

Carlos Pedraza-Ali

List of Publications by Year in descending order

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178
papers

18,168
citations

11608

70
h-index

14702

127
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189
all docs

189
docs citations

189
times ranked

13058
citing authors

#	ARTICLE	IF	CITATIONS
1	Deep ocean metagenomes provide insight into the metabolic architecture of bathypelagic microbial communities. <i>Communications Biology</i> , 2021, 4, 604.	2.0	107
2	Time travel in microorganisms. <i>Systematic and Applied Microbiology</i> , 2021, 44, 126227.	1.2	11
3	Proteorhodopsin Phototrophy in Antarctic Coastal Waters. <i>MSphere</i> , 2021, 6, e0052521.	1.3	2
4	Compendium of 530 metagenome-assembled bacterial and archaeal genomes from the polar Arctic Ocean. <i>Nature Microbiology</i> , 2021, 6, 1561-1574.	5.9	57
5	Diversity and distribution of marine heterotrophic bacteria from a large culture collection. <i>BMC Microbiology</i> , 2020, 20, 207.	1.3	27
6	Roadmap for naming uncultivated Archaea and Bacteria. <i>Nature Microbiology</i> , 2020, 5, 987-994.	5.9	115
7	<i>Thalassocella blandensis</i> gen. nov., sp. nov., a novel member of the family Cellvibrionaceae. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 1231-1239.	0.8	19
8	<i>Mesonia oceanica</i> sp. nov., isolated from oceans during the Tara oceans expedition, with a preference for mesopelagic waters. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 4329-4338.	0.8	11
9	Microdiversity ensures the maintenance of functional microbial communities under changing environmental conditions. <i>ISME Journal</i> , 2019, 13, 2969-2983.	4.4	121
10	The influence of temperature and pH on bacterial community composition of microbial mats in hot springs from Costa Rica. <i>MicrobiologyOpen</i> , 2019, 8, e893.	1.2	42
11	Delineation of ecologically distinct units of marine Bacteroidetes in the Northwestern Mediterranean Sea. <i>Molecular Ecology</i> , 2019, 28, 2846-2859.	2.0	31
12	Functional annotation of orthologs in metagenomes: a case study of genes for the transformation of oceanic dimethylsulfoniopropionate. <i>ISME Journal</i> , 2019, 13, 1183-1197.	4.4	24
13	Active Crossfire Between Cyanobacteria and Cyanophages in Phototrophic Mat Communities Within Hot Springs. <i>Frontiers in Microbiology</i> , 2018, 9, 2039.	1.5	29
14	Draft genome sequences of <i>Cylindrospermopsis raciborskii</i> strains CS-508 and MVCC14, isolated from freshwater bloom events in Australia and Uruguay. <i>Standards in Genomic Sciences</i> , 2018, 13, 26.	1.5	4
15	Diurnal Changes in Active Carbon and Nitrogen Pathways Along the Temperature Gradient in Porcelana Hot Spring Microbial Mat. <i>Frontiers in Microbiology</i> , 2018, 9, 2353.	1.5	36
16	The interactive microbial ocean. <i>Nature Microbiology</i> , 2017, 2, 16255.	5.9	15
17	Exploring Microdiversity in Novel <i>Kordia</i> sp. (Bacteroidetes) with Proteorhodopsin from the Tropical Indian Ocean via Single Amplified Genomes. <i>Frontiers in Microbiology</i> , 2017, 8, 1317.	1.5	7
18	Composition and Interactions among Bacterial, Microeukaryotic, and T4-like Viral Assemblages in Lakes from Both Polar Zones. <i>Frontiers in Microbiology</i> , 2016, 7, 337.	1.5	12

