James I Prosser

List of Publications by Year
in descending order

229 papers	30,738 citations	3933 88 h-index	4885 168 g-index
240 all docs	240 docs citations	240 times ranked	19896 citing authors

#	Article	IF	Citations
1	Archaea predominate among ammonia-oxidizing prokaryotes in soils. Nature, 2006, 442, 806-809.	27.8	2,144
2	The influence of soil pH on the diversity, abundance and transcriptional activity of ammonia oxidizing archaea and bacteria. Environmental Microbiology, 2008, 10, 2966-2978.	3.8	1,104
3	Soil bacterial networks are less stable under drought than fungal networks. Nature Communications, 2018, 9, 3033.	12.8	992
4	Plant host habitat and root exudates shape soil bacterial community structure. ISME Journal, 2008, 2, 1221-1230.	9.8	958
5	Archaeal and bacterial ammonia-oxidisers in soil: the quest for niche specialisation and differentiation. Trends in Microbiology, 2012, 20, 523-531.	7.7	853
6	The role of ecological theory in microbial ecology. Nature Reviews Microbiology, 2007, 5, 384-392.	28.6	796
7	Analysis of ammonia-oxidizing bacteria of the beta subdivision of the class Proteobacteria in coastal sand dunes by denaturing gradient gel electrophoresis and sequencing of PCR-amplified 16S ribosomal DNA fragments. Applied and Environmental Microbiology, 1997, 63, 1489-1497.	3.1	711
8	Growth, activity and temperature responses of ammoniaâ€oxidizing archaea and bacteria in soil microcosms. Environmental Microbiology, 2008, 10, 1357-1364.	3.8	658
9	Autotrophic Nitrification in Bacteria. Advances in Microbial Physiology, 1990, 30, 125-181.	2.4	636
10	The ecological coherence of high bacterial taxonomic ranks. Nature Reviews Microbiology, 2010, 8, 523-529.	28.6	562
11	Ammonia concentration determines differential growth of ammonia-oxidising archaea and bacteria in soil microcosms. ISME Journal, 2011, 5, 1067-1071.	9.8	543
12	Relative contributions of archaea and bacteria to aerobic ammonia oxidation in the environment. Environmental Microbiology, 2008, 10, 2931-2941.	3.8	531
13	Temperature sensitivity of soil respiration rates enhanced by microbial community response. Nature, 2014, 513, 81-84.	27.8	528
14	Microbes as Engines of Ecosystem Function: When Does Community Structure Enhance Predictions of Ecosystem Processes?. Frontiers in Microbiology, 2016, 7, 214.	3.5	479
15	Cultivation of an obligate acidophilic ammonia oxidizer from a nitrifying acid soil. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 15892-15897.	7.1	464
16	Bacterial diversity promotes community stability and functional resilience after perturbation. Environmental Microbiology, 2005, 7, 301-313.	3.8	429
17	Niche specialization of terrestrial archaeal ammonia oxidizers. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 21206-21211.	7.1	402
18	Molecular diversity of soil and marine 16S rRNA gene sequences related to beta-subgroup ammonia-oxidizing bacteria. Applied and Environmental Microbiology, 1996, 62, 4147-4154.	3.1	378

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19	Molecular Analysis of Bacterial Community Structure and Diversity in Unimproved and Improved Upland Grass Pastures. Applied and Environmental Microbiology, 1999, 65, 1721-1730.	3.1	370
20	Links between Ammonia Oxidizer Community Structure, Abundance, and Nitrification Potential in Acidic Soils. Applied and Environmental Microbiology, 2011, 77, 4618-4625.	3.1	357
21	Archaea rather than bacteria control nitrification in two agricultural acidic soils. FEMS Microbiology Ecology, 2010, 74, 566-574.	2.7	346
