

Nicole S Webster

List of Publications by Year in descending order

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155
papers

16,580
citations

25423

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121
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docs citations

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times ranked

14393
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#	ARTICLE	IF	CITATIONS
1	Reef location has a greater impact than coral bleaching severity on the microbiome of <i>Pocillopora acuta</i> . <i>Coral Reefs</i> , 2022, 41, 63-79.	0.9	11
2	Comparative metabolomic analysis reveals shared and unique chemical interactions in sponge holobionts. <i>Microbiome</i> , 2022, 10, 22.	4.9	11
3	Novel reference transcriptomes for the sponges <i>Carteriospongia foliascens</i> and <i>Cliona orientalis</i> and associated algal symbiont <i>Gerakladium endoclonum</i> . <i>Coral Reefs</i> , 2021, 40, 9-13.	0.9	3
4	A genomic view of the microbiome of coral reef demosponges. <i>ISME Journal</i> , 2021, 15, 1641-1654.	4.4	67
5	Subcellular view of host-microbiome nutrient exchange in sponges: insights into the ecological success of an early metazoan-microbe symbiosis. <i>Microbiome</i> , 2021, 9, 44.	4.9	32
6	Taxonomic, functional and expression analysis of viral communities associated with marine sponges. <i>PeerJ</i> , 2021, 9, e10715.	0.9	10
7	Microbiome-mediated mechanisms contributing to the environmental tolerance of reef invertebrate species. <i>Marine Biology</i> , 2021, 168, 1.	0.7	19
8	Testing cophylogeny between coral reef invertebrates and their bacterial and archaeal symbionts. <i>Molecular Ecology</i> , 2021, 30, 3768-3782.	2.0	11
9	Benthic micro- and macro-community succession and coral recruitment under overfishing and nutrient enrichment. <i>Ecology</i> , 2021, 102, e03536.	1.5	12
10	Integrating novel tools to elucidate the metabolic basis of microbial symbiosis in reef holobionts. <i>Marine Biology</i> , 2021, 168, 1.	0.7	8
11	Microbial Surface Biofilm Responds to the Growth-Reproduction-Senescence Cycle of the Dominant Coral Reef Macroalgae <i>Sargassum</i> spp.. <i>Life</i> , 2021, 11, 1199.	1.1	0
12	Evidence for genetic structuring and limited dispersal ability in the Great Barrier Reef sponge <i>Carteriospongia foliascens</i> . <i>Coral Reefs</i> , 2020, 39, 39-46.	0.9	5
13	Viral ecogenomics across the Porifera. <i>Microbiome</i> , 2020, 8, 144.	4.9	21
14	Simulated future conditions of ocean warming and acidification disrupt the microbiome of the calcifying foraminifera <i>Marginopora vertebralis</i> across life stages. <i>Environmental Microbiology Reports</i> , 2020, 12, 693-701.	1.0	5
15	Cross-generational effects of climate change on the microbiome of a photosynthetic sponge. <i>Environmental Microbiology</i> , 2020, 22, 4732-4744.	1.8	21
16	Spatial patterns of microbial communities across surface waters of the Great Barrier Reef. <i>Communications Biology</i> , 2020, 3, 442.	2.0	30
17	Assessing the strength and sensitivity of the core microbiota approach on a highly diverse sponge reef. <i>Environmental Microbiology</i> , 2020, 22, 3985-3999.	1.8	12
18	Editorial: special issue on sponge microbiome. <i>FEMS Microbiology Ecology</i> , 2020, 96, .	1.3	4

