

# Gerhard J Herndl

## List of Publications by Year in descending order

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238  
papers

22,672  
citations

9756

73  
h-index

10127

140  
g-index

254  
all docs

254  
docs citations

254  
times ranked

16196  
citing authors

#	ARTICLE	IF	CITATIONS
1	Prokaryotic Life in the Deep Ocean's Water Column. <i>Annual Review of Marine Science</i> , 2023, 15, 461-483.	5.1	20
2	Phylogenetically and functionally diverse microorganisms reside under the Ross Ice Shelf. <i>Nature Communications</i> , 2022, 13, 117.	5.8	17
3	Microbial Consortia of Putative Degradors of Low-Density Polyethylene-Associated Compounds in the Ocean. <i>MSystems</i> , 2022, 7, e0141521.	1.7	7
4	Phylogeny and Metabolic Potential of the Candidate Phylum SAR324. <i>Biology</i> , 2022, 11, 599.	1.3	8
5	Extracellular Enzymatic Activities of Oceanic Pelagic Fungal Strains and the Influence of Temperature. <i>Journal of Fungi</i> (Basel, Switzerland), 2022, 8, 571.	1.5	4
6	Recognizing the complexity of soil organic carbon dynamics in vegetated coastal habitats. <i>Global Change Biology</i> , 2021, 27, 3-4.	4.2	1
7	What Is Refractory Organic Matter in the Ocean?. <i>Frontiers in Marine Science</i> , 2021, 8, .	1.2	31
8	The importance of jellyfish-microbe interactions for biogeochemical cycles in the ocean. <i>Limnology and Oceanography</i> , 2021, 66, 2011-2032.	1.6	20
9	Correcting a major error in assessing organic carbon pollution in natural waters. <i>Science Advances</i> , 2021, 7, .	4.7	37
10	Enzyme promiscuity in natural environments: alkaline phosphatase in the ocean. <i>ISME Journal</i> , 2021, 15, 3375-3383.	4.4	30
11	Selective DNA and Protein Isolation From Marine Macrophyte Surfaces. <i>Frontiers in Microbiology</i> , 2021, 12, 665999.	1.5	3
12	Potential and expression of carbohydrate utilization by marine fungi in the global ocean. <i>Microbiome</i> , 2021, 9, 106.	4.9	28
13	Reviews and syntheses: Heterotrophic fixation of inorganic carbon – significant but invisible flux in environmental carbon cycling. <i>Biogeosciences</i> , 2021, 18, 3689-3700.	1.3	37
14	Adapting an Ergosterol Extraction Method with Marine Yeasts for the Quantification of Oceanic Fungal Biomass. <i>Journal of Fungi</i> (Basel, Switzerland), 2021, 7, 690.	1.5	8
15	Microbes mediating the sulfur cycle in the Atlantic Ocean and their link to chemolithoautotrophy. <i>Environmental Microbiology</i> , 2021, 23, 7152-7167.	1.8	3
16	Seasonal Dynamics of Epiphytic Microbial Communities on Marine Macrophyte Surfaces. <i>Frontiers in Microbiology</i> , 2021, 12, 671342.	1.5	11
17	Seasonality combined with the orientation of surfaces influences the microbial community structure of biofilms in the deep Mediterranean Sea. <i>Deep-Sea Research Part II: Topical Studies in Oceanography</i> , 2020, 171, 104703.	0.6	7
18	Functional Seasonality of Free-Living and Particle-Associated Prokaryotic Communities in the Coastal Adriatic Sea. <i>Frontiers in Microbiology</i> , 2020, 11, 584222.	1.5	9