22	Autotrophic ammonia oxidation by soil thaumarchaea. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 17240-17245.	7.1	305
23	Impact of Protozoan Grazing on Bacterial Community Structure in Soil Microcosms. Applied and Environmental Microbiology, 2002, 68, 6094-6105.	3.1	300
24	Nitrosospira spp. can produce nitrous oxide via a nitrifier denitrification pathway. Environmental Microbiology, 2006, 8, 214-222.	3.8	287
25	Decline of soil microbial diversity does not influence the resistance and resilience of key soil microbial functional groups following a model disturbance. Environmental Microbiology, 2007, 9, 2211-2219.	3.8	286
26	The consequences of niche and physiological differentiation of archaeal and bacterial ammonia oxidisers for nitrous oxide emissions. ISME Journal, 2018, 12, 1084-1093.	9.8	274
27	Impacts of Soil Faunal Community Composition on Model Grassland Ecosystems. Science, 2002, 298, 615-618.	12.6	260
28	Maintenance of soil functioning following erosion of microbial diversity. Environmental Microbiology, 2006, 8, 2162-2169.	3.8	251
29	Numerical Analysis of Grassland Bacterial Community Structure under Different Land Management Regimens by Using 16S Ribosomal DNA Sequence Data and Denaturing Gradient Gel Electrophoresis Banding Patterns. Applied and Environmental Microbiology, 2001, 67, 4554-4559.	3.1	247
30	Differential photoinhibition of bacterial and archaeal ammonia oxidation. FEMS Microbiology Letters, 2012, 327, 41-46.	1.8	245
31	Analysis of Î²-Subgroup Proteobacterial Ammonia Oxidizer Populations in Soil by Denaturing Gradient Gel Electrophoresis Analysis and Hierarchical Phylogenetic Probing. Applied and Environmental Microbiology, 1998, 64, 2958-2965.	3.1	245
32	Archaea produce lower yields of N₂O than bacteria during aerobic ammonia oxidation in soil. Environmental Microbiology, 2017, 19, 4829-4837.	3.8	243
33	Identification of active methylotroph populations in an acidic forest soil by stable-isotope probing c cThe GenBank accession numbers for the sequences reported in this paper are AY080911â€"AY080961.. Microbiology (United Kingdom), 2002, 148, 2331-2342.	1.8	238
34	Potential bias of fungal 18S rDNA and internal transcribed spacer polymerase chain reaction primers for estimating fungal biodiversity in soil. Environmental Microbiology, 2003, 5, 36-47.	3.8	235
35	Growth of ammonia-oxidizing archaea in soil microcosms is inhibited by acetylene. FEMS Microbiology Ecology, 2009, 70, 99-108.	2.7	235
36	Dispersing misconceptions and identifying opportunities for the use of 'omics' in soil microbial ecology. Nature Reviews Microbiology, 2015, 13, 439-446.	28.6	234

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37	Environmentally-acquired bacteria influence microbial diversity and natural innate immune responses at gut surfaces. BMC Biology, 2009, 7, 79.	3.8	228
38	Replicate or lie. Environmental Microbiology, 2010, 12, 1806-1810.	3.8	227
39	Nitrous oxide production by ammonia oxidizers: Physiological diversity, niche differentiation and potential mitigation strategies. Global Change Biology, 2020, 26, 103-118.	9.5	227
40	EFFECTS OF GRAZING ON MICROBIAL FUNCTIONAL GROUPS INVOLVED IN SOIL N DYNAMICS. Ecological Monographs, 2005, 75, 65-80.	5.4	201
41	Comparative Diversity of Ammonia Oxidizer 16S rRNA Gene Sequences in Native, Tilled, and Successional Soils. Applied and Environmental Microbiology, 1999, 65, 2994-3000.	3.1	200
42	Isolation of â€˜<i>Candidatus</i>Nitrosocosmicus franklandusâ€™, a novel ureolytic soil archaeal ammonia oxidiser with tolerance to high ammonia concentration. FEMS Microbiology Ecology, 2016, 92, fiw057.	2.7	197