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19	Visualizing the invisible: class excursions to ignite children's enthusiasm for microbes. <i>Microbial Biotechnology</i> , 2020, 13, 844-887.	2.0	26
20	Coral Reef Microorganisms in a Changing Climate. <i>IScience</i> , 2020, 23, 100972.	1.9	52
21	Comparative genome-centric analysis reveals seasonal variation in the function of coral reef microbiomes. <i>ISME Journal</i> , 2020, 14, 1435-1450.	4.4	40
22	Gene correlation networks reveal the transcriptomic response to elevated nitrogen in a photosynthetic sponge. <i>Molecular Ecology</i> , 2020, 29, 1452-1462.	2.0	4
23	Characterization of a sponge microbiome using an integrative genome-centric approach. <i>ISME Journal</i> , 2020, 14, 1100-1110.	4.4	98
24	Diverse coral reef invertebrates exhibit patterns of phyllosymbiosis. <i>ISME Journal</i> , 2020, 14, 2211-2222.	4.4	43
25	Microbiome dynamics in the tissue and mucus of acroporid corals differ in relation to host and environmental parameters. <i>PeerJ</i> , 2020, 8, e9644.	0.9	30
26	Comparative Genomics Reveals Ecological and Evolutionary Insights into Sponge-Associated <i>Thaumarchaeota</i> . <i>MSystems</i> , 2019, 4, .	1.7	59
27	Characterization of a thaumarchaeal symbiont that drives incomplete nitrification in the tropical sponge <i>Lantheella basta</i> . <i>Environmental Microbiology</i> , 2019, 21, 3831-3854.	1.8	50
28	Thermal stress modifies the marine sponge virome. <i>Environmental Microbiology Reports</i> , 2019, 11, 690-698.	1.0	13
29	Spotlight on how microbes influence their host's behavior. <i>Environmental Microbiology</i> , 2019, 21, 3185-3187.	1.8	2
30	Changes in the metabolic potential of the sponge microbiome under ocean acidification. <i>Nature Communications</i> , 2019, 10, 4134.	5.8	55
31	Marine biofilms constitute a bank of hidden microbial diversity and functional potential. <i>Nature Communications</i> , 2019, 10, 517.	5.8	100
32	Host-Microbe Coevolution: Applying Evidence from Model Systems to Complex Marine Invertebrate Holobionts. <i>MBio</i> , 2019, 10, .	1.8	88
33	Scientists' warning to humanity: microorganisms and climate change. <i>Nature Reviews Microbiology</i> , 2019, 17, 569-586.	13.6	1,138
34	Microbial indicators of environmental perturbations in coral reef ecosystems. <i>Microbiome</i> , 2019, 7, 94.	4.9	126
35	Modularity and predicted functions of the global sponge-microbiome network. <i>Nature Communications</i> , 2019, 10, 992.	5.8	94
36	The urgent need for microbiology literacy in society. <i>Environmental Microbiology</i> , 2019, 21, 1513-1528.	1.8	99

