

Newton Gomes

List of Publications by Year in descending order

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137
papers

6,399
citations

57758

44
h-index

74163

75
g-index

141
all docs

141
docs citations

141
times ranked

8176
citing authors

#	ARTICLE	IF	CITATIONS
1	Antimicrobial Photodynamic Therapy: Study of Bacterial Recovery Viability and Potential Development of Resistance after Treatment. <i>Marine Drugs</i> , 2010, 8, 91-105.	4.6	340
2	Title is missing!. <i>Plant and Soil</i> , 2001, 232, 167-180.	3.7	266
3	Dynamics of Fungal Communities in Bulk and Maize Rhizosphere Soil in the Tropics. <i>Applied and Environmental Microbiology</i> , 2003, 69, 3758-3766.	3.1	248
4	Trends in the Discovery of New Marine Natural Products from Invertebrates over the Last Two Decades – Where and What Are We Bioprospecting?. <i>PLoS ONE</i> , 2012, 7, e30580.	2.5	217
5	Cnidarians as a Source of New Marine Bioactive Compounds – An Overview of the Last Decade and Future Steps for Bioprospecting. <i>Marine Drugs</i> , 2011, 9, 1860-1886.	4.6	210
6	Wavelength dependence of biological damage induced by UV radiation on bacteria. <i>Archives of Microbiology</i> , 2013, 195, 63-74.	2.2	205
7	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
8	Phage Therapy and Photodynamic Therapy: Low Environmental Impact Approaches to Inactivate Microorganisms in Fish Farming Plants. <i>Marine Drugs</i> , 2009, 7, 268-313.	4.6	127
9	Exploring the diversity of bacterial communities in sediments of urban mangrove forests. <i>FEMS Microbiology Ecology</i> , 2008, 66, 96-109.	2.7	124
10	Rhizosphere Communities of Genetically Modified Zeaxanthin-Accumulating Potato Plants and Their Parent Cultivar Differ Less than Those of Different Potato Cultivars. <i>Applied and Environmental Microbiology</i> , 2009, 75, 3859-3865.	3.1	122
11	Taking Root: Enduring Effect of Rhizosphere Bacterial Colonization in Mangroves. <i>PLoS ONE</i> , 2010, 5, e14065.	2.5	121
12	Phage Therapy as an Approach to Prevent <i>Vibrio anguillarum</i> Infections in Fish Larvae Production. <i>PLoS ONE</i> , 2014, 9, e114197.	2.5	117
13	Effects of the inoculant strain <i>Pseudomonas putida</i> KT2442 (pNF142) and of naphthalene contamination on the soil bacterial community. <i>FEMS Microbiology Ecology</i> , 2005, 54, 21-33.	2.7	110
14	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
15	Intracellular Signal Triggered by Cholera Toxin in <i>Saccharomyces boulardii</i> and <i>Saccharomyces cerevisiae</i> . <i>Applied and Environmental Microbiology</i> , 1998, 64, 564-568.	3.1	109
16	Mechanisms of photodynamic inactivation of a Gram-negative recombinant bioluminescent bacterium by cationic porphyrins. <i>Photochemical and Photobiological Sciences</i> , 2011, 10, 1659-1669.	2.9	106
17	Polycyclic aromatic hydrocarbons in deep sea sediments: Microbe – pollutant interactions in a remote environment. <i>Science of the Total Environment</i> , 2015, 526, 312-328.	8.0	99
18	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