43	Effects of management regime and plant species on the enzyme activity and genetic structure of N-fixing, denitrifying and nitrifying bacterial communities in grassland soils. Environmental Microbiology, 2006, 8, 1005-1016.	3.8	196
44	Grassland Management Regimens Reduce Small-Scale Heterogeneity and Species Diversity of Î²-Proteobacterial Ammonia Oxidizer Populations. Applied and Environmental Microbiology, 2002, 68, 20-30.	3.1	187
45	Molecular analysis of enrichment cultures of marine ammonia oxidisers. FEMS Microbiology Letters, 1994, 120, 363-367.	1.8	180
46	Autotrophic Ammonia Oxidation at Low pH through Urea Hydrolysis. Applied and Environmental Microbiology, 2001, 67, 2952-2957.	3.1	180
47	Thaumarchaeal Ammonia Oxidation in an Acidic Forest Peat Soil Is Not Influenced by Ammonium Amendment. Applied and Environmental Microbiology, 2010, 76, 7626-7634.	3.1	180
48	Changes in the community structure and activity of betaproteobacterial ammonia-oxidizing sediment bacteria along a freshwater-marine gradient. Environmental Microbiology, 2006, 8, 684-696.	3.8	172
49	Effects of Agronomic Treatments on Structure and Function of Ammonia-Oxidizing Communities. Applied and Environmental Microbiology, 2000, 66, 5410-5418.	3.1	166
50	Diversity of fungi in organic soils under a moorland - Scots pine (Pinus sylvestris L.) gradient. Environmental Microbiology, 2003, 5, 1121-1132.	3.8	166
51	Stimulation of thaumarchaeal ammonia oxidation by ammonia derived from organic nitrogen but not added inorganic nitrogen. FEMS Microbiology Ecology, 2012, 80, 114-123.	2.7	160
52	Cell density-regulated recovery of starved biofilm populations of ammonia-oxidizing bacteria. Applied and Environmental Microbiology, 1997, 63, 2281-2286.	3.1	160
53	Molecular marker systems for detection of genetically engineered micro-organisms in the environment. Microbiology (United Kingdom), 1994, 140, 5-17.	1.8	156
54	Links between ammonia oxidizer species composition, functional diversity and nitrification kinetics in grassland soils. Environmental Microbiology, 2005, 7, 676-684.	3.8	156

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55	Altitude ammonia-oxidizing bacteria and archaea in soils of Mount Everest. FEMS Microbiology Ecology, 2009, 70, 208-217.	2.7	155
56	Growth Mechanisms and Growth Kinetics of Filamentous Microorganisms. Critical Reviews in Biotechnology, 1991, 10, 253-274.	9.0	154
57	Spatial structure in soil chemical and microbiological properties in an upland grassland. FEMS Microbiology Ecology, 2004, 49, 191-205.	2.7	154
58	Nitrogen Cycling and Community Structure of Proteobacterial Î²-Subgroup Ammonia-Oxidizing Bacteria within Polluted Marine Fish Farm Sediments. Applied and Environmental Microbiology, 1999, 65, 213-220.	3.1	153
59	Stable isotope probing analysis of the influence of liming on root exudate utilization by soil microorganisms. Environmental Microbiology, 2005, 7, 828-838.	3.8	153
60	Correlation of Methane Production and Functional Gene Transcriptional Activity in a Peat Soil. Applied and Environmental Microbiology, 2009, 75, 6679-6687.	3.1	152
61	Community Structure of Ammonia-Oxidizing Bacteria within Anoxic Marine Sediments. Applied and Environmental Microbiology, 2003, 69, 1359-1371.	3.1	151
62	Bacterial Biodiversity-Ecosystem Functioning Relations Are Modified by Environmental Complexity. PLoS ONE, 2010, 5, e10834.	2.5	149
63	Primary succession of soil Crenarchaeota across a receding glacier foreland. Environmental Microbiology, 2005, 7, 337-347.	3.8	145