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37	The Effects of Crude Oil and Dispersant on the Larval Sponge Holobiont. <i>MSystems</i> , 2019, 4, .	1.7	11
38	Minimum Information about an Uncultivated Virus Genome (MIUViG). <i>Nature Biotechnology</i> , 2019, 37, 29-37.	9.4	414
39	Disentangling the effect of host-genotype and environment on the microbiome of the coral <i>Acropora tenuis</i> . <i>PeerJ</i> , 2019, 7, e6377.	0.9	60
40	Elucidating the sponge stress response; lipids and fatty acids can facilitate survival under future climate scenarios. <i>Global Change Biology</i> , 2018, 24, 3130-3144.	4.2	32
41	Elevated seawater temperature disrupts the microbiome of an ecologically important bioeroding sponge. <i>Molecular Ecology</i> , 2018, 27, 2124-2137.	2.0	81
42	Reef invertebrate viromics: diversity, host specificity and functional capacity. <i>Environmental Microbiology</i> , 2018, 20, 2125-2141.	1.8	41
43	Microbial conservation in the Anthropocene. <i>Environmental Microbiology</i> , 2018, 20, 1925-1928.	1.8	19
44	Sponges to Be Winners under Near-Future Climate Scenarios. <i>BioScience</i> , 2018, 68, 955-968.	2.2	85
45	In situ responses of the sponge microbiome to ocean acidification. <i>FEMS Microbiology Ecology</i> , 2018, 94, .	1.3	6
46	Establishing microbial baselines to identify indicators of coral reef health. <i>Microbiology Australia</i> , 2018, 39, 42.	0.1	23
47	The bioeroding sponge <i>Cliona orientalis</i> will not tolerate future projected ocean warming. <i>Scientific Reports</i> , 2018, 8, 8302.	1.6	26
48	Exploring the diversity-stability paradigm using sponge microbial communities. <i>Scientific Reports</i> , 2018, 8, 8425.	1.6	66
49	Climate change alterations to ecosystem dominance: how might sponge-dominated reefs function?. <i>Ecology</i> , 2018, 99, 1920-1931.	1.5	56
50	Crown-of-Thorns Sea Star <i>Acanthaster cf. solaris</i> Has Tissue-Characteristic Microbiomes with Potential Roles in Health and Reproduction. <i>Applied and Environmental Microbiology</i> , 2018, 84, .	1.4	29
51	Effects of ocean acidification on the settlement and metamorphosis of marine invertebrate and fish larvae: a review. <i>Marine Ecology - Progress Series</i> , 2018, 606, 237-257.	0.9	54
52	Morphological characterization of virus-like particles in coral reef sponges. <i>PeerJ</i> , 2018, 6, e5625.	0.9	27
53	Evaluating the core microbiota in complex communities: A systematic investigation. <i>Environmental Microbiology</i> , 2017, 19, 1450-1462.	1.8	187
54	Microbiome analysis of a disease affecting the deep-sea sponge <i>Geodia barretti</i> . <i>FEMS Microbiology Ecology</i> , 2017, 93, .	1.3	36

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55	Microbial indicators as a diagnostic tool for assessing water quality and climate stress in coral reef ecosystems. <i>Marine Biology</i> , 2017, 164, 1.	0.7	101
56	Prevalent and persistent viral infection in cultures of the coral algal endosymbiont <i>Symbiodinium</i> . <i>Coral Reefs</i> , 2017, 36, 773-784.	0.9	36
57	Conceptual and methodological advances for holobiont research. <i>Environmental Microbiology Reports</i> , 2017, 9, 30-32.	1.0	2
58	Sponge Disease and Climate Change. , 2017, , 411-428.		15
59	A communal catalogue reveals Earth's multiscale microbial diversity. <i>Nature</i> , 2017, 551, 457-463.	13.7	1,942
60	The sponge microbiome project. <i>GigaScience</i> , 2017, 6, 1-7.	3.3	193
61	Redefining the sponge-symbiont acquisition paradigm: sponge microbes exhibit chemotaxis towards host-derived compounds. <i>Environmental Microbiology Reports</i> , 2017, 9, 750-755.	1.0	20
62	The contribution of microbial biotechnology to mitigating coral reef degradation. <i>Microbial Biotechnology</i> , 2017, 10, 1236-1243.	2.0	101
63	A decadal analysis of bioeroding sponge cover on the inshore Great Barrier Reef. <i>Scientific Reports</i> , 2017, 7, 2706.	1.6	19
64	Effects of combined dredging-related stressors on sponges: a laboratory approach using realistic scenarios. <i>Scientific Reports</i> , 2017, 7, 5155.	1.6	16
65	The response of a boreal deep-sea sponge holobiont to acute thermal stress. <i>Scientific Reports</i> , 2017, 7, 1660.	1.6	67
66	Effects of suspended sediments on the sponge holobiont with implications for dredging management. <i>Scientific Reports</i> , 2017, 7, 4925.	1.6	52
67	Effects of sediment smothering on the sponge holobiont with implications for dredging management. <i>Scientific Reports</i> , 2017, 7, 5156.	1.6	29
68	Interactive effects of temperature and CO_2 on sponges: from the cradle to the grave. <i>Global Change Biology</i> , 2017, 23, 2031-2046.	4.2	79
69	Predicting the HMA-LMA Status in Marine Sponges by Machine Learning. <i>Frontiers in Microbiology</i> , 2017, 8, 752.	1.5	175
70	Microbial contributions to the persistence of coral reefs. <i>ISME Journal</i> , 2017, 11, 2167-2174.	4.4	173
71	Sediment tolerance mechanisms identified in sponges using advanced imaging techniques. <i>PeerJ</i> , 2017, 5, e3904.	0.9	19
72	Coral-associated viral communities show high levels of diversity and host auxiliary functions. <i>PeerJ</i> , 2017, 5, e4054.	0.9	34