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19	Quantifying the Relative Importance of Phylogeny and Environmental Preferences As Drivers of Gene Content in Prokaryotic Microorganisms. <i>Frontiers in Microbiology</i> , 2016, 7, 433.	1.5	19
20	Age-Related Differences in the Gastrointestinal Microbiota of Chinstrap Penguins (<i>Pygoscelis</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 702	1.1	79
21	Global genetic capacity for mixotrophy in marine picocyanobacteria. <i>ISME Journal</i> , 2016, 10, 2946-2957.	4.4	82
22	Marine Bacterial and Archaeal Ion-Pumping Rhodopsins: Genetic Diversity, Physiology, and Ecology. <i>Microbiology and Molecular Biology Reviews</i> , 2016, 80, 929-954.	2.9	173
23	The vast unknown microbial biosphere. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 6585-6587.	3.3	29
24	Probing the Rare Biosphere of the North-West Mediterranean Sea: An Experiment with High Sequencing Effort. <i>PLoS ONE</i> , 2016, 11, e0159195.	1.1	17
25	Seasonal patterns in phytoplankton photosynthetic parameters and primary production at a coastal NW Mediterranean site. <i>Scientia Marina</i> , 2016, 80, 63-77.	0.3	57
26	Winter diversity and expression of proteorhodopsin genes in a polar ocean. <i>ISME Journal</i> , 2015, 9, 1835-1845.	4.4	22
27	Diversity of planktonic microorganisms in the Arctic Ocean. <i>Progress in Oceanography</i> , 2015, 139, 233-243.	1.5	52
28	Winter bloom of a rare betaproteobacterium in the Arctic Ocean. <i>Frontiers in Microbiology</i> , 2014, 5, 425.	1.5	43
29	The phylogenetic and ecological context of cultured and whole genome-sequenced planktonic bacteria from the coastal NW Mediterranean Sea. <i>Systematic and Applied Microbiology</i> , 2014, 37, 216-228.	1.2	22
30	Stimulation of growth by proteorhodopsin phototrophy involves regulation of central metabolic pathways in marine planktonic bacteria. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, E3650-8.	3.3	87
31	Seasonal changes in substrate utilization patterns by bacterioplankton in the Amundsen Gulf (western Arctic). <i>Polar Biology</i> , 2014, 37, 1321-1329.	0.5	7
32	Polar marine biology science in Portugal and Spain: Recent advances and future perspectives. <i>Journal of Sea Research</i> , 2013, 83, 9-29.	0.6	15
33	Bacterial composition of microbial mats in hot springs in Northern Patagonia: variations with seasons and temperature. <i>Extremophiles</i> , 2013, 17, 123-136.	0.9	75
34	Ecology of marine Bacteroidetes: a comparative genomics approach. <i>ISME Journal</i> , 2013, 7, 1026-1037.	4.4	614
35	Rare Biosphere. , 2013, , 345-352.		2
36	Distribution of Microbial Arsenic Reduction, Oxidation and Extrusion Genes along a Wide Range of Environmental Arsenic Concentrations. <i>PLoS ONE</i> , 2013, 8, e78890.	1.1	97

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37	Taxonomic composition of the particle-associated and free-living bacterial assemblages in the Northwest Mediterranean Sea analyzed by pyrosequencing of the 16S rRNA. <i>MicrobiologyOpen</i> , 2013, 2, 541-552.	1.2	151
38	Pole-to-pole biogeography of surface and deep marine bacterial communities. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 17633-17638.	3.3	283
39	Role for urea in nitrification by polar marine Archaea. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 17989-17994.	3.3	253
40	Patterns and architecture of genomic islands in marine bacteria. <i>BMC Genomics</i> , 2012, 13, 347.	1.2	84
41	The Rare Bacterial Biosphere. <i>Annual Review of Marine Science</i> , 2012, 4, 449-466.	5.1	580
42	Enrichment of arsenic transforming and resistant heterotrophic bacteria from sediments of two salt lakes in Northern Chile. <i>Extremophiles</i> , 2012, 16, 523-538.	0.9	49
43	High cyanobacterial <i>nifH</i> gene diversity in Arctic seawater and sea ice brine. <i>Environmental Microbiology Reports</i> , 2012, 4, 360-366.	1.0	67
44	Picoplankton seasonal variation and community structure in the northeast Adriatic coastal zone. <i>FEMS Microbiology Ecology</i> , 2012, 82, 678-691.	1.3	25
45	A Holistic Approach to Marine Eco-Systems Biology. <i>PLoS Biology</i> , 2011, 9, e1001177.	2.6	353
46	Transcriptome Fingerprinting Analysis: An Approach to Explore Gene Expression Patterns in Marine Microbial Communities. <i>PLoS ONE</i> , 2011, 6, e22950.	1.1	3
47	Biogenic carbon flows through the planktonic food web of the Amundsen Gulf (Arctic Ocean): A synthesis of field measurements and inverse modeling analyses. <i>Progress in Oceanography</i> , 2011, 91, 410-436.	1.5	138
48	Genomics of the Proteorhodopsin-Containing Marine Flavobacterium <i>Dokdonia</i> sp. Strain MED134. <i>Applied and Environmental Microbiology</i> , 2011, 77, 8676-8686.	1.4	56
49	High bicarbonate assimilation in the dark by Arctic bacteria. <i>ISME Journal</i> , 2010, 4, 1581-1590.	4.4	131
50	Spatial patterns of bacterial richness and evenness in the NW Mediterranean Sea explored by pyrosequencing of the 16S rRNA. <i>Aquatic Microbial Ecology</i> , 2010, 61, 221-233.	0.9	100
51	Evaluation of DNA extraction methods from complex phototrophic biofilms. <i>Biofouling</i> , 2010, 26, 349-357.	0.8	24
52	<i>Bermanella marisrubri</i> gen. nov., sp. nov., a genome-sequenced gammaproteobacterium from the Red Sea. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2009, 59, 373-377.	0.8	24
53	Differential response of grazing and bacterial heterotrophic production to experimental warming in Antarctic waters. <i>Aquatic Microbial Ecology</i> , 2009, 54, 101-112.	0.9	31
54	Influence of primer mismatch and microdiversity on DGGE results: a case study with SAR11. <i>Aquatic Microbial Ecology</i> , 2009, 54, 211-216.	0.9	8