64	Links between Plant and Rhizoplane Bacterial Communities in Grassland Soils, Characterized Using Molecular Techniques. Applied and Environmental Microbiology, 2005, 71, 6784-6792.	3.1	144
65	Soil pH regulates the abundance and diversity of Group 1.1c Crenarchaeota. FEMS Microbiology Ecology, 2009, 70, 367-376.	2.7	143
66	Characterisation of terrestrial acidophilic archaeal ammonia oxidisers and their inhibition and stimulation by organic compounds. FEMS Microbiology Ecology, 2014, 89, 542-552.	2.7	141
67	Bacterial Origin and Community Composition in the Barley Phytosphere as a Function of Habitat and Presowing Conditions. Applied and Environmental Microbiology, 2000, 66, 4372-4377.	3.1	140
68	Luminescence-based nonextractive technique for in situ detection of Escherichia coli in soil. Applied and Environmental Microbiology, 1990, 56, 3368-3374.	3.1	140
69	Influence of Inorganic Nitrogen Management Regime on the Diversity of Nitrite-Oxidizing Bacteria in Agricultural Grassland Soils. Applied and Environmental Microbiology, 2005, 71, 8323-8334.	3.1	139
70	A Model for Hyphal Growth and Branching. Journal of General Microbiology, 1979, 111, 153-164.	2.3	136
71	Effects of aboveground grazing on coupling among nitrifier activity, abundance and community structure. ISME Journal, 2008, 2, 221-232.	9.8	134
72	Environmental and spatial characterisation of bacterial community composition in soil to inform sampling strategies. Soil Biology and Biochemistry, 2009, 41, 2292-2298.	8.8	130

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73	Diversity of Bacteria Associated with Natural AphidPopulations. Applied and Environmental Microbiology, 2003, 69, 7216-7223.	3.1	129
74	The Family Nitrosomonadaceae. , 2014, , 901-918.		127
75	Phylogenetic Differences between Particle-Associated and Planktonic Ammonia-Oxidizing Bacteria of the Î² Subdivision of the Class <i>Proteobacteria</i> in the Northwestern Mediterranean Sea. Applied and Environmental Microbiology, 1999, 65, 779-786.	3.1	127
76	Characterisation of bacterial communities associated with toxic and non-toxic dinoflagellates: Alexandrium spp. and Scrippsiella trochoidea. FEMS Microbiology Ecology, 2001, 37, 161-173.	2.7	126
77	Luminescence-based detection of activity of starved and viable but nonculturable bacteria. Applied and Environmental Microbiology, 1994, 60, 1308-1316.	3.1	124
78	Molecular and functional diversity in soil micro-organisms. Plant and Soil, 2002, 244, 9-17.	3.7	120
79	Establishment of Normal Gut Microbiota Is Compromised under Excessive Hygiene Conditions. PLoS ONE, 2011, 6, e28284.	2.5	120
80	Molecular Analysis of a Bacterial Chitinolytic Community in an Upland Pasture. Applied and Environmental Microbiology, 2002, 68, 5042-5050.	3.1	118
81	Identifying Potential Mechanisms Enabling Acidophily in the Ammonia-Oxidizing Archaeon â€œCandidatus Nitrosotalea devanaterraâ€. Applied and Environmental Microbiology, 2016, 82, 2608-2619.	3.1	117
82	Chemotaxonomic characterisation of the thaumarchaeal lipidome. Environmental Microbiology, 2017, 19, 2681-2700.	3.8	117
83	Ecosystem processes and interactions in a morass of diversity. FEMS Microbiology Ecology, 2012, 81, 507-519.	2.7	111
84	Role of Pore Size Location in Determining Bacterial Activity during Predation by Protozoa in Soil. Applied and Environmental Microbiology, 1995, 61, 3537-3543.	3.1	108
85	Ammoniaâ€oxidising archaea living at low pH: Insights from comparative genomics. Environmental Microbiology, 2017, 19, 4939-4952.	3.8	107