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73	HoloVir: A Workflow for Investigating the Diversity and Function of Viruses in Invertebrate Holobionts. <i>Frontiers in Microbiology</i> , 2016, 7, 822.	1.5	49
74	Host-associated coral reef microbes respond to the cumulative pressures of ocean warming and ocean acidification. <i>Scientific Reports</i> , 2016, 6, 19324.	1.6	161
75	Effects of light attenuation on the sponge holobiont- implications for dredging management. <i>Scientific Reports</i> , 2016, 6, 39038.	1.6	56
76	The Sponge Hologenome. <i>MBio</i> , 2016, 7, e00135-16.	1.8	269
77	Insights into the Coral Microbiome: Underpinning the Health and Resilience of Reef Ecosystems. <i>Annual Review of Microbiology</i> , 2016, 70, 317-340.	2.9	600
78	Diversity, structure and convergent evolution of the global sponge microbiome. <i>Nature Communications</i> , 2016, 7, 11870.	5.8	594
79	Acute ecotoxicology of natural oil and gas condensate to coral reef larvae. <i>Scientific Reports</i> , 2016, 6, 21153.	1.6	58
80	Appearance matters: sedimentation effects on different sponge morphologies. <i>Journal of the Marine Biological Association of the United Kingdom</i> , 2016, 96, 481-492.	0.4	30
81	Using a thermistor flowmeter with attached video camera for monitoring sponge excurrent speed and oscular behaviour. <i>PeerJ</i> , 2016, 4, e2761.	0.9	33
82	Increased seawater temperature increases the abundance and alters the structure of natural <i>Vibrio</i> populations associated with the coral <i>Pocillopora damicornis</i> . <i>Frontiers in Microbiology</i> , 2015, 6, 432.	1.5	142
83	Chemotaxis by natural populations of coral reef bacteria. <i>ISME Journal</i> , 2015, 9, 1764-1777.	4.4	60
84	Natural volcanic CO ₂ seeps reveal future trajectories for host-microbial associations in corals and sponges. <i>ISME Journal</i> , 2015, 9, 894-908.	4.4	268
85	The Pathogen of the Great Barrier Reef Sponge <i>Rhopaloeides odorabile</i> Is a New Strain of <i>Pseudoalteromonas agarivorans</i> Containing Abundant and Diverse Virulence-Related Genes. <i>Marine Biotechnology</i> , 2015, 17, 463-478.	1.1	51
86	Biogeographic variation in the microbiome of the ecologically important sponge, <i>Carteriospongia foliascens</i> . <i>PeerJ</i> , 2015, 3, e1435.	0.9	42
87	Cooperation, communication, and co-evolution: grand challenges in microbial symbiosis research. <i>Frontiers in Microbiology</i> , 2014, 5, 164.	1.5	30
88	Eutrophication has no short-term effect on the <i>Cymbastela stipitata</i> holobiont. <i>Frontiers in Microbiology</i> , 2014, 5, 216.	1.5	60
89	Draft Genome Sequence of <i>Pseudoalteromonas</i> sp. Strain NW 4327 (MTCC 11073, DSM 25418), a Pathogen of the Great Barrier Reef Sponge <i>Rhopaloeides odorabile</i> . <i>Genome Announcements</i> , 2014, 2, .	0.8	6
90	Sponge larval settlement cues: the role of microbial biofilms in a warming ocean. <i>Scientific Reports</i> , 2014, 4, 4072.	1.6	93

