Microbeâ€“pollutant interactions in a remote environment. <i>Science of the Total Environment</i> , 2015, 526, 312-328.	8.0	99
59	Composition and Predictive Functional Analysis of Bacterial Communities in Seawater, Sediment and Sponges in the Spermonde Archipelago, Indonesia. <i>Microbial Ecology</i> , 2015, 70, 889-903.	2.8	59
60	Molecular Analysis of Skin Bacterial Assemblages from Codfish and Pollock after Dry-Salted Fish Production. <i>Journal of Food Protection</i> , 2015, 78, 983-989.	1.7	4
61	Bacterial community composition and predicted functional ecology of sponges, sediment and seawater from the thousand islands reef complex, West Java, Indonesia. <i>FEMS Microbiology Ecology</i> , 2015, 91, .	2.7	109
62	Habitat and water quality variables as predictors of community composition in an Indonesian coral reef: a multi-taxon study in the Spermonde Archipelago. <i>Science of the Total Environment</i> , 2015, 537, 139-151.	8.0	43
63	The putative functional ecology and distribution of archaeal communities in sponges, sediment and seawater in a coral reef environment. <i>Molecular Ecology</i> , 2015, 24, 409-423.	3.9	44
64	Unraveling the interactive effects of climate change and oil contamination on laboratoryâ€“simulated estuarine benthic communities. <i>Global Change Biology</i> , 2015, 21, 1871-1886.	9.5	28
65	The <sc>PREDICTS</sc> database: a global database of how local terrestrial biodiversity responds to human impacts. <i>Ecology and Evolution</i> , 2014, 4, 4701-4735.	1.9	178
66	Assessing variation in bacterial composition between the rhizospheres of two mangrove tree species. <i>Estuarine, Coastal and Shelf Science</i> , 2014, 139, 40-45.	2.1	30
67	Composition of Archaea in Seawater, Sediment, and Sponges in the Kepulauan Seribu Reef System, Indonesia. <i>Microbial Ecology</i> , 2014, 67, 553-567.	2.8	51
68	Interannual variability in the biochemical composition of newly hatched larvae of the spider crab <i>Maja brachydactyla</i> (Decapoda, Majidae). <i>Marine Ecology</i> , 2014, 35, 298-307.	1.1	8
69	Halophyte plant colonization as a driver of the composition of bacterial communities in salt marshes chronically exposed to oil hydrocarbons. <i>FEMS Microbiology Ecology</i> , 2014, 90, 647-662.	2.7	23
70	Coral reefs next to a major conurbation: a study of temporal change (1985-2011) in coral cover and composition in the reefs of Jakarta, Indonesia. <i>Marine Ecology - Progress Series</i> , 2014, 501, 89-98.	1.9	40
71	The coral-killing sponge <i>Terpios hoshinota</i> invades Indonesia. <i>Coral Reefs</i> , 2013, 32, 755-755.	2.2	34
72	Effect of light, temperature and diet on the fatty acid profile of the tropical sea anemone <i>Aiptasia pallida</i>. <i>Aquaculture Nutrition</i> , 2013, 19, 818-826.	2.7	15

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73	Contrasting habitats occupied by sibling spider crabs <i>Maja squinado</i> and <i>Maja brachydactyla</i> (<i>Brachyura</i> , <i>Majidae</i>) can influence the biochemical variability displayed by newly hatched larvae. <i>Journal of Plankton Research</i> , 2013, 35, 684-688.	1.8	2
74	Richness and composition of sediment bacterial assemblages in an Atlantic port environment. <i>Science of the Total Environment</i> , 2013, 452-453, 172-180.	8.0	16
75	Development and validation of an experimental life support system for assessing the effects of global climate change and environmental contamination on estuarine and coastal marine benthic communities. <i>Global Change Biology</i> , 2013, 19, 2584-2595.	9.5	18