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19	Microbial Processing of Jellyfish Detritus in the Ocean. <i>Frontiers in Microbiology</i> , 2020, 11, 590995.	1.5	19
20	Nitrogen Isotope Fractionation During Archaeal Ammonia Oxidation: Coupled Estimates From Measurements of Residual Ammonium and Accumulated Nitrite. <i>Frontiers in Microbiology</i> , 2020, 11, 1710.	1.5	10
21	Mesozooplankton taurine production and prokaryotic uptake in the northern Adriatic Sea. <i>Limnology and Oceanography</i> , 2020, 65, 2730-2747.	1.6	4
22	Putative degraders of low-density polyethylene-derived compounds are ubiquitous members of plastic-associated bacterial communities in the marine environment. <i>Environmental Microbiology</i> , 2020, 22, 4779-4793.	1.8	21
23	Nitrifier adaptation to low energy flux controls inventory of reduced nitrogen in the dark ocean. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 4823-4830.	3.3	72
24	Hiding in Plain Sight: The Globally Distributed Bacterial Candidate Phylum PAUC34f. <i>Frontiers in Microbiology</i> , 2020, 11, 376.	1.5	5
25	Linking extracellular enzymes to phylogeny indicates a predominantly particle-associated lifestyle of deep-sea prokaryotes. <i>Science Advances</i> , 2020, 6, eaaz4354.	4.7	63
26	Relative Importance of Phosphodiesterase vs. Phosphomonoesterase (Alkaline Phosphatase) Activities for Dissolved Organic Phosphorus Hydrolysis in Epi- and Mesopelagic Waters. <i>Frontiers in Earth Science</i> , 2020, 8, .	0.8	12
27	Effects of the Invasion of <i>Caulerpa cylindracea</i> in a <i>Cymodocea nodosa</i> Meadow in the Northern Adriatic Sea. <i>Frontiers in Marine Science</i> , 2020, 7, .	1.2	5
28	Dynamics of environmental conditions during the decline of a <i>Cymodocea nodosa</i> meadow. <i>Biogeosciences</i> , 2020, 17, 3299-3315.	1.3	6
29	Highly variable mRNA half-life time within marine bacterial taxa and functional genes. <i>Environmental Microbiology</i> , 2019, 21, 3873-3884.	1.8	21
30	Seasonal dynamics of marine snow-associated and free-living demethylating bacterial communities in the coastal northern Adriatic Sea. <i>Environmental Microbiology Reports</i> , 2019, 11, 699-707.	1.0	19
31	Ammonia-oxidizing archaea release a suite of organic compounds potentially fueling prokaryotic heterotrophy in the ocean. <i>Environmental Microbiology</i> , 2019, 21, 4062-4075.	1.8	71
32	Towards Integrating Evolution, Metabolism, and Climate Change Studies of Marine Ecosystems. <i>Trends in Ecology and Evolution</i> , 2019, 34, 1022-1033.	4.2	28
33	Viral Communities in the Global Deep Ocean Conveyor Belt Assessed by Targeted Viromics. <i>Frontiers in Microbiology</i> , 2019, 10, 1801.	1.5	21
34	Niche Differentiation of Aerobic and Anaerobic Ammonia Oxidizers in a High Latitude Deep Oxygen Minimum Zone. <i>Frontiers in Microbiology</i> , 2019, 10, 2141.	1.5	44
35	Uneven host cell growth causes lysogenic virus induction in the Baltic Sea. <i>PLoS ONE</i> , 2019, 14, e0220716.	1.1	4
36	Taurine Is a Major Carbon and Energy Source for Marine Prokaryotes in the North Atlantic Ocean off the Iberian Peninsula. <i>Microbial Ecology</i> , 2019, 78, 299-312.	1.4	59