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91	Phenology of sexual reproduction in the common coral reef sponge, <i>Carteriospongia foliascens</i> . <i>Coral Reefs</i> , 2014, 33, 381.	0.9	20
92	Variability in Microbial Community Composition and Function Between Different Niches Within a Coral Reef. <i>Microbial Ecology</i> , 2014, 67, 540-552.	1.4	68
93	The influence of habitat on post-settlement processes, larval production and recruitment in a common coral reef sponge. <i>Journal of Experimental Marine Biology and Ecology</i> , 2014, 461, 162-172.	0.7	21
94	Down under the tunic: bacterial biodiversity hotspots and widespread ammonia-oxidizing archaea in coral reef ascidians. <i>ISME Journal</i> , 2014, 8, 575-588.	4.4	88
95	Combining morphometrics with molecular taxonomy: How different are similar foliose keratose sponges from the Australian tropics?. <i>Molecular Phylogenetics and Evolution</i> , 2014, 73, 23-39.	1.2	20
96	Larval Behaviours and Their Contribution to the Distribution of the Intertidal Coral Reef Sponge <i>Carteriospongia foliascens</i> . <i>PLoS ONE</i> , 2014, 9, e98181.	1.1	30
97	Ocean acidification reduces induction of coral settlement by crustose coralline algae. <i>Global Change Biology</i> , 2013, 19, 303-315.	4.2	125
98	Small core communities and high variability in bacteria associated with the introduced ascidian <i>Styela plicata</i> . <i>Symbiosis</i> , 2013, 59, 35-46.	1.2	24
99	A complex life cycle in a warming planet: gene expression in thermally stressed sponges. <i>Molecular Ecology</i> , 2013, 22, 1854-1868.	2.0	59
100	Marine microbial symbiosis heats up: the phylogenetic and functional response of a sponge holobiont to thermal stress. <i>ISME Journal</i> , 2013, 7, 991-1002.	4.4	266
101	Near-future ocean acidification causes differences in microbial associations within diverse coral reef taxa. <i>Environmental Microbiology Reports</i> , 2013, 5, 243-251.	1.0	64
102	Could some coral reefs become sponge reefs as our climate changes?. <i>Global Change Biology</i> , 2013, 19, 2613-2624.	4.2	261
103	Coral Reef Bacterial Communities. , 2013, , 163-187.		58
104	Coral reef invertebrate microbiomes correlate with the presence of photosymbionts. <i>ISME Journal</i> , 2013, 7, 1452-1458.	4.4	146
105	"Sponge-specific" bacteria are widespread (but rare) in diverse marine environments. <i>ISME Journal</i> , 2013, 7, 438-443.	4.4	161
106	Cryptic speciation and phylogeographic relationships in the elephant ear sponge <i>Lanthella basta</i> (Porifera, Lanthellidae) from northern Australia. <i>Zoological Journal of the Linnean Society</i> , 2012, 166, 225-235.	1.0	22
107	Assessing the complex sponge microbiota: core, variable and species-specific bacterial communities in marine sponges. <i>ISME Journal</i> , 2012, 6, 564-576.	4.4	508
108	Functional equivalence and evolutionary convergence in complex communities of microbial sponge symbionts. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, E1878-87.	3.3	361