86	Cultivation-independent in situ molecular analysis of bacteria involved in degradation of pentachlorophenol in soil. Environmental Microbiology, 2005, 7, 1349-1360.	3.8	104
87	Abiotic Conversion of Extracellular NH₂OH Contributes to N₂O Emission during Ammonia Oxidation. Environmental Science & Technology, 2017, 51, 13122-13132.	10.0	104
88	Coupling of diversification and pH adaptation during the evolution of terrestrial Thaumarchaeota. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 9370-9375.	7.1	98
89	The impact of zero-valent iron nanoparticles on a river water bacterial community. Journal of Hazardous Materials, 2010, 184, 73-80.	12.4	97
90	The impact of grassland management on archaeal community structure in upland pasture rhizosphere soil. Environmental Microbiology, 2003, 5, 152-162.	3.8	96

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91	pH as a Driver for Ammonia-Oxidizing Archaea in Forest Soils. Microbial Ecology, 2015, 69, 879-883.	2.8	95
92	Phylogenetic congruence and ecological coherence in terrestrial Thaumarchaeota. ISME Journal, 2016, 10, 85-96.	9.8	94
93	Identity of active methanotrophs in landfill cover soil as revealed by DNA-stable isotope probing. FEMS Microbiology Ecology, 2007, 62, 12-23.	2.7	92
94	Spatial Analysis of Archaeal Community Structure inGrasslandSoil. Applied and Environmental Microbiology, 2003, 69, 7420-7429.	3.1	91
95	Links between methane flux and transcriptional activities of methanogens and methane oxidizers in a blanket peat bog. FEMS Microbiology Ecology, 2010, 73, no-no.	2.7	91
96	Impact of cultivation on characterisation of species composition of soil bacterial communities. FEMS Microbiology Ecology, 2001, 35, 37-48.	2.7	88
97	Relationship between assemblages of mycorrhizal fungi and bacteria on grass roots. Environmental Microbiology, 2008, 10, 534-541.	3.8	86
98	Effect of nitrification inhibitors on the growth and activity of Nitrosotalea devanaterra in culture and soil. Soil Biology and Biochemistry, 2013, 62, 129-133.	8.8	86
99	Kinetics of NH₃â€oxidation, NOâ€turnover, N₂Oâ€production and electron flow during oxygen depletion in model bacterial and archaeal ammonia oxidisers. Environmental Microbiology, 2017, 19, 4882-4896.	3.8	86
100	Cultivation-based and molecular approaches to characterisation of terrestrial and aquatic nitrifiers. Antonie Van Leeuwenhoek, 2002, 81, 165-179.	1.7	85
101	Steady state and transient growth of autotrophic nitrifying bacteria. Archives of Microbiology, 1987, 147, 73-79.	2.2	81
102	Studying plantâ€“microbe interactions using stable isotope technologies. Current Opinion in Biotechnology, 2006, 17, 98-102.	6.6	78
103	Detection of a Single Genetically Modified Bacterial Cell in Soil by Using Charge Coupled Device-Enhanced Microscopy. Applied and Environmental Microbiology, 1992, 58, 2444-2448.	3.1	78
104	Differential effects of microorganism-invertebrate interactions on benthic nitrogen cycling. FEMS Microbiology Ecology, 2012, 82, 11-22.	2.7	76
105	Plant nitrogen-use strategy as a driver of rhizosphere archaeal and bacterial ammonia oxidiser abundance. FEMS Microbiology Ecology, 2016, 92, fiw091.	2.7	76
106	Abundance and community structure of sulfate reducing prokaryotes in a paddy soil of southern China under different fertilization regimes. Soil Biology and Biochemistry, 2009, 41, 687-694.	8.8	74
107	Plasmid and chromosomally encoded luminescence marker systems for detection of Pseudomonas fluorescens in soil. Molecular Ecology, 1993, 2, 47-54.	3.9	73
108	Ammonium supply rate influences archaeal and bacterial ammonia oxidizers in a wetland soil vertical profile. FEMS Microbiology Ecology, 2010, 74, 302-315.	2.7	72