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55	Novelty and spatio-temporal heterogeneity in the bacterial diversity of hypersaline Lake Tebenquiche (Salar de Atacama). <i>Extremophiles</i> , 2008, 12, 491-504.	0.9	118
56	Metabolic diversity of heterotrophic bacterioplankton over winter and spring in the coastal Arctic Ocean. <i>Environmental Microbiology</i> , 2008, 10, 942-949.	1.8	68
57	Winter-to-summer changes in the composition and single-cell activity of near-surface Arctic prokaryotes. <i>Environmental Microbiology</i> , 2008, 10, 2444-2454.	1.8	145
58	A Lagrangian biogeochemical study of an eddy in the Northeast Atlantic. <i>Progress in Oceanography</i> , 2008, 76, 366-398.	1.5	19
59	Number and phylogenetic affiliation of bacteria assimilating dimethylsulfoniopropionate and leucine in the ice-covered coastal Arctic Ocean. <i>Journal of Marine Systems</i> , 2008, 74, 957-963.	0.9	18
60	Unveiling new microbial eukaryotes in the surface ocean. <i>Current Opinion in Microbiology</i> , 2008, 11, 213-218.	2.3	162
61	Genome analysis of the proteorhodopsin-containing marine bacterium <i>Polaribacter</i> sp. MED152 (Flavobacteria). <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 8724-8729.	3.3	231
62	Seasonal changes in planktonic bacterivory rates under the ice-covered coastal Arctic Ocean. <i>Limnology and Oceanography</i> , 2008, 53, 2427-2438.	1.6	58
63	Carbon dioxide fixation in the dark by photosynthetic bacteria in sulfide-rich stratified lakes with oxic-anoxic interfaces. <i>Limnology and Oceanography</i> , 2008, 53, 1193-1203.	1.6	57
64	Comparison of Different Denaturing Gradient Gel Electrophoresis Primer Sets for the Study of Marine Bacterioplankton Communities. <i>Applied and Environmental Microbiology</i> , 2007, 73, 5962-5967.	1.4	102
65	<i>Reinekea blandensis</i> sp. nov., a marine, genome-sequenced gammaproteobacterium. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2007, 57, 2370-2375.	0.8	21
66	<i>Neptuniibacter caesariensis</i> gen. nov., sp. nov., a novel marine genome-sequenced gammaproteobacterium. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2007, 57, 1000-1006.	0.8	58
67	Microbial Precipitation of Arsenic Sulfides in Andean Salt Flats. <i>Geomicrobiology Journal</i> , 2007, 24, 111-123.	1.0	63
68	Predictions for the Future of Microbial Oceanography. <i>Oceanography</i> , 2007, 20, 166-171.	0.5	3
69	Light stimulates growth of proteorhodopsin-containing marine Flavobacteria. <i>Nature</i> , 2007, 445, 210-213.	13.7	349
70	DISTRIBUTION, PHYLOGENY, AND GROWTH OF COLD-ADAPTED PICOPRASINOPHYTES IN ARCTIC SEAS. <i>Journal of Phycology</i> , 2007, 43, 78-89.	1.0	296
71	Seasonality in bacterial diversity in north-west Mediterranean coastal waters: assessment through clone libraries, fingerprinting and FISH. <i>FEMS Microbiology Ecology</i> , 2007, 60, 98-112.	1.3	195
72	Dipping into the Rare Biosphere. <i>Science</i> , 2007, 315, 192-193.	6.0	129