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37	Resolving the paradox: Continuous cell-free alkaline phosphatase activity despite high phosphate concentrations. <i>Marine Chemistry</i> , 2019, 214, 103671.	0.9	18
38	Proteomic Response of Three Marine Ammonia-Oxidizing Archaea to Hydrogen Peroxide and Their Metabolic Interactions with a Heterotrophic Alphaproteobacterium. <i>MSystems</i> , 2019, 4, .	1.7	57
39	Global Structuring of Phylogenetic and Functional Diversity of Pelagic Fungi by Depth and Temperature. <i>Frontiers in Marine Science</i> , 2019, 6, .	1.2	39
40	The composition of bacterial communities associated with plastic biofilms differs between different polymers and stages of biofilm succession. <i>PLoS ONE</i> , 2019, 14, e0217165.	1.1	190
41	Jellyfish-Associated Microbiome in the Marine Environment: Exploring Its Biotechnological Potential. <i>Marine Drugs</i> , 2019, 17, 94.	2.2	39
42	Ideas and perspectives: Is dark carbon fixation relevant for oceanic primary production estimates?. <i>Biogeosciences</i> , 2019, 16, 3793-3799.	1.3	36
43	Estimating Carbon Flux From Optically Recording Total Particle Volume at Depths Below the Primary Pycnocline. <i>Frontiers in Marine Science</i> , 2019, 6, .	1.2	2
44	Differential Response of <i>Cafeteria roenbergensis</i> to Different Bacterial and Archaeal Prey Characteristics. <i>Microbial Ecology</i> , 2019, 78, 1-5.	1.4	16
45	<i>Nitrosopumilus adriaticus</i> sp. nov. and <i>Nitrosopumilus piranensis</i> sp. nov., two ammonia-oxidizing archaea from the Adriatic Sea and members of the class Nitrososphaeria. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 69, 1892-1902.	0.8	64
46	Dissolved organic carbon leaching from plastics stimulates microbial activity in the ocean. <i>Nature Communications</i> , 2018, 9, 1430.	5.8	402
47	Mixing alters the lytic activity of viruses in the dark ocean. <i>Ecology</i> , 2018, 99, 700-713.	1.5	14
48	Organic matter processing by microbial communities throughout the Atlantic water column as revealed by metaproteomics. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E400-E408.	3.3	146
49	Metagenomic insights into zooplankton-associated bacterial communities. <i>Environmental Microbiology</i> , 2018, 20, 492-505.	1.8	57
50	High dark inorganic carbon fixation rates by specific microbial groups in the Atlantic off the Galician coast (NW Iberian margin). <i>Environmental Microbiology</i> , 2018, 20, 602-611.	1.8	22
51	Host Differentiation and Compartmentalization of Microbial Communities in the Azooxanthellate Cupcorals <i>Tubastrea coccinea</i> and <i>Rhizopsammia goesi</i> in the Caribbean. <i>Frontiers in Marine Science</i> , 2018, 5, .	1.2	25
52	Seasonal variations in extracellular enzymatic activity in marine snow-associated microbial communities and their impact on the surrounding water. <i>FEMS Microbiology Ecology</i> , 2018, 94, .	1.3	15
53	Dialysis Bag Incubation as a Nonradiolabeling Technique to Estimate Bacterioplankton Production In Situ. , 2018, , 553-556.		3
54	Microbiome variation in corals with distinct depth distribution ranges across a shallow-to-mesophotic gradient (15-85m). <i>Coral Reefs</i> , 2017, 36, 447-452.	0.9	34

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55	SAR202 Genomes from the Dark Ocean Predict Pathways for the Oxidation of Recalcitrant Dissolved Organic Matter. <i>MBio</i> , 2017, 8, .	1.8	168
56	Chemotaxonomic characterisation of the thaumarchaeal lipidome. <i>Environmental Microbiology</i> , 2017, 19, 2681-2700.	1.8	117
57	Crustacean zooplankton release copious amounts of dissolved organic matter as taurine in the ocean. <i>Limnology and Oceanography</i> , 2017, 62, 2745-2758.	1.6	44
58	Major role of nitrite-oxidizing bacteria in dark ocean carbon fixation. <i>Science</i> , 2017, 358, 1046-1051.	6.0	229
59	Extracting DNA from ocean microplastics: a method comparison study. <i>Analytical Methods</i> , 2017, 9, 1521-1526.	1.3	46
60	Eukaryotic microbes, principally fungi and labyrinthulomycetes, dominate biomass on bathypelagic marine snow. <i>ISME Journal</i> , 2017, 11, 362-373.	4.4	169
61	High viral abundance as a consequence of low viral decay in the Baltic Sea redoxcline. <i>PLoS ONE</i> , 2017, 12, e0178467.	1.1	12
62	Depth Dependent Relationships between Temperature and Ocean Heterotrophic Prokaryotic Production. <i>Frontiers in Marine Science</i> , 2016, 3, .	1.2	37
63	Geographic Distribution of Archaeal Ammonia Oxidizing Ecotypes in the Atlantic Ocean. <i>Frontiers in Microbiology</i> , 2016, 7, 77.	1.5	84
64	Prokaryotic Responses to Ammonium and Organic Carbon Reveal Alternative CO <sub>2</sub> Fixation Pathways and Importance of Alkaline Phosphatase in the Mesopelagic North Atlantic. <i>Frontiers in Microbiology</i> , 2016, 7, 1670.	1.5	47
65	Connectivity between surface and deep waters determines prokaryotic diversity in the North Atlantic Deep Water. <i>Environmental Microbiology</i> , 2016, 18, 2052-2063.	1.8	58
66	Dimethylsulfoniopropionate in corals and its interrelations with bacterial assemblages in coral surface mucus. <i>Environmental Chemistry</i> , 2016, 13, 252.	0.7	28
67	Springtime dynamics, productivity and activity of prokaryotes in two Arctic fjords. <i>Polar Biology</i> , 2016, 39, 1749-1763.	0.5	21
68	Large-scale distribution of microbial and viral populations in the South Atlantic Ocean. <i>Environmental Microbiology Reports</i> , 2016, 8, 305-315.	1.0	38
69	Erythromycin and GC7 fail as domain-specific inhibitors for bacterial and archaeal activity in the open ocean. <i>Aquatic Microbial Ecology</i> , 2016, 77, 99-110.	0.9	2
70	Dragon kings of the deep sea: marine particles deviate markedly from the common number-size spectrum. <i>Scientific Reports</i> , 2016, 6, 22633.	1.6	58
71	The microbiome of coral surface mucus has a key role in mediating holobiont health and survival upon disturbance. <i>ISME Journal</i> , 2016, 10, 2280-2292.	4.4	280
72	Physiological and genomic characterization of two novel marine thaumarchaeal strains indicates niche differentiation. <i>ISME Journal</i> , 2016, 10, 1051-1063.	4.4	160