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109	Effect of earthworms on the community structure of active methanotrophic bacteria in a landfill cover soil. ISME Journal, 2008, 2, 92-104.	9.8	71
110	Resource availability influences the diversity of a functional group of heterotrophic soil bacteria. Environmental Microbiology, 2008, 10, 2245-2256.	3.8	71
111	Nutrient Amendments in Soil DNA Stable Isotope Probing Experiments Reduce the Observed Methanotroph Diversity. Applied and Environmental Microbiology, 2007, 73, 798-807.	3.1	70
112	Ammonia oxidation is not required for growth of GroupÂ1.1c soil Thaumarchaeota. FEMS Microbiology Ecology, 2015, 91, .	2.7	70
113	Phylogeny of nitrite reductase (nirK) and nitric oxide reductase (norB) genes fromNitrosospiraspecies isolated from soil. FEMS Microbiology Letters, 2007, 266, 83-89.	1.8	69
114	Quantitative analysis of ammonia oxidising bacteria using competitive PCR. FEMS Microbiology Ecology, 2000, 32, 167-175.	2.7	63
115	The ribulose-1,5-bisphosphate carboxylase/oxygenase gene cluster of Methylococcus capsulatus (Bath). Archives of Microbiology, 2002, 177, 279-289.	2.2	63
116	Characterization of rhizosphere colonization by luminescent Enterobacter cloacae at the population and single-cell levels. Applied and Environmental Microbiology, 1995, 61, 2950-2957.	3.1	63
117	Differential response of nonadapted ammonia-oxidising archaea and bacteria to drying-rewetting stress. FEMS Microbiology Ecology, 2014, 90, n/a-n/a.	2.7	61
118	Temperature responses of soil ammonia-oxidising archaea depend on pH. Soil Biology and Biochemistry, 2017, 106, 61-68.	8.8	58
119	Matric potential and the survival and activity of a Pseudomonas fluorescens inoculum in soil. Soil Biology and Biochemistry, 1995, 27, 881-892.	8.8	57
120	Luminescence-Based Systems for Detection of Bacteria in the Environment. Critical Reviews in Biotechnology, 1996, 16, 157-183.	9.0	57
121	The application of high-throughput sequencing technology to analysis of amoA phylogeny and environmental niche specialisation of terrestrial bacterial ammonia-oxidisers. Environmental Microbiomes, 2019, 14, 3.	5.0	53
122	Quantification of the presence and activity of specific microorganisms in nature. Molecular Biotechnology, 1997, 7, 103-120.	2.4	52
123	Stable Isotope Probing Analysis of Interactions between Ammonia Oxidizers. Applied and Environmental Microbiology, 2010, 76, 2468-2477.	3.1	50
124	Selective inhibition of ammonia oxidising archaea by simvastatin stimulates growth of ammonia oxidising bacteria. Soil Biology and Biochemistry, 2020, 141, 107673.	8.8	49
125	Models of Microbial Interactions in the Soil. CRC Critical Reviews in Microbiology, 1976, 4, 463-498.	4.8	48
126	Potential luminescence as an indicator of activation of genetically-modified Pseudomonas fluorescens in liquid culture and in soil. Soil Biology and Biochemistry, 1994, 26, 747-755.	8.8	46

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127	Crenarchaeal community assembly and microdiversity in developing soils at two sites associated with deglaciation. Environmental Microbiology, 2006, 8, 1382-1393.	3.8	46
128	Apical hyphal extension in Streptomyces coelicolor A3(2). Journal of General Microbiology, 1990, 136, 1077-1084.	2.3	45
129	Differential response of archaeal and bacterial communities to nitrogen inputs and pH changes in upland pasture rhizosphere soil. Environmental Microbiology, 2004, 6, 861-867.	3.8	44