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109	Thermal stress responses in the bacterial biosphere of the Great Barrier Reef sponge, <i>Rhopaloeides odorabile</i> . Environmental Microbiology, 2012, 14, 3232-3246.	1.8	93
110	Qualitative variation in colour morphotypes of lanthella basta (Porifera: Verongida). Hydrobiologia, 2012, 687, 191-203.	1.0	12
111	The marine sponge lanthella basta can recover from stress-induced tissue regression. Hydrobiologia, 2012, 687, 227-235.	1.0	26
112	Behavioral and morphological changes caused by thermal stress in the Great Barrier Reef sponge <i>Rhopaloeides odorabile</i> . Journal of Experimental Marine Biology and Ecology, 2012, 416-417, 55-60.	0.7	57
113	Marine sponges and their microbial symbionts: love and other relationships. Environmental Microbiology, 2012, 14, 335-346.	1.8	491
114	Sponge-specific clusters revisited: a comprehensive phylogeny of sponge-associated microorganisms. Environmental Microbiology, 2012, 14, 517-524.	1.8	253
115	Same, same but different: symbiotic bacterial associations in GBR sponges. Frontiers in Microbiology, 2012, 3, 444.	1.5	52
116	Crustose Coralline Algae and a Cnidarian Neuropeptide Trigger Larval Settlement in Two Coral Reef Sponges. PLoS ONE, 2012, 7, e30386.	1.1	48
117	Thermal and Sedimentation Stress Are Unlikely Causes of Brown Spot Syndrome in the Coral Reef Sponge, lanthella basta. PLoS ONE, 2012, 7, e39779.	1.1	58
118	Sponge-Microbe Associations Survive High Nutrients and Temperatures. PLoS ONE, 2012, 7, e52220.	1.1	72
119	The larval sponge holobiont exhibits high thermal tolerance. Environmental Microbiology Reports, 2011, 3, 756-762.	1.0	66
120	Strict thermal threshold identified by quantitative PCR in the sponge <i>Rhopaloeides odorabile</i> . Marine Ecology - Progress Series, 2011, 431, 97-105.	0.9	37
121	Elevated seawater temperature causes a microbial shift on crustose coralline algae with implications for the recruitment of coral larvae. ISME Journal, 2011, 5, 759-770.	4.4	145
122	Bacterial Community Dynamics in the Marine Sponge <i>Rhopaloeides odorabile</i> Under In Situ and Ex Situ Cultivation. Marine Biotechnology, 2011, 13, 296-304.	1.1	52
123	The marine sponge lanthella basta can recover from stress-induced tissue regression. , 2011, , 227-235.		2
124	Incongruence between the distribution of a common coral reef sponge and photosynthesis. Marine Ecology - Progress Series, 2011, 423, 95-100.	0.9	8
125	Qualitative variation in colour morphotypes of lanthella basta (Porifera: Verongida). , 2011, , 191-203.		0
126	Deep sequencing reveals exceptional diversity and modes of transmission for bacterial sponge symbionts. Environmental Microbiology, 2010, 12, 2070-2082.	1.8	394