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73	Archaeal and Bacterial Communities Associated with the Surface Mucus of Caribbean Corals Differ in Their Degree of Host Specificity and Community Turnover Over Reefs. <i>PLoS ONE</i> , 2016, 11, e0144702.	1.1	30
74	Conservation of dissolved organic matter molecular composition during mixing of the deep water masses of the northeast Atlantic Ocean. <i>Marine Chemistry</i> , 2015, 177, 288-297.	0.9	51
75	Response to Comment on "Dilution limits dissolved organic carbon utilization in the deep ocean". <i>Science</i> , 2015, 350, 1483-1483.	6.0	11
76	Dilution limits dissolved organic carbon utilization in the deep ocean. <i>Science</i> , 2015, 348, 331-333.	6.0	230
77	Macroecological patterns of archaeal ammonia oxidizers in the Atlantic Ocean. <i>Molecular Ecology</i> , 2015, 24, 4931-4942.	2.0	34
78	Production and degradation of fluorescent dissolved organic matter in surface waters of the eastern north Atlantic ocean. <i>Deep-Sea Research Part I: Oceanographic Research Papers</i> , 2015, 96, 28-37.	0.6	43
79	Potential impacts of black carbon on the marine microbial community. <i>Aquatic Microbial Ecology</i> , 2015, 75, 27-42.	0.9	25
80	Resolving the abundance and air-sea fluxes of airborne microorganisms in the North Atlantic Ocean. <i>Frontiers in Microbiology</i> , 2014, 5, 557.	1.5	76
81	Fracture zones in the Mid Atlantic Ridge lead to alterations in prokaryotic and viral parameters in deep-water masses. <i>Frontiers in Microbiology</i> , 2014, 5, 264.	1.5	17
82	Drivers shaping the diversity and biogeography of total and active bacterial communities in the South China Sea. <i>Molecular Ecology</i> , 2014, 23, 2260-2274.	2.0	194
83	Comparison of Deep-Water Viromes from the Atlantic Ocean and the Mediterranean Sea. <i>PLoS ONE</i> , 2014, 9, e100600.	1.1	42
84	Linkage between copepods and bacteria in the North Atlantic Ocean. <i>Aquatic Microbial Ecology</i> , 2014, 72, 215-225.	0.9	41
85	Seasonal variation in marine-snow-associated and ambient-water prokaryotic communities in the northern Adriatic Sea. <i>Aquatic Microbial Ecology</i> , 2014, 73, 211-224.	0.9	41
86	Archaeal <i>amoA</i> gene diversity points to distinct biogeography of ammonia-oxidizing <i>Crenarchaeota</i> in the ocean. <i>Environmental Microbiology</i> , 2013, 15, 1647-1658.	1.8	169
87	Bacterial Versus Archaeal Origin of Extracellular Enzymatic Activity in the Northeast Atlantic Deep Waters. <i>Microbial Ecology</i> , 2013, 65, 277-288.	1.4	45
88	Microbial control of the dark end of the biological pump. <i>Nature Geoscience</i> , 2013, 6, 718-724.	5.4	276
89	Diversity and distribution of microbial eukaryotes in the deep tropical and subtropical North Atlantic Ocean. <i>Deep-Sea Research Part I: Oceanographic Research Papers</i> , 2013, 78, 58-69.	0.6	22
90	Temporal dynamics in the free-living bacterial community composition in the coastal North Sea. <i>FEMS Microbiology Ecology</i> , 2013, 83, 413-424.	1.3	31