130	Carbon flow in an upland grassland: effect of liming on the flux of recently photosynthesized carbon to rhizosphere soil. Global Change Biology, 2004, 10, 2100-2108.	9.5	43
131	Protection of Nitrosomonas europaea colonizing clay minerals from inhibition by nitrapyrin. Journal of General Microbiology, 1991, 137, 1923-1929.	2.3	42
132	Molecular analysis of methanogenic archaeal communities in managed and natural upland pasture soils. Global Change Biology, 2003, 9, 1451-1457.	9.5	42
133	Mineralization of Chitin in an estuarine sediment: The importance of the Chitosan pathway. Biochemical Systematics and Ecology, 1991, 19, 395-400.	1.3	41
134	Kinetics of Filamentous Growth and Branching. , 1995, , 301-318.		41
135	Rhizosphere bacterial community composition responds to arbuscular mycorrhiza, but not to reductions in microbial activity induced by foliar cutting. FEMS Microbiology Ecology, 2008, 64, 78-89.	2.7	41
136	The Role of Microbial Community Composition in Controlling Soil Respiration Responses to Temperature. PLoS ONE, 2016, 11, e0165448.	2.5	41
137	Right and left handed helicity of chitin microfibrils in stipe cells inCoprinus cinereus. Protoplasma, 1991, 165, 64-70.	2.1	40
138	Effect of anoxia and high sulphide concentrations on heterotrophic microbial communities in reduced surface sediments (Black Spots) in sandy intertidal flats of the German Wadden Sea. FEMS Microbiology Ecology, 2003, 44, 291-301.	2.7	40
139	Comparison of PCR primer-based strategies for characterization of ammonia oxidizer communities in environmental samples. FEMS Microbiology Ecology, 2006, 56, 482-493.	2.7	40
140	The life beneath our feet. Nature, 2013, 494, 40-41.	27.8	40
141	Inhibition of Ammonium Oxidation by Nitrapyrin in Soil and Liquid Culture. Applied and Environmental Microbiology, 1986, 52, 782-787.	3.1	40
142	Surface attachment of nitrifying bacteria and their inhibition by potassium ethyl xanthate. Microbial Ecology, 1987, 14, 129-139.	2.8	39
143	The contribution of ammonia-oxidizing archaea and bacteria to gross nitrification under different substrate availability. Soil Biology and Biochemistry, 2021, 160, 108353.	8.8	39
144	A model for bacterial conjugal gene transfer on solid surfaces. FEMS Microbiology Ecology, 2003, 44, 67-78.	2.7	37

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145	The influence of synthetic sheep urine on ammonia oxidizing bacterial communities in grassland soil. FEMS Microbiology Ecology, 2006, 56, 444-454.	2.7	37
146	Putting science back into microbial ecology: a question of approach. Philosophical Transactions of the Royal Society B: Biological Sciences, 2020, 375, 20190240.	4.0	37
147	The surface growth and activity ofNitrobacter. Microbial Ecology, 1988, 15, 21-39.	2.8	35
148	Afforestation of moorland leads to changes in crenarchaeal community structure. FEMS Microbiology Ecology, 2007, 60, 51-59.	2.7	35
149	PCR profiling of ammonia-oxidizer communities in acidic soils subjected to nitrogen and sulphur deposition. FEMS Microbiology Ecology, 2007, 61, 305-316.	2.7	35
150	Stream drying drives microbial ammonia oxidation and firstâ€flush nitrate export. Ecology, 2016, 97, 2192-2198.	3.2	35
151	Diversity of Endospore-forming Bacteria in Soil: Characterization and Driving Mechanisms. Soil Biology, 2011, , 31-59.	0.8	35
152	Role of functionally dominant species in varying environmental regimes: evidence for the performance-enhancing effect of biodiversity. BMC Ecology, 2012, 12, 14.	3.0	34
153	Flux and turnover of fixed carbon in soil microbial biomass of limed and unlimed plots of an upland grassland ecosystem. Environmental Microbiology, 2005, 7, 544-552.	3.8	31