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19	Evaluation of resistance development and viability recovery by a non-enveloped virus after repeated cycles of aPDT. <i>Antiviral Research</i> , 2011, 91, 278-282.	4.1	89
20	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
21	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
22	<i>Pseudomonas</i> community structure and antagonistic potential in the rhizosphere: insights gained by combining phylogenetic and functional gene-based analyses. <i>Environmental Microbiology</i> , 2007, 9, 2260-2273.	3.8	86
23	The sponge microbiome within the greater coral reef microbial metacommunity. <i>Nature Communications</i> , 2019, 10, 1644.	12.8	86
24	Survival of gfp-tagged antagonistic bacteria in the rhizosphere of tomato plants and their effects on the indigenous bacterial community. <i>FEMS Microbiology Ecology</i> , 2006, 56, 207-218.	2.7	83
25	Photodynamic inactivation of <i>Penicillium chrysogenum</i> conidia by cationic porphyrins. <i>Photochemical and Photobiological Sciences</i> , 2011, 10, 1735-1743.	2.9	82
26	Impact of organic and inorganic nanomaterials in the soil microbial community structure. <i>Science of the Total Environment</i> , 2012, 424, 344-350.	8.0	80
27	A new semi-nested PCR protocol to amplify large 18S rRNA gene fragments for PCR-DGGE analysis of soil fungal communities. <i>Journal of Microbiological Methods</i> , 2006, 65, 63-75.	1.6	75
28	Diversity of ndo Genes in Mangrove Sediments Exposed to Different Sources of Polycyclic Aromatic Hydrocarbon Pollution. <i>Applied and Environmental Microbiology</i> , 2007, 73, 7392-7399.	3.1	74
29	Phage therapy to control multidrug-resistant <i>Pseudomonas aeruginosa</i> skin infections: in vitro and ex vivo experiments. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2012, 31, 3241-3249.	2.9	73
30	Bacteriophages with Potential for Inactivation of Fish Pathogenic Bacteria: Survival, Host Specificity and Effect on Bacterial Community Structure. <i>Marine Drugs</i> , 2011, 9, 2236-2255.	4.6	72
31	Biological control of <i>Aeromonas salmonicida</i> infection in juvenile Senegalese sole (<i>Solea</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 5	3.5	71
32	Contribution of reactive oxygen species to UV-B-induced damage in bacteria. <i>Journal of Photochemistry and Photobiology B: Biology</i> , 2012, 117, 40-46.	3.8	70
33	Short-term effects of amoxicillin on bacterial communities in manured soil. <i>FEMS Microbiology Ecology</i> , 2007, 62, 290-302.	2.7	68
34	Photodynamic Antimicrobial Chemotherapy in Aquaculture: Photoinactivation Studies of <i>Vibrio fischeri</i> . <i>PLoS ONE</i> , 2011, 6, e20970.	2.5	67
35	Diversity and antagonistic potential of <i>Pseudomonas</i> spp. associated to the rhizosphere of maize grown in a subtropical organic farm. <i>Soil Biology and Biochemistry</i> , 2006, 38, 2434-2447.	8.8	64
36	The impact of shrimp farming effluent on bacterial communities in mangrove waters, CearÃ¡, Brazil. <i>Marine Pollution Bulletin</i> , 2006, 52, 1725-1734.	5.0	62

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37	Influence of environmental variables in the efficiency of phage therapy in aquaculture. <i>Microbial Biotechnology</i> , 2014, 7, 401-413.	4.2	62
38	Mangrove microniches determine the structural and functional diversity of enriched petroleum hydrocarbon-degrading consortia. <i>FEMS Microbiology Ecology</i> , 2010, 74, 276-290.	2.7	61
39	Phthalocyanine Thioâ€Pyridinium Derivatives as Antibacterial Photosensitizers^{â€}. <i>Photochemistry and Photobiology</i> , 2012, 88, 537-547.	2.5	60
40	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
41	Hydrocarbon contamination and plant species determine the phylogenetic and functional diversity of endophytic degrading bacteria. <i>Molecular Ecology</i> , 2014, 23, 1392-1404.	3.9	55
42	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
43	Occurrence and diversity of naphthalene dioxygenase genes in soil microbial communities from the Maritime Antarctic. <i>Environmental Microbiology</i> , 2009, 11, 700-714.	3.8	51
44	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
45	Effects of UV-B Radiation on the Structural and Physiological Diversity of Bacterioneuston and Bacterioplankton. <i>Applied and Environmental Microbiology</i> , 2012, 78, 2066-2069.	3.1	48
46	Effects of Cd- and Zn-enriched sewage sludge on soil bacterial and fungal communities. <i>Ecotoxicology and Environmental Safety</i> , 2010, 73, 1255-1263.	6.0	45
47	Evidence for Selective Bacterial Community Structuring in the Freshwater Sponge <i>Ephydatia fluviatilis</i> . <i>Microbial Ecology</i> , 2013, 65, 232-244.	2.8	44
48	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
49	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
50	Nucleic acid changes during photodynamic inactivation of bacteria by cationic porphyrins. <i>Bioorganic and Medicinal Chemistry</i> , 2013, 21, 4311-4318.	3.0	42
51	Biodegradation of 17 β -estradiol by bacteria isolated from deep sea sediments in aerobic and anaerobic media. <i>Journal of Hazardous Materials</i> , 2017, 323, 359-366.	12.4	42
52	Evaluating seasonal dynamics of bacterial communities in marine fish aquaculture: a preliminary study before applying phage therapy. <i>Journal of Environmental Monitoring</i> , 2011, 13, 1053.	2.1	41
53	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
54	Microbe-Assisted Phytoremediation of Hydrocarbons in Estuarine Environments. <i>Microbial Ecology</i> , 2015, 69, 1-12.	2.8	38