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91	Abundance and distribution of archaeal acetyl-CoA/propionyl-CoA carboxylase genes indicative for putatively chemoautotrophic Archaea in the tropical Atlantic's interior. <i>FEMS Microbiology Ecology</i> , 2013, 84, 461-473.	1.3	21
92	Comparison between MICROCARD-FISH and 16 S rRNA gene clone libraries to assess the active versus total bacterial community in the coastal Arctic. <i>Environmental Microbiology Reports</i> , 2013, 5, 272-281.	1.0	21
93	Spatial patterns of bacterial and archaeal communities along the Romanche Fracture Zone (tropical) Tj ETQq1 1 0.784314 rgBT /Over 16	1.3	16
94	Thick-shelled, grazer-protected diatoms decouple ocean carbon and silicon cycles in the iron-limited Antarctic Circumpolar Current. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 20633-20638.	3.3	216
95	Prevalence of strong vertical CO <sub>2</sub> and O <sub>2</sub> variability in the top meters of the ocean. <i>Global Biogeochemical Cycles</i> , 2013, 27, 941-949.	1.9	15
96	Impact of water mass mixing on the biogeochemistry and microbiology of the Northeast Atlantic Deep Water. <i>Global Biogeochemical Cycles</i> , 2013, 27, 1151-1162.	1.9	18
97	Development and deployment of a point-source digital inline holographic microscope for the study of plankton and particles to a depth of 6000 m. <i>Limnology and Oceanography: Methods</i> , 2013, 11, 28-40.	1.0	71
98	Major Effect of Hydrogen Peroxide on Bacterioplankton Metabolism in the Northeast Atlantic. <i>PLoS ONE</i> , 2013, 8, e61051.	1.1	23
99	Microbial Functioning and Community Structure Variability in the Mesopelagic and Epipelagic Waters of the Subtropical Northeast Atlantic Ocean. <i>Applied and Environmental Microbiology</i> , 2012, 78, 3309-3316.	1.4	21
100	Links between viruses and prokaryotes throughout the water column along a North Atlantic latitudinal transect. <i>ISME Journal</i> , 2012, 6, 1566-1577.	4.4	90
101	Direct observations of diel biological CO <sub>2</sub> and CH <sub>4</sub> fixation on the Scotian Shelf, northwestern Atlantic Ocean. <i>Biogeosciences</i> , 2012, 9, 2301-2309.	1.3	10
102	Deep carbon export from a Southern Ocean iron-fertilized diatom bloom. <i>Nature</i> , 2012, 487, 313-319.	13.7	367
103	Sunlight Effects on the Osmotrophic Uptake of DMSP-Sulfur and Leucine by Polar Phytoplankton. <i>PLoS ONE</i> , 2012, 7, e45545.	1.1	21
104	Differentiating leucine incorporation of Archaea and Bacteria throughout the water column of the eastern Atlantic using metabolic inhibitors. <i>Aquatic Microbial Ecology</i> , 2012, 66, 247-256.	0.9	9
105	Potential for Chemolithoautotrophy Among Ubiquitous Bacteria Lineages in the Dark Ocean. <i>Science</i> , 2011, 333, 1296-1300.	6.0	510
106	Contribution of <i>Crenarchaeota</i> and <i>Bacteria</i> to autotrophy in the North Atlantic interior. <i>Environmental Microbiology</i> , 2011, 13, 1524-1533.	1.8	42
107	Changes in viral and bacterial communities during the ice-melting season in the coastal Arctic (Kongsfjorden, Ny-Ålesund). <i>Environmental Microbiology</i> , 2011, 13, 1827-1841.	1.8	37
108	Water mass-specificity of bacterial communities in the North Atlantic revealed by massively parallel sequencing. <i>Molecular Ecology</i> , 2011, 20, 258-274.	2.0	243