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73	Marine microbial diversity: can it be determined?. Trends in Microbiology, 2006, 14, 257-263.	3.5	612
74	Seasonal changes in bacterioplankton nutrient limitation and their effects on bacterial community composition in the NW Mediterranean Sea. Aquatic Microbial Ecology, 2006, 44, 241-252.	0.9	163
75	Response of Southern Ocean phytoplankton and bacterioplankton production to short-term experimental warming. Limnology and Oceanography, 2006, 51, 1791-1800.	1.6	56
76	Distribution and abundance of uncultured heterotrophic flagellates in the world oceans. Environmental Microbiology, 2006, 8, 1515-1522.	1.8	219
77	Spatial and temporal variation in marine bacterioplankton diversity as shown by RFLP fingerprinting of PCR amplified 16S rDNA. FEMS Microbiology Ecology, 2006, 24, 27-40.	1.3	123
78	Growth of uncultured heterotrophic flagellates in unamended seawater incubations. Aquatic Microbial Ecology, 2006, 45, 171-180.	0.9	50
79	Diversity and Distribution of Marine Microbial Eukaryotes in the Arctic Ocean and Adjacent Seas. Applied and Environmental Microbiology, 2006, 72, 3085-3095.	1.4	258
80	Leeuwenhoekiella blandensis sp. nov., a genome-sequenced marine member of the family Flavobacteriaceae. International Journal of Systematic and Evolutionary Microbiology, 2006, 56, 1489-1493.	0.8	57
81	Genomics and marine microbial ecology. International Microbiology, 2006, 9, 191-7.	1.1	24
82	Phylogenetic and functional diversity of bacterioplankton during Alexandrium spp. blooms. FEMS Microbiology Ecology, 2005, 54, 257-267.	1.3	43
83	Partitioning of CO2 Incorporation Among Planktonic Microbial Guilds and Estimation of In Situ Specific Growth Rates. Microbial Ecology, 2005, 50, 230-241.	1.4	32
84	Dimethylsulfoniopropionate Turnover Is Linked to the Composition and Dynamics of the Bacterioplankton Assemblage during a Microcosm Phytoplankton Bloom. Applied and Environmental Microbiology, 2005, 71, 7650-7660.	1.4	69
85	Late summer community composition and abundance of photosynthetic picoeukaryotes in Norwegian and Barents Seas. Limnology and Oceanography, 2005, 50, 1677-1686.	1.6	177
86	Diversity of Microbial Communities: The Case of Solar Salterns. , 2005, , 71-90.		7
87	Use of Microautoradiography Combined with Fluorescence In Situ Hybridization To Determine Dimethylsulfoniopropionate Incorporation by Marine Bacterioplankton Taxa. Applied and Environmental Microbiology, 2004, 70, 4648-4657.	1.4	86
88	Phylogenetic and Ecological Analysis of Novel Marine Stramenopiles. Applied and Environmental Microbiology, 2004, 70, 3528-3534.	1.4	321
89	Distribution of eukaryotic picoplankton assemblages across hydrographic fronts in the Southern Ocean, studied by denaturing gradient gel electrophoresis. Limnology and Oceanography, 2004, 49, 1022-1034.	1.6	51
90	Diversity of Picoplanktonic Prasinophytes Assessed by Direct Nuclear SSU rDNA Sequencing of Environmental Samples and Novel Isolates Retrieved from Oceanic and Coastal Marine Ecosystems. Protist, 2004, 155, 193-214.	0.6	235