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55	Applicability of photodynamic antimicrobial chemotherapy as an alternative to inactivate fish pathogenic bacteria in aquaculture systems. <i>Photochemical and Photobiological Sciences</i> , 2011, 10, 1691-1700.	2.9	36
56	Mangrove bacterial richness. <i>Communicative and Integrative Biology</i> , 2011, 4, 419-423.	1.4	35
57	The gorgonian coral <i>Eunicella labiata</i> hosts a distinct prokaryotic consortium amenable to cultivation. <i>FEMS Microbiology Ecology</i> , 2017, 93, .	2.7	35
58	Photodynamic oxidation of <i>Staphylococcus warneri</i> membrane phospholipids: new insights based on lipidomics. <i>Rapid Communications in Mass Spectrometry</i> , 2013, 27, 1607-1618.	1.5	34
59	Jellyfish-associated bacterial communities and bacterioplankton in Indonesian Marine lakes. <i>FEMS Microbiology Ecology</i> , 2016, 92, fiw064.	2.7	32
60	Compositional analysis of bacterial communities in seawater, sediment, and sponges in the Misool coral reef system, Indonesia. <i>Marine Biodiversity</i> , 2018, 48, 1889-1901.	1.0	32
61	Bacteriome Structure, Function, and Probiotics in Fish Larviculture: The Good, the Bad, and the Gaps. <i>Annual Review of Animal Biosciences</i> , 2021, 9, 423-452.	7.4	31
62	Short-term variability of abundance, diversity and activity of estuarine bacterioplankton and bacterioplankton. <i>Journal of Plankton Research</i> , 2009, 31, 1545-1555.	1.8	30
63	Ultracentrifugation as a direct method to concentrate viruses in environmental waters: virus-like particle enumeration as a new approach to determine the efficiency of recovery. <i>Journal of Environmental Monitoring</i> , 2012, 14, 64-70.	2.1	30
64	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
65	Bacterial and microeukaryotic plankton communities in a semi-intensive aquaculture system of sea bass (<i>Dicentrarchus labrax</i>): A seasonal survey. <i>Aquaculture</i> , 2019, 503, 59-69.	3.5	29
66	Unraveling the interactive effects of climate change and oil contamination on laboratory-simulated estuarine benthic communities. <i>Global Change Biology</i> , 2015, 21, 1871-1886.	9.5	28
67	An optimized protocol for simultaneous extraction of DNA and RNA from soils. <i>Brazilian Journal of Microbiology</i> , 2004, 35, 230-234.	2.0	25
68	Effects of Monospecific Banks of Salt Marsh Vegetation on Sediment Bacterial Communities. <i>Microbial Ecology</i> , 2010, 60, 167-179.	2.8	25
69	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
70	Relation between bacterial activity in the surface microlayer and estuarine hydrodynamics. <i>FEMS Microbiology Ecology</i> , 2011, 77, 636-646.	2.7	24
71	A bacterial consortium isolated from an Icelandic fumarole displays exceptionally high levels of sulfate reduction and metals resistance. <i>Journal of Hazardous Materials</i> , 2011, 187, 362-370.	12.4	24
72	Prokaryotes in salt marsh sediments of Ria de Aveiro: Effects of halophyte vegetation on abundance and diversity. <i>Estuarine, Coastal and Shelf Science</i> , 2012, 110, 61-68.	2.1	24