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109	DIEL IN SITU PICOPHYTOPLANKTON CELL DEATH CYCLES COUPLED WITH CELL DIVISION <sup>1</sup> . Journal of Phycology, 2011, 47, 1247-1257.	1.0	16
110	The microbial carbon pump and the oceanic recalcitrant dissolved organic matter pool. Nature Reviews Microbiology, 2011, 9, 555-555.	13.6	73
111	An overview of the structure and function of microbial biofilms, with special emphasis on heterotrophic aquatic microbial communities. African Journal of Aquatic Science, 2011, 36, 1-10.	0.5	6
112	Abundance of eukaryotic microbes in the deep subtropical North Atlantic. Aquatic Microbial Ecology, 2011, 65, 103-115.	0.9	15
113	Spatial patterns of bacterial abundance, activity and community composition in relation to water masses in the eastern Mediterranean Sea. Aquatic Microbial Ecology, 2010, 59, 185-195.	0.9	36
114	Mesoscale eddies: hotspots of prokaryotic activity and differential community structure in the ocean. ISME Journal, 2010, 4, 975-988.	4.4	86
115	Links between viral and prokaryotic communities throughout the water column in the (sub)tropical Atlantic Ocean. ISME Journal, 2010, 4, 1431-1442.	4.4	47
116	Relevance of a crenarchaeotal subcluster related to <i>Candidatus</i> Nitrosopumilus maritimus to ammonia oxidation in the suboxic zone of the central Baltic Sea. ISME Journal, 2010, 4, 1496-1508.	4.4	110
117	Microbial production of recalcitrant dissolved organic matter: long-term carbon storage in the global ocean. Nature Reviews Microbiology, 2010, 8, 593-599.	13.6	1,278
118	Role of macroscopic particles in deep-sea oxygen consumption. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 8287-8291.	3.3	79
119	Significance of non-sinking particulate organic carbon and dark CO <sub>2</sub> fixation to heterotrophic carbon demand in the mesopelagic northeast Atlantic. Geophysical Research Letters, 2010, 37, .	1.5	64
120	Biogeochemical relationships between ultrafiltered dissolved organic matter and picoplankton activity in the Eastern Mediterranean Sea. Deep-Sea Research Part II: Topical Studies in Oceanography, 2010, 57, 1460-1477.	0.6	48
121	Emerging concepts on microbial processes in the bathypelagic ocean – ecology, biogeochemistry, and genomics. Deep-Sea Research Part II: Topical Studies in Oceanography, 2010, 57, 1519-1536.	0.6	153
122	Major contribution of autotrophy to microbial carbon cycling in the deep North Atlantic's interior. Deep-Sea Research Part II: Topical Studies in Oceanography, 2010, 57, 1572-1580.	0.6	152
123	High dissolved extracellular enzymatic activity in the deep central Atlantic Ocean. Aquatic Microbial Ecology, 2010, 58, 287-302.	0.9	90
124	Seasonal dynamics of dissolved organic matter and microbial activity in the coastal North Sea. Aquatic Microbial Ecology, 2010, 60, 85-95.	0.9	33
125	Prokaryotic carbon utilization in the dark ocean: growth efficiency, leucine-to-carbon conversion factors, and their relation. Aquatic Microbial Ecology, 2010, 60, 227-232.	0.9	39
126	Synechococcus and Prochlorococcus cell death induced by UV radiation and the penetration of lethal UVR in the Mediterranean Sea. Marine Ecology - Progress Series, 2010, 399, 27-37.	0.9	57