76	El Niño and Biodiversity. , 2013, , 155-163.		1
77	Habitat- and host-related variation in sponge bacterial symbiont communities in Indonesian waters. <i>FEMS Microbiology Ecology</i> , 2013, 85, 465-482.	2.7	87
78	Interactive effects of global climate change and pollution on marine microbes: the way ahead. <i>Ecology and Evolution</i> , 2013, 3, 1808-1818.	1.9	39
79	Lock, Stock and Two Different Barrels: Comparing the Genetic Composition of Morphotypes of the Indo-Pacific Sponge <i>Xestospongia testudinaria</i> . <i>PLoS ONE</i> , 2013, 8, e74396.	2.5	27
80	Molecular Analysis of Bacterial Communities and Detection of Potential Pathogens in a Recirculating Aquaculture System for <i>Scophthalmus maximus</i> and <i>Solea senegalensis</i> . <i>PLoS ONE</i> , 2013, 8, e80847.	2.5	90
81	Sponge species composition, abundance, and cover in marine lakes and coastal mangroves in Berau, Indonesia. <i>Marine Ecology - Progress Series</i> , 2013, 481, 105-120.	1.9	39
82	Ragworm fatty acid profiles reveals habitat and trophic interactions with halophytes and with mercury. <i>Marine Pollution Bulletin</i> , 2012, 64, 2528-2534.	5.0	2
83	Denaturing Gradient Gel Electrophoresis and Barcoded Pyrosequencing Reveal Unprecedented Archaeal Diversity in Mangrove Sediment and Rhizosphere Samples. <i>Applied and Environmental Microbiology</i> , 2012, 78, 5520-5528.	3.1	204
84	Impact of sampling depth and plant species on local environmental conditions, microbiological parameters and bacterial composition in a mercury contaminated salt marsh. <i>Marine Pollution Bulletin</i> , 2012, 64, 263-271.	5.0	16
85	Assessment of Variation in Bacterial Composition among Microhabitats in a Mangrove Environment Using DGGE Fingerprints and Barcoded Pyrosequencing. <i>PLoS ONE</i> , 2012, 7, e29380.	2.5	88
86	Indonesia: Threats to the Country's Biodiversity. , 2011, , 187-197.		10
87	Morphometric variation in two intertidal littorinid gastropods. <i>Contributions To Zoology</i> , 2011, 80, 201-211.	0.5	7
88	Sea surface temperature and ocean colour (MODIS/AQUA) space and time variability in Indonesian Sea coral reef systems from 2002 to 2011. <i>Proceedings of SPIE</i> , 2011, , .	0.8	0
89	Mangrove bacterial richness. <i>Communicative and Integrative Biology</i> , 2011, 4, 419-423.	1.4	35
90	Mangrove bacterial richness. <i>Communicative and Integrative Biology</i> , 2011, 4, 419-23.	1.4	12

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91	Providing a common diet to different marine decapods does not standardize the fatty acid profiles of their larvae: a warning sign for experimentation using invertebrate larvae produced in captivity. <i>Marine Biology</i> , 2010, 157, 2427-2434.	1.5	15
92	Taking Root: Enduring Effect of Rhizosphere Bacterial Colonization in Mangroves. <i>PLoS ONE</i> , 2010, 5, e14065.	2.5	121
93	Butterfly species and traits associated with selectively logged forest in Borneo. <i>Basic and Applied Ecology</i> , 2009, 10, 237-245.	2.7	12
94	Ecology and conservation status of endemic freshwater crabs in Lake Tanganyika, Africa. <i>Biodiversity and Conservation</i> , 2009, 18, 1555-1573.	2.6	23
95	Sponge community composition in the Derawan Islands, NE Kalimantan, Indonesia. <i>Marine Ecology - Progress Series</i> , 2009, 396, 169-180.	1.9	50
96	Ecological correlates of species differences in the Lake Tanganyika crab radiation. <i>Hydrobiologia</i> , 2008, 615, 81-94.	2.0	19
97	A mowing experiment to evaluate the influence of management on the activity of host ants of <i>Maculinea</i> butterflies. <i>Journal of Insect Conservation</i> , 2008, 12, 617-627.	1.4	31