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73	Uptake of free and complexed silver ions by different strains of <i>Rhodotorula mucilaginosa</i> . <i>Brazilian Journal of Microbiology</i> , 2002, 33, 62-66.	2.0	23
74	Diversity in UV sensitivity and recovery potential among bacterioneuston and bacterioplankton isolates. <i>Letters in Applied Microbiology</i> , 2011, 52, 360-366.	2.2	23
75	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
76	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
77	<i>Clavispora opuntiae</i> and other yeasts associated with the moth <i>Sigelgaita</i> sp. in the cactus <i>Pilosocereus arrabidaei</i> of Rio de Janeiro, Brazil. <i>Antonie Van Leeuwenhoek</i> , 1992, 62, 267-272.	1.7	21
78	The UV responses of bacterioneuston and bacterioplankton isolates depend on the physiological condition and involve a metabolic shift. <i>FEMS Microbiology Ecology</i> , 2012, 80, 646-658.	2.7	21
79	Can Volatile Organic Metabolites Be Used to Simultaneously Assess Microbial and Mite Contamination Level in Cereal Grains and Coffee Beans?. <i>PLoS ONE</i> , 2013, 8, e59338.	2.5	21
80	Impact of freshwater inflow on bacterial abundance and activity in the estuarine system Ria de Aveiro. <i>Estuarine, Coastal and Shelf Science</i> , 2014, 138, 107-120.	2.1	20
81	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
82	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
83	An ecotoxicological analysis of the sediment quality in a European Atlantic harbor emphasizes the current limitations of the Water Framework Directive. <i>Marine Pollution Bulletin</i> , 2013, 72, 197-204.	5.0	19
84	Isolation of Surfactant-Resistant <i>Pseudomonads</i> from the Estuarine Surface Microlayer. <i>Journal of Microbiology and Biotechnology</i> , 2012, 22, 283-291.	2.1	19
85	The performance of <i>Fraxinus angustifolia</i> as a helper for metal phytoremediation programs and its relation to the endophytic bacterial communities. <i>Geoderma</i> , 2013, 202-203, 171-182.	5.1	18
86	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
87	Bacterial production of biosurfactants under microaerobic and anaerobic conditions. <i>Reviews in Environmental Science and Biotechnology</i> , 2017, 16, 239-272.	8.1	17
88	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
89	Seasonal patterns of bacterioplankton composition in a semi-intensive European seabass (<i>Dicentrarchus labrax</i>) aquaculture system. <i>Aquaculture</i> , 2018, 490, 240-250.	3.5	17
90	Sponge Prokaryote Communities in Taiwanese Coral Reef and Shallow Hydrothermal Vent Ecosystems. <i>Microbial Ecology</i> , 2018, 75, 239-254.	2.8	17

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91	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
92	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
93	Title is missing!. <i>World Journal of Microbiology and Biotechnology</i> , 1997, 14, 149-149.	3.6	15
94	Characterization of bacterioplankton communities from a hatchery recirculating aquaculture system (RAS) for juvenile sole (<i>Solea senegalensis</i>) production. <i>PLoS ONE</i> , 2019, 14, e0211209.	2.5	15
95	Effects of ultraviolet radiation on the abundance, diversity and activity of bacterioneuston and bacterioplankton: insights from microcosm studies. <i>Aquatic Sciences</i> , 2011, 73, 63-77.	1.5	14
96	Optimization of preservation and processing of sea anemones for microbial community analysis using molecular tools. <i>Scientific Reports</i> , 2014, 4, 6986.	3.3	13
97	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
98	Exploring hydrocarbonoclastic bacterial communities in the estuarine surface microlayer. <i>Aquatic Microbial Ecology</i> , 2011, 64, 185-195.	1.8	12
99	Mangrove bacterial richness. <i>Communicative and Integrative Biology</i> , 2011, 4, 419-23.	1.4	12
100	Effect of chemical stress and ultraviolet radiation in the bacterial communities of zebrafish embryos. <i>Environmental Pollution</i> , 2016, 208, 626-636.	7.5	11
101	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
102	Humic substances modulate fish bacterial communities in a marine recirculating aquaculture system. <i>Aquaculture</i> , 2021, 544, 737121.	3.5	11
103	Cyano-metal complexes uptake by <i>Aspergillus niger</i> . <i>Biotechnology Letters</i> , 1999, 21, 487-490.	2.2	10
104	Thiocyanate degradation by pure and mixed cultures of microorganisms. <i>Brazilian Journal of Microbiology</i> , 2004, 35, 333-336.	2.0	10
105	Development of a molecular methodology for fast detection of <i>Photobacterium damsela</i> subspecies in water samples. <i>Aquaculture</i> , 2015, 435, 137-142.	3.5	10
106	Micro-eukaryotic plankton diversity in an intensive aquaculture system for production of <i>Scophthalmus maximus</i> and <i>Solea senegalensis</i> . <i>Aquaculture</i> , 2018, 490, 321-328.	3.5	10
107	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
108	Role of Transition Metals in UV-Induced Damage to Bacteria. <i>Photochemistry and Photobiology</i> , 2013, 89, 640-648.	2.5	9