154	Sheep-urine-induced changes in soil microbial community structure. FEMS Microbiology Ecology, 2006, 56, 310-320.	2.7	29
155	Revisiting plant biological nitrification inhibition efficiency using multiple archaeal and bacterial ammonia-oxidising cultures. Biology and Fertility of Soils, 2022, 58, 241-249.	4.3	29
156	Differences between Betaproteobacterial Ammonia-Oxidizing Communities in Marine Sediments and Those in Overlying Water. Applied and Environmental Microbiology, 2004, 70, 3789-3793.	3.1	28
157	Soil Nitrifiers and Nitrification. , 2014, , 347-383.		28
158	Experimental verification of a mathematical model for pelleted growth of Streptomyces coelicolor A3(2) in submerged batch culture. Microbiology (United Kingdom), 1996, 142, 639-648.	1.8	28
159	Use and abuse of potential rates in soil microbiology. Soil Biology and Biochemistry, 2021, 157, 108242.	8.8	26
160	Novel method for the study of the population dynamics of a genetically modified microorganism in the gut of the earthworm Lumbricus terrestris. Biology and Fertility of Soils, 1993, 15, 55-59.	4.3	25
161	Influence of repeated prescribed burning on incorporation of 13C from cellulose by forest soil fungi as determined by RNA stable isotope probing. Soil Biology and Biochemistry, 2009, 41, 467-472.	8.8	25
162	Effect of arbuscular mycorrhizal colonisation on the growth and phosphorus nutrition of Populus euramericana c.v. Ghoy. Biomass and Bioenergy, 2011, 35, 4605-4612.	5.7	25

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163	The role of the earthworm Lumbricus terrestris in the transport of bacterial inocula through soil. Biology and Fertility of Soils, 1996, 23, 132-139.	4.3	24
164	Functional principal component data analysis: A new method for analysing microbial community fingerprints. Journal of Microbiological Methods, 2009, 79, 89-95.	1.6	24
165	Survival and Activity of lux-Marked Aeromonas salmonicida in Seawater. Applied and Environmental Microbiology, 1995, 61, 3494-3498.	3.1	24
166	Luminometric measurement of population activity of genetically modified Pseudomonas fluorescens in the soil. FEMS Microbiology Letters, 1992, 99, 217-220.	1.8	22
167	Activity of the ammonia oxidising bacteria is responsible for zinc tolerance development of the ammonia oxidising community in soil: A stable isotope probing study. Soil Biology and Biochemistry, 2013, 58, 244-247.	8.8	21
168	Experimental testing of hypotheses for temperatureâ€•and <scp>pH</scp>â€based niche specialization of ammonia oxidizing archaea and bacteria. Environmental Microbiology, 2020, 22, 4032-4045.	3.8	21
169	Conceptual challenges in microbial community ecology. Philosophical Transactions of the Royal Society B: Biological Sciences, 2020, 375, 20190241.	4.0	21
170	Chronic Environmental Perturbation Influences Microbial Community Assembly Patterns. Environmental Science & Technology, 2022, 56, 2300-2311.	10.0	21
171	Degradation of metalaxyl-M in contrasting soils is influenced more by differences in physicochemical characteristics than in microbial community composition after re-inoculation of sterilised soils. Soil Biology and Biochemistry, 2010, 42, 1123-1131.	8.8	20
172	Luminescence based detection of Erwinia carotovora subsp. Carotovora in soil. Soil Biology and Biochemistry, 1992, 24, 961-967.	8.8	19
173	Effect of temperature and plasmid carriage on nonculturability in organisms targeted for release. FEMS Microbiology Ecology, 1995, 17, 229-237.	2.7	19
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