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127	Viral and Flagellate Control of Prokaryotic Production and Community Structure in Offshore Mediterranean Waters. <i>Applied and Environmental Microbiology</i> , 2009, 75, 4801-4812.	1.4	60
128	Mesoscale variability modulates seasonal changes in the trophic structure of nano- and picoplankton communities across the NW Africa-Canary Islands transition zone. <i>Progress in Oceanography</i> , 2009, 83, 180-188.	1.5	35
129	Spatial distribution of <i>Bacteria</i> and <i>Archaea</i> and <i>amoA</i> gene copy numbers throughout the water column of the Eastern Mediterranean Sea. <i>ISME Journal</i> , 2009, 3, 147-158.	4.4	134
130	Heterotrophic prokaryotic production in ultraoligotrophic alpine karst aquifers and ecological implications. <i>FEMS Microbiology Ecology</i> , 2009, 68, 287-299.	1.3	55
131	Prokaryotic extracellular enzymatic activity in relation to biomass production and respiration in the meso- and bathypelagic waters of the (sub)tropical Atlantic. <i>Environmental Microbiology</i> , 2009, 11, 1998-2014.	1.8	117
132	Deep-sea bacterial communities in sediments and guts of deposit-feeding holothurians in Portuguese canyons (NE Atlantic). <i>Deep-Sea Research Part I: Oceanographic Research Papers</i> , 2009, 56, 1834-1843.	0.6	47
133	Microbial oceanography of the dark ocean's pelagic realm. <i>Limnology and Oceanography</i> , 2009, 54, 1501-1529.	1.6	437
134	Evidence of prokaryotic metabolism on suspended particulate organic matter in the dark waters of the subtropical North Atlantic. <i>Limnology and Oceanography</i> , 2009, 54, 182-193.	1.6	120
135	Diversity of Archaea and detection of crenarchaeotal <i>amoA</i> genes in the rivers Rhine and $\ddot{A}$ t. <i>Aquatic Microbial Ecology</i> , 2009, 55, 189-201.	0.9	42
136	Role of mesoscale cyclonic eddies in the distribution and activity of Archaea and Bacteria in the South China Sea. <i>Aquatic Microbial Ecology</i> , 2009, 56, 65-79.	0.9	39
137	Latitudinal trends of <i>Crenarchaeota</i> and <i>Bacteria</i> in the meso- and bathypelagic water masses of the Eastern North Atlantic. <i>Environmental Microbiology</i> , 2008, 10, 110-124.	1.8	104
138	Relationship of Geographic Distance, Depth, Temperature, and Viruses with Prokaryotic Communities in the Eastern Tropical Atlantic Ocean. <i>Microbial Ecology</i> , 2008, 56, 383-389.	1.4	27
139	Dynamics and diversity of newly produced virioplankton in the North Sea. <i>ISME Journal</i> , 2008, 2, 924-936.	4.4	35
140	Major gradients in putatively nitrifying and non-nitrifying Archaea in the deep North Atlantic. <i>Nature</i> , 2008, 456, 788-791.	13.7	259
141	Abundance and activity of <i>Chloroflexi</i> -type SAR202 bacterioplankton in the meso- and bathypelagic waters of the (sub)tropical Atlantic. <i>Environmental Microbiology</i> , 2008, 10, 1903-1911.	1.8	99
142	<i>Epsilonproteobacteria</i> Represent the Major Portion of Chemoautotrophic Bacteria in Sulfidic Waters of Pelagic Redoxclines of the Baltic and Black Seas. <i>Applied and Environmental Microbiology</i> , 2008, 74, 7546-7551.	1.4	131
143	Towards a better understanding of microbial carbon flux in the sea*. <i>Aquatic Microbial Ecology</i> , 2008, 53, 21-38.	0.9	81
144	Dissolved organic matter and bacterial production and respiration in the sea's surface microlayer of the open Atlantic and the western Mediterranean Sea. <i>Limnology and Oceanography</i> , 2008, 53, 122-136.	1.6	110

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145	Regulation of aquatic microbial processes: the "microbial loop" of the sunlit surface waters and the dark ocean dissected. <i>Aquatic Microbial Ecology</i> , 2008, 53, 59-68.	0.9	44
146	Contribution of Crenarchaeota and Euryarchaeota to the prokaryotic plankton in the coastal northwestern Black Sea. <i>Journal of Plankton Research</i> , 2007, 29, 699-706.	0.8	44
147	Contribution of Crenarchaeota and Euryarchaeota to the prokaryotic plankton in the coastal northwestern Black Sea. <i>Journal of Plankton Research</i> , 2007, 30, 93-93.	0.8	0
148	Viral Abundance, Decay, and Diversity in the Meso- and Bathypelagic Waters of the North Atlantic. <i>Applied and Environmental Microbiology</i> , 2007, 73, 4429-4438.	1.4	105
149	Microbes and the Dissipation of Energy and Respiration: From Cells to Ecosystems. <i>Oceanography</i> , 2007, 20, 89-100.	0.5	125
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