98	Relating variation in species composition to environmental variables: a multi-taxon study in an Indonesian coral reef complex. <i>Aquatic Sciences</i> , 2008, 70, 419-431.	1.5	47
99	An analysis of sponge diversity and distribution at three taxonomic levels in the Thousand Islands/Jakarta Bay reef complex, West Java, Indonesia. <i>Marine Ecology</i> , 2008, 29, 205-215.	1.1	53
100	Relating species traits to environmental variables in Indonesian coral reef sponge assemblages. <i>Marine and Freshwater Research</i> , 2007, 58, 240.	1.3	36
101	Environmental associations of sponges in the Spermonde Archipelago, Indonesia. <i>Journal of the Marine Biological Association of the United Kingdom</i> , 2007, 87, 1669-1676.	0.8	31
102	Analysis of evolutionary, biogeographical and taxonomic patterns of nucleotide composition in demosponge rRNA. <i>Journal of the Marine Biological Association of the United Kingdom</i> , 2007, 87, 1607-1614.	0.8	6
103	BIRD SPECIES AND TRAITS ASSOCIATED WITH LOGGED AND UNLOGGED FOREST IN BORNEO. , 2007, 17, 1184-1197.		97
104	El Niño and Biodiversity. , 2007, , 1-11.		0
105	Sponge diversity and community composition in Irish bathyal coral reefs. <i>Contributions To Zoology</i> , 2007, 76, 121-142.	0.5	56
106	ARES: software to compare allelic richness between uneven samples. <i>Molecular Ecology Notes</i> , 2007, 7, 579-582.	1.7	34
107	Relating coral species traits to environmental conditions in the Jakarta Bay/Pulau Seribu reef system, Indonesia. <i>Estuarine, Coastal and Shelf Science</i> , 2007, 73, 816-826.	2.1	65
108	Relating species traits of foraminifera to environmental variables in the Spermonde Archipelago, Indonesia. <i>Marine Ecology - Progress Series</i> , 2007, 334, 73-82.	1.9	30

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109	Ecological differentiation between the Sardinian endemic <i>Maniola</i> sp. and the pan-European <i>M. jurtina</i> . <i>Biological Journal of the Linnean Society</i> , 2006, 89, 561-574.	1.6	13
110	Beta diversity of tropical marine benthic assemblages in the Spermonde Archipelago, Indonesia. <i>Marine Ecology</i> , 2006, 27, 76-88.	1.1	67
111	Parallel responses of species and genetic diversity to El Niño Southern Oscillation-induced environmental destruction. <i>Ecology Letters</i> , 2006, 9, 304-310.	6.4	63
112	Short-term impact of disturbance on genetic diversity and structure of Indonesian populations of the butterfly <i>Drupadia</i> in East Kalimantan. <i>Molecular Ecology</i> , 2006, 15, 2069-2081.	3.9	14
113	Burning and logging differentially affect endemic vs. widely distributed butterfly species in Borneo. <i>Diversity and Distributions</i> , 2006, 12, 409-416.	4.1	29
114	Butterfly, seedling, sapling and tree diversity and composition in a fire-affected Bornean rainforest. <i>Austral Ecology</i> , 2006, 31, 46-57.	1.5	16
115	Coral diversity across a disturbance gradient in the Pulau Seribu reef complex off Jakarta, Indonesia. <i>Biodiversity and Conservation</i> , 2006, 15, 3653-3674.	2.6	58
116	Diversity Patterns of Bornean Butterfly Assemblages. <i>Biodiversity and Conservation</i> , 2006, 15, 517-538.	2.6	19
117	RANGE-RESTRICTED, SPECIALIST BORNEAN BUTTERFLIES ARE LESS LIKELY TO RECOVER FROM ENSO-INDUCED DISTURBANCE. <i>Ecology</i> , 2006, 87, 2330-2337.	3.2	76
118	Short-Term Impact of 1997/1998 ENSO-Induced Disturbance on Abundance and Genetic Variation in a Tropical Butterfly. <i>Journal of Heredity</i> , 2006, 97, 367-380.	2.4	15