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109	Heterotrophic activities of neustonic and planktonic bacterial communities in an estuarine environment (Ria de Aveiro). <i>Journal of Plankton Research</i> , 2014, 36, 230-242.	1.8	9
110	Microcosm evaluation of the impact of oil contamination and chemical dispersant addition on bacterial communities and sediment remediation of an estuarine port environment. <i>Journal of Applied Microbiology</i> , 2019, 127, 134-149.	3.1	9
111	Title is missing!. <i>World Journal of Microbiology and Biotechnology</i> , 2000, 16, 107-108.	3.6	8
112	Contribution of chemical water properties to the differential responses of bacterioneuston and bacterioplankton to ultraviolet-B radiation. <i>FEMS Microbiology Ecology</i> , 2014, 87, 517-535.	2.7	8
113	Biosurfactant Production in Sub-Oxic Conditions Detected in Hydrocarbon-Degrading Isolates from Marine and Estuarine Sediments. <i>International Journal of Environmental Research and Public Health</i> , 2020, 17, 1746.	2.6	8
114	Environmental controls on estuarine nitrifying communities along a salinity gradient. <i>Aquatic Microbial Ecology</i> , 2017, 80, 167-180.	1.8	8
115	Bacterial biomass production in an estuarine system: high variability of leucine conversion factors and changes in bacterial community structure during incubation. <i>Aquatic Microbial Ecology</i> , 2011, 62, 299-310.	1.8	6
116	Independent and interactive effects of reduced seawater pH and oil contamination on subsurface sediment bacterial communities. <i>Environmental Science and Pollution Research</i> , 2018, 25, 32756-32766.	5.3	6
117	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
118	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
119	Petroleum contamination and bioaugmentation in bacterial rhizosphere communities from <i>Avicennia schaueriana</i> . <i>Brazilian Journal of Microbiology</i> , 2018, 49, 757-769.	2.0	6
120	SELECTIVE CULTURES FOR THE ISOLATION OF BIOSURFACTANT PRODUCING BACTERIA: COMPARISON OF DIFFERENT COMBINATIONS OF ENVIRONMENTAL INOCULA AND HYDROPHOBIC CARBON SOURCES. <i>Preparative Biochemistry and Biotechnology</i> , 2013, 43, 237-255.	1.9	5
121	Bacterial communities from corals cultured ex situ remain stable under different light regimes – Relevance for in toto aquaculture. <i>Aquaculture</i> , 2016, 450, 258-261.	3.5	5
122	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
123	Maintenance of <i>Rhodotorula rubra</i> isolated from liquid samples of gold mine effluents. <i>Folia Microbiologica</i> , 1995, 40, 487-489.	2.3	3
124	Bacterial communities associated with the decomposition of <i>Fucus vesiculosus</i> in transitional waters. <i>Estuarine, Coastal and Shelf Science</i> , 2012, 110, 116-124.	2.1	3
125	Growth conditions influence UVB sensitivity and oxidative damage in an estuarine bacterial isolate. <i>Photochemical and Photobiological Sciences</i> , 2013, 12, 974-986.	2.9	3
126	Baseline information on prokaryotic and microeukaryotic plankton communities inside and outside of Indonesian marine lakes. <i>Journal of Sea Research</i> , 2019, 148-149, 23-32.	1.6	3

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127	Comparison of bacterial communities associated with <i>Xestospongia testudinaria</i> , sediment and seawater in a Singaporean coral reef ecosystem. <i>Journal of the Marine Biological Association of the United Kingdom</i> , 2019, 99, 331-342.	0.8	3
128	Effects of the Inoculant Strain <i>Pseudomonas</i> sp. SPN31 nah + and of 2-Methylnaphthalene Contamination on the Rhizosphere and Endosphere Bacterial Communities of <i>Halimione portulacoides</i> . <i>Current Microbiology</i> , 2017, 74, 575-583.	2.2	2
129	Degradation of thiocyanate by immobilized cells of mixed and pure cultures. <i>Process Metallurgy</i> , 1999, 9, 819-824.	0.1	1
130	Influence of incubation conditions on bacterial production estimates in an estuarine system. <i>Aquatic Ecology</i> , 2014, 48, 327-336.	1.5	1
131	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
132	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
133	Biosorption of free and complexed cadmium ions by <i>Aspergillus niger</i> . <i>Process Metallurgy</i> , 1999, , 513-519.	0.1	0
134	Enhancement of gold-cyanide biosorption by L-cysteine. <i>Process Metallurgy</i> , 1999, 9, 493-502.	0.1	0
135	Effect of different culture conditions on the structural diversity of prokaryote communities in the sediment of earth ponds stocked with gilthead seabream <i>Sparus aurata</i> (Linnaeus, 1758). <i>Aquaculture Research</i> , 2015, 46, 1760-1769.	1.8	0
136	Involvement of oxidative stress in UV-induced impairment of bacterial activity and culturability. , 2010, , .		0
137	Draft Genome Sequence of <i>Vibrio mediterranei</i> Strain CyArs1. <i>Microbiology Resource Announcements</i> , 2022, , e0015522.	0.6	0