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91	Distribution of prokaryotic genetic diversity in athalassohaline lakes of the Atacama Desert, Northern Chile. <i>FEMS Microbiology Ecology</i> , 2004, 48, 57-69.	1.3	163
92	High-diversity biofilm for the oxidation of sulfide-containing effluents. <i>Applied Microbiology and Biotechnology</i> , 2004, 64, 726-734.	1.7	45
93	Diversity of planktonic photoautotrophic microorganisms along a salinity gradient as depicted by microscopy, flow cytometry, pigment analysis and DNA-based methods. <i>FEMS Microbiology Ecology</i> , 2004, 49, 281-293.	1.3	98
94	Picoeukaryotic diversity in an oligotrophic coastal site studied by molecular and culturing approaches. <i>FEMS Microbiology Ecology</i> , 2004, 50, 231-243.	1.3	204
95	Control of heterotrophic prokaryotic abundance and growth rate in hypersaline planktonic environments. <i>Aquatic Microbial Ecology</i> , 2004, 34, 193-206.	0.9	66
96	Trophic Ecology of Solar Salterns. , 2004, , 33-48.		20
97	Seasonal changes in the taxonomic composition of bacterioplankton in a coastal oligotrophic system. <i>Aquatic Microbial Ecology</i> , 2003, 31, 163-174.	0.9	183
98	Unveiling the Organisms behind Novel Eukaryotic Ribosomal DNA Sequences from the Ocean. <i>Applied and Environmental Microbiology</i> , 2002, 68, 4554-4558.	1.4	176
99	Microheterogeneity in 16S Ribosomal DNA-Defined Bacterial Populations from a Stratified Planktonic Environment Is Related to Temporal Changes and to Ecological Adaptations. <i>Applied and Environmental Microbiology</i> , 2002, 68, 1706-1714.	1.4	124
100	Coupled dynamics of dimethylsulfoniopropionate and dimethylsulfide cycling and the microbial food web in surface waters of the North Atlantic. <i>Limnology and Oceanography</i> , 2002, 47, 53-61.	1.6	184
101	Changes in bacterial and archaeal assemblages in an equatorial river induced by the water eutrophication of Petit Saut dam reservoir (French Guiana). <i>Aquatic Microbial Ecology</i> , 2002, 26, 209-221.	0.9	77
102	Prokaryotic plankton biomass and heterotrophic production in western Antarctic waters during the 1995-1996 Austral summer. <i>Deep-Sea Research Part II: Topical Studies in Oceanography</i> , 2002, 49, 805-825.	0.6	48
103	Viral distribution and activity in Antarctic waters. <i>Deep-Sea Research Part II: Topical Studies in Oceanography</i> , 2002, 49, 827-845.	0.6	88
104	Distribution of microbial biomass and importance of protists in regulating prokaryotic assemblages in three areas close to the Antarctic Peninsula in spring and summer 1995/96. <i>Deep-Sea Research Part II: Topical Studies in Oceanography</i> , 2002, 49, 847-867.	0.6	28
105	Dissolved Primary Production and the Strength of Phytoplankton? Bacterioplankton Coupling in Contrasting Marine Regions. <i>Microbial Ecology</i> , 2002, 44, 217-223.	1.4	151
106	Studying marine microorganisms from space. <i>International Microbiology</i> , 2002, 5, 195-200.	1.1	4
107	Changes in archaeal, bacterial and eukaryal assemblages along a salinity gradient by comparison of genetic fingerprinting methods in a multipond solar saltern. <i>Environmental Microbiology</i> , 2002, 4, 338-348.	1.8	433
108	Prokaryotic genetic diversity throughout the salinity gradient of a coastal solar saltern. <i>Environmental Microbiology</i> , 2002, 4, 349-360.	1.8	287