119	Diversity patterns of Bornean butterfly assemblages. , 2006, , 503-524.		4
120	Sponge beta diversity in the Spermonde Archipelago, SW Sulawesi, Indonesia. <i>Marine Ecology - Progress Series</i> , 2006, 309, 131-142.	1.9	78
121	The impact of logging on the abundance, species richness and community composition of butterfly guilds in Borneo. <i>Journal of Applied Entomology</i> , 2005, 129, 52-59.	1.8	11
122	Variation in the diversity and composition of benthic taxa as a function of distance offshore, depth and exposure in the Spermonde Archipelago, Indonesia. <i>Estuarine, Coastal and Shelf Science</i> , 2005, 65, 557-570.	2.1	94
123	Vegetation responses to burning in a rain forest in Borneo. <i>Plant Ecology</i> , 2005, 177, 145-163.	1.6	26
124	Associations of Bird Species Richness and Community Composition with Local and Landscape-scale Environmental Factors in Borneo. <i>Landscape Ecology</i> , 2005, 20, 989-1001.	4.2	45
125	Butterfly, spider, and plant communities in different land-use types in Sardinia, Italy. <i>Biodiversity and Conservation</i> , 2005, 14, 1281-1300.	2.6	47
126	Community composition and species richness of parasitoids infesting <i>Yponomeuta</i> species in the Netherlands. <i>Contributions To Zoology</i> , 2004, 73, 255-261.	0.5	3

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127	Effects of Host Species and Size on Brood Size and Larval Mortality of the Parasitoid, <i>Ageniaspis fuscicollis</i> (Dalman) (Hymenoptera, Encyrtidae). <i>Environmental Entomology</i> , 2004, 33, 528-534.	1.4	7
128	Assessing the Use of Butterflies as Indicators of Logging in Borneo at Three Taxonomic Levels. <i>Journal of Economic Entomology</i> , 2004, 97, 429-435.	1.8	43
129	The Sudden Death of a Coral Reef. <i>Science</i> , 2004, 303, 1293b-1294.	12.6	13
130	Changes in rain forest butterfly diversity following major ENSO-induced fires in Borneo. <i>Global Ecology and Biogeography</i> , 2004, 13, 129-140.	5.8	78
131	Beta diversity of rock-restricted cichlid fishes in Lake Malawi: importance of environmental and spatial factors. <i>Ecography</i> , 2004, 27, 601-610.	4.5	36
132	Diversity and community composition of butterflies and odonates in an ENSO-induced fire affected habitat mosaic: a case study from East Kalimantan, Indonesia. <i>Oikos</i> , 2004, 105, 426-448.	2.7	64
133	Butterfly response to severe ENSO-induced forest fires in Borneo. <i>Ecological Entomology</i> , 2004, 29, 666-676.	2.2	33
134	How does the taxonomic status of allopatric populations influence species richness within African cichlid fish assemblages?. <i>Journal of Biogeography</i> , 2004, 31, 93-102.	3.0	65
135	Butterfly species richness and community composition in forests affected by ENSO-induced burning and habitat isolation in Borneo. <i>Journal of Tropical Ecology</i> , 2004, 20, 359-367.	1.1	25
136	Assessing the Use of Butterflies as Indicators of Logging in Borneo at Three Taxonomic Levels. <i>Journal of Economic Entomology</i> , 2004, 97, 429-435.	1.8	11
137	An examination of scale of assessment, logging and ENSO-induced fires on butterfly diversity in Borneo. <i>Oecologia</i> , 2003, 135, 313-321.	2.0	107
138	Diversity patterns in butterfly communities of the Greek nature reserve Dadia. <i>Biological Conservation</i> , 2003, 114, 427-436.	4.1	48
139	Genetic and ecological differentiation between the butterfly sisterspecies <i>Colias alfacariensis</i> and <i>Colias hyale</i> . <i>Contributions To Zoology</i> , 2002, 71, 131-139.	0.5	5