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127	Exploring the Role of Microorganisms in the Disease-Like Syndrome Affecting the Sponge <i>Lanthella basta</i> . Applied and Environmental Microbiology, 2010, 76, 5736-5744.	1.4	56
128	Prevalence of tissue necrosis and brown spot lesions in a common marine sponge. Marine and Freshwater Research, 2010, 61, 484.	0.7	25
129	Impact of global climate change on marine bacterial symbioses and disease. Microbiology Australia, 2009, 30, 78.	0.1	1
130	Purification and Characterization of a Collagenolytic Enzyme from a Pathogen of the Great Barrier Reef Sponge, <i>Rhopaloeides odorabile</i> . PLoS ONE, 2009, 4, e7177.	1.1	35
131	What do we really know about sponge-microbial symbioses?. ISME Journal, 2009, 3, 1-3.	4.4	92
132	<i>Bacillus insecticides</i> are not acutely harmful to corals and sponges. Marine Ecology - Progress Series, 2009, 381, 157-165.	0.9	7
133	Temperature thresholds for bacterial symbiosis with a sponge. ISME Journal, 2008, 2, 830-842.	4.4	226
134	Shifts in microbial and chemical patterns within the marine sponge <i>Aplysina aerophoba</i> during a disease outbreak. Environmental Microbiology, 2008, 10, 3366-3376.	1.8	112
135	Microbiological aspects of phyllosoma rearing of the ornate rock lobster <i>Panulirus ornatus</i> . Aquaculture, 2007, 268, 274-287.	1.7	31
136	Sponge disease: a global threat?. Environmental Microbiology, 2007, 9, 1363-1375.	1.8	190
137	Bacterial community structure associated with the Antarctic soft coral, <i>Alcyonium antarcticum</i> . FEMS Microbiology Ecology, 2007, 59, 81-94.	1.3	83
138	Vibrionaceae infection in phyllosomas of the tropical rock lobster <i>Panulirus ornatus</i> as detected by fluorescence in situ hybridisation. Aquaculture, 2006, 255, 173-178.	1.7	17
139	Biofilm development within a larval rearing tank of the tropical rock lobster, <i>Panulirus ornatus</i> . Aquaculture, 2006, 260, 27-38.	1.7	51
140	Contamination in sediments, bivalves and sponges of McMurdo Sound, Antarctica. Environmental Pollution, 2006, 143, 456-467.	3.7	145
141	Site-specific variation in Antarctic marine biofilms established on artificial surfaces. Environmental Microbiology, 2006, 8, 1177-1190.	1.8	80
142	Recruitment of Antarctic marine eukaryotes onto artificial surfaces. Polar Biology, 2006, 30, 1-10.	0.5	7
143	Diverse microbial communities inhabit Antarctic sponges. Environmental Microbiology, 2004, 6, 288-300.	1.8	251
144	TBT contamination identified in Antarctic marine sediments. Marine Pollution Bulletin, 2004, 48, 1142-1144.	2.3	51

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145	Metamorphosis of a Scleractinian Coral in Response to Microbial Biofilms. <i>Applied and Environmental Microbiology</i> , 2004, 70, 1213-1221.	1.4	287
146	Microbial community dynamics in a larval aquaculture system of the tropical rock lobster, <i>Panulirus ornatus</i> . <i>Aquaculture</i> , 2004, 242, 31-51.	1.7	67
147	The effects of antifoulant-paint-contaminated sediments on coral recruits and branchlets. <i>Marine Biology</i> , 2003, 143, 651-657.	0.7	42
148	PARALYTIC SHELLFISH TOXINS ARE RESTRICTED TO FEW SPECIES AMONG AUSTRALIA'S TAXONOMIC DIVERSITY OF CULTURED MICROALGAE1. <i>Journal of Phycology</i> , 2003, 39, 663-667.	1.0	33
149	Understanding ship-grounding impacts on a coral reef: potential effects of anti-foulant paint contamination on coral recruitment. <i>Marine Pollution Bulletin</i> , 2002, 44, 111-117.	2.3	98
150	A spongin-boring α -proteobacterium is the etiological agent of disease in the Great Barrier Reef sponge <i>Rhopaloeides odorabile</i> . <i>Marine Ecology - Progress Series</i> , 2002, 232, 305-309.	0.9	110
151	Phylogenetic Diversity of Bacteria Associated with the Marine Sponge <i>Rhopaloeides odorabile</i> . <i>Applied and Environmental Microbiology</i> , 2001, 67, 434-444.	1.4	322
152	The culturable microbial community of the Great Barrier Reef sponge <i>Rhopaloeides odorabile</i> is dominated by an α -Proteobacterium. <i>Marine Biology</i> , 2001, 138, 843-851.	0.7	247
153	The effects of copper on the microbial community of a coral reef sponge. <i>Environmental Microbiology</i> , 2001, 3, 19-31.	1.8	95
154	Detection and Phylogenetic Analysis of Novel Crenarchaeote and Euryarchaeote 16S Ribosomal RNA Gene Sequences from a Great Barrier Reef Sponge. <i>Marine Biotechnology</i> , 2001, 3, 0600-0608.	1.1	73
155	Metamorphosis of broadcast spawning corals in response to bacteria isolated from crustose algae. <i>Marine Ecology - Progress Series</i> , 2001, 223, 121-131.	0.9	213