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109	Regulation of bacterial assemblages in oligotrophic plankton systems: results from experimental and empirical approaches. <i>Antonie Van Leeuwenhoek</i> , 2002, 81, 435-452.	0.7	111
110	Seasonal and spatial variations in the nutrient limitation of bacterioplankton growth in the northwestern Mediterranean. <i>Aquatic Microbial Ecology</i> , 2002, 27, 47-56.	0.9	98
111	Spatial distribution of microbial biomass and activity (bacterivory and bacterial production) in the northern Weddell Sea during the austral summer (January 1994). <i>Aquatic Microbial Ecology</i> , 2002, 29, 107-121.	0.9	31
112	Partitioning of phytoplanktonic organic carbon production and bacterial production along a coastal-offshore gradient in the NE Atlantic during different hydrographic regimes. <i>Aquatic Microbial Ecology</i> , 2002, 29, 239-252.	0.9	44
113	Study of Genetic Diversity of Eukaryotic Picoplankton in Different Oceanic Regions by Small-Subunit rRNA Gene Cloning and Sequencing. <i>Applied and Environmental Microbiology</i> , 2001, 67, 2932-2941.	1.4	470
114	Application of Denaturing Gradient Gel Electrophoresis (DGGE) To Study the Diversity of Marine Picoeukaryotic Assemblages and Comparison of DGGE with Other Molecular Techniques. <i>Applied and Environmental Microbiology</i> , 2001, 67, 2942-2951.	1.4	473
115	Changes in marine bacterioplankton phylogenetic composition during incubations designed to measure biogeochemically significant parameters. <i>Limnology and Oceanography</i> , 2001, 46, 1181-1188.	1.6	162
116	In Situ Assessment on the Physiological State of Purple and Green Sulfur Bacteria through the Analyses of Pigment and 5S rRNA Content. <i>Microbial Ecology</i> , 2001, 42, 427-437.	1.4	13
117	Unexpected diversity of small eukaryotes in deep-sea Antarctic plankton. <i>Nature</i> , 2001, 409, 603-607.	13.7	838
118	Composition and temporal dynamics of planktonic archaeal assemblages from anaerobic sulfurous environments studied by 16S rDNA denaturing gradient gel electrophoresis and sequencing. <i>Aquatic Microbial Ecology</i> , 2001, 25, 237-246.	0.9	58
119	Primary production in estuarine oxic/anoxic interfaces: contribution of microbial dark CO ₂ fixation in the Ebro River Salt Wedge Estuary. <i>Marine Ecology - Progress Series</i> , 2001, 215, 49-56.	0.9	31
120	Dissolved and particulate primary production and bacterial production in offshore Antarctic waters during austral summer: coupled or uncoupled?. <i>Marine Ecology - Progress Series</i> , 2001, 222, 25-39.	0.9	66
121	Dissolved and suspended organic carbon in the Atlantic sector of the Southern Ocean. Stock dynamics in upper ocean waters. <i>Marine Ecology - Progress Series</i> , 2001, 223, 27-38.	0.9	10
122	The microbial food web along salinity gradients. , 2000, 32, 143-155.		136
123	Comparative analysis shows that bacterivory, not viral lysis, controls the abundance of heterotrophic prokaryotic plankton. <i>FEMS Microbiology Ecology</i> , 2000, 32, 157-165.	1.3	46
124	Comparison of pure cultures and natural assemblages of planktonic photosynthetic sulfur bacteria by low molecular mass RNA fingerprinting. <i>FEMS Microbiology Ecology</i> , 2000, 32, 25-34.	1.3	17
125	The microbial food web along salinity gradients. <i>FEMS Microbiology Ecology</i> , 2000, 32, 143-155.	1.3	95
126	Spatial differences in bacterioplankton composition along the Catalan coast (NW Mediterranean) assessed by molecular fingerprinting. <i>FEMS Microbiology Ecology</i> , 2000, 33, 51-59.	1.3	187

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127	5S rRNA fingerprints of marine bacteria, halophilic archaea and natural prokaryotic assemblages along a salinity gradient. <i>FEMS Microbiology Ecology</i> , 2000, 34, 113-119.	1.3	39
128	A Few Cosmopolitan Phylotypes Dominate Planktonic Archaeal Assemblages in Widely Different Oceanic Provinces. <i>Applied and Environmental Microbiology</i> , 2000, 66, 1777-1787.	1.4	311
129	Identification of and Spatio-Temporal Differences between Microbial Assemblages from Two Neighboring Sulfurous Lakes: Comparison by Microscopy and Denaturing Gradient Gel Electrophoresis. <i>Applied and Environmental Microbiology</i> , 2000, 66, 499-508.	1.4	392
130	Bacterial Community Structure Associated with a Dimethylsulfoniopropionate-Producing North Atlantic Algal Bloom. <i>Applied and Environmental Microbiology</i> , 2000, 66, 4237-4246.	1.4	402
131	Biological turnover of DMS, DMSP and DMSO in contrasting open-sea waters. <i>Marine Ecology - Progress Series</i> , 2000, 203, 1-11.	0.9	109
132	Role of vertical mixing in controlling the oceanic production of dimethyl sulphide. <i>Nature</i> , 1999, 402, 396-399.	13.7	191
133	Bacterioplankton and phytoplankton biomass and production during summer stratification in the northwestern Mediterranean Sea. <i>Deep-Sea Research Part I: Oceanographic Research Papers</i> , 1999, 46, 985-1019.	0.6	108
134	Short-term variability in the open ocean cycle of dimethylsulfide. <i>Global Biogeochemical Cycles</i> , 1999, 13, 1173-1181.	1.9	110
135	Distribution of viruses and their potential effect on bacterioplankton in an oligotrophic marine system. <i>Aquatic Microbial Ecology</i> , 1999, 19, 205-213.	0.9	49
136	Identification of phototrophic sulfur bacteria through the analysis of 16S rRNA band patterns. <i>Archives of Microbiology</i> , 1998, 170, 269-278.	1.0	13
137	Diel variations in bacterial heterotrophic activity and growth in the northwestern Mediterranean Sea. <i>Marine Ecology - Progress Series</i> , 1998, 164, 107-124.	0.9	170
138	Viral lysis and bacterivory as prokaryotic loss factors along a salinity gradient. <i>Aquatic Microbial Ecology</i> , 1996, 11, 215-227.	0.9	220
139	Microbial plankton across Drake Passage. <i>Polar Biology</i> , 1996, 16, 613-622.	0.5	12
140	Components, structure and fluxes of the microbial food web in a small, stratified lake. <i>Aquatic Microbial Ecology</i> , 1996, 11, 279-288.	0.9	12
141	Microbial plankton across Drake Passage. <i>Polar Biology</i> , 1996, 16, 613-622.	0.5	1
142	Predation by ciliates on a metalimnetic <i>Cryptomonas</i> population: feeding rates, impact and effects of vertical migration. <i>Journal of Plankton Research</i> , 1995, 17, 2131-2154.	0.8	41
143	Occurrence and transformation of dissolved dimethyl sulfur species in stratified seawater (western Tj ETQq1 1 0.784314 rgBT /Overload	0.9	41
144	The problem of species aggregation in food webs. <i>Microbial Ecology</i> , 1994, 28, 201-203.	1.4	0

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145	Impact of <i>Daphnia pulex</i> on a metalimnetic microbial community. <i>Journal of Plankton Research</i> , 1994, 16, 1379-1399.	0.8	27
146	A method to determine integrated predation in stratified waters. <i>Limnology and Oceanography</i> , 1994, 39, 248-262.	1.6	4
147	Thymidine incorporation in Lake Cis: Problems in estimating bacterial secondary production across oxic-anoxic interfaces. <i>FEMS Microbiology Ecology</i> , 1994, 14, 53-64.	1.3	4
148	Effects of Temperature, Sulfide, and Food Abundance on Growth and Feeding of Anaerobic Ciliates. <i>Applied and Environmental Microbiology</i> , 1994, 60, 1317-1324.	1.4	12
149	Role of Anaerobic Ciliates in Planktonic Food Webs: Abundance, Feeding, and Impact on Bacteria in the Field. <i>Applied and Environmental Microbiology</i> , 1994, 60, 1325-1334.	1.4	41
150	Physiological ecology of a metalimnetic <i>Cryptomonas</i> population: relationships to light, sulfide and nutrients. <i>Journal of Plankton Research</i> , 1993, 15, 255-275.	0.8	45
151	Diversity of bacterioplankton. <i>Trends in Ecology and Evolution</i> , 1993, 8, 86-90.	4.2	30
152	Heterotrophic bacterial production in systems of the northern Spanish Mediterranean Region. <i>Verhandlungen Der Internationalen Vereinigung Fur Theoretische Und Angewandte Limnologie International Association of Theoretical and Applied Limnology</i> , 1993, 25, 739-742.	0.1	2
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