Hang-Wei Hu

List of Publications by Year
in descending order

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Version: 2024-02-01

266 papers	19,428 citations	11651 70 h-index	15732 125 g-index
271 all docs	271 docs citations	271 times ranked	12460 citing authors

#	Article	IF	Citations
1	Quantitative analyses of the abundance and composition of ammoniaâ€oxidizing bacteria and ammoniaâ€oxidizing archaea of a Chinese upland red soil under longâ€term fertilization practices. Environmental Microbiology, 2007, 9, 2364-2374.	3.8	877
2	An overview of microplastic and nanoplastic pollution in agroecosystems. Science of the Total Environment, 2018, 627, 1377-1388.	8.0	846
3	Ammonia-oxidizing archaea have more important role than ammonia-oxidizing bacteria in ammonia oxidation of strongly acidic soils. ISME Journal, 2012, 6, 1032-1045.	9.8	614
4	Multiple elements of soil biodiversity drive ecosystem functions across biomes. Nature Ecology and Evolution, 2020, 4, 210-220.	7.8	543
5	Microbial regulation of terrestrial nitrous oxide formation: understanding the biological pathways for prediction of emission rates. FEMS Microbiology Reviews, 2015, 39, 729-749.	8.6	530
6	Abundance and composition of ammoniaâ€oxidizing bacteria and ammoniaâ€oxidizing archaea communities of an alkaline sandy loam. Environmental Microbiology, 2008, 10, 1601-1611.	3.8	508
7	Ammonia-oxidizing bacteria and archaea grow under contrasting soil nitrogen conditions. FEMS Microbiology Ecology, 2010, 72, 386-394.	2.7	419
8	Host selection shapes crop microbiome assembly and network complexity. New Phytologist, 2021, 229, 1091-1104.	7.3	349
9	Ammoniaâ€oxidizing archaea: important players in paddy rhizosphere soil?. Environmental Microbiology, 2008, 10, 1978-1987.	3.8	340
10	Microbial regulation of the soil carbon cycle: evidence from geneâ€“enzyme relationships. ISME Journal, 2016, 10, 2593-2604.	9.8	324
11	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
12	Effects of Cd and Pb on soil microbial community structure and activities. Environmental Science and Pollution Research, 2010, 17, 288-296.	5.3	304
13	Transfer of antibiotic resistance from manure-amended soils to vegetable microbiomes. Environment International, 2019, 130, 104912.	10.0	278
14	Protist communities are more sensitive to nitrogen fertilization than other microorganisms in diverse agricultural soils. Microbiome, 2019, 7, 33.	11.1	278
15	Current insights into the autotrophic thaumarchaeal ammonia oxidation in acidic soils. Soil Biology and Biochemistry, 2012, 55, 146-154.	8.8	268
16	Rare microbial taxa as the major drivers of ecosystem multifunctionality in long-term fertilized soils. Soil Biology and Biochemistry, 2020, 141, 107686.	8.8	247
17	Long-Term Nickel Contamination Increases the Occurrence of Antibiotic Resistance Genes in Agricultural Soils. Environmental Science & Technology, 2017, 51, 790-800.	10.0	240
18	pH-dependent distribution of soil ammonia oxidizers across a large geographical scale as revealed by high-throughput pyrosequencing. Journal of Soils and Sediments, 2013, 13, 1439-1449.	3.0	219

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19	Fieldâ€based evidence for copper contamination induced changes of antibiotic resistance in agricultural soils. Environmental Microbiology, 2016, 18, 3896-3909.	3.8	216
20	Plant diversity represents the prevalent determinant of soil fungal community structure across temperate grasslands in northern China. Soil Biology and Biochemistry, 2017, 110, 12-21.	8.8	202
21	Comammoxâ€”a newly discovered nitrification process in the terrestrial nitrogen cycle. Journal of Soils and Sediments, 2017, 17, 2709-2717.	3.0	194
22	A review of ammonia-oxidizing bacteria and archaea in Chinese soils. Frontiers in Microbiology, 2012, 3, 296.	3.5	191
23	Aerobic composting reduces antibiotic resistance genes in cattle manure and the resistome dissemination in agricultural soils. Science of the Total Environment, 2018, 612, 1300-1310.	8.0	190
24	Differences in soil bacterial diversity: driven by contemporary disturbances or historical contingencies?. ISME Journal, 2008, 2, 254-264.	9.8	182
25	Long-term fertilization regimes affect bacterial community structure and diversity of an agricultural soil in northern China. Journal of Soils and Sediments, 2008, 8, 43-50.	3.0	177
26	Microbial composition and diversity of an upland red soil under long-term fertilization treatments as revealed by culture-dependent and culture-independent approaches. Journal of Soils and Sediments, 2008, 8, 349-358.	3.0	170
27	Antibiotic resistance genes and associated bacterial communities in agricultural soils amended with different sources of animal manures. Soil Biology and Biochemistry, 2018, 126, 91-102.	8.8	170
28	Altitudinal Distribution Patterns of Soil Bacterial and Archaeal Communities Along Mt. Shegyla on the Tibetan Plateau. Microbial Ecology, 2015, 69, 135-145.	2.8	166
29	Temporal succession of soil antibiotic resistance genes following application of swine, cattle and poultry manures spiked with or without antibiotics. Environmental Pollution, 2017, 231, 1621-1632.	7.5	166
30	Plant developmental stage drives the differentiation in ecological role of the maize microbiome. Microbiome, 2021, 9, 171.	11.1	164
31	Impact of long-term fertilization practices on the abundance and composition of soil bacterial communities in Northeast China. Applied Soil Ecology, 2010, 46, 119-124.	4.3	158
32	Altitude ammonia-oxidizing bacteria and archaea in soils of Mount Everest. FEMS Microbiology Ecology, 2009, 70, 208-217.	2.7	155
33	Comammox Nitrospira play an active role in nitrification of agricultural soils amended with nitrogen fertilizers. Soil Biology and Biochemistry, 2019, 138, 107609.	8.8	143
34	Multivariate geostatistical analysis of heavy metals in topsoils from Beijing, China. Journal of Soils and Sediments, 2008, 8, 51-58.	3.0	136
35	New insights into the role of microbial community composition in driving soil respiration rates. Soil Biology and Biochemistry, 2018, 118, 35-41.	8.8	134
36	Rare taxa maintain the stability of crop mycobiomes and ecosystem functions. Environmental Microbiology, 2021, 23, 1907-1924.	3.8	132

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37	Niche separation of comammox Nitrospira and canonical ammonia oxidizers in an acidic subtropical forest soil under long-term nitrogen deposition. Soil Biology and Biochemistry, 2018, 126, 114-122.	8.8	129
38	Nitrous oxide emissions from grazed grassland as affected by a nitrification inhibitor, dicyandiamide, and relationships with ammonia-oxidizing bacteria and archaea. Journal of Soils and Sediments, 2010, 10, 943-954.	3.0	122
39	Analysis of the Microbial Community Structure by Monitoring an Hg Methylation Gene (hgcA) in Paddy Soils along an Hg Gradient. Applied and Environmental Microbiology, 2014, 80, 2874-2879.	3.1	119
40	Contrasting patterns and drivers of soil bacterial and fungal diversity across a mountain gradient. Environmental Microbiology, 2020, 22, 3287-3301.	3.8	119
41	Responses of ammonia-oxidizing bacteria and archaea to nitrogen fertilization and precipitation increment in a typical temperate steppe in Inner Mongolia. Applied Soil Ecology, 2013, 68, 36-45.	4.3	116
42	Speciation, transportation, and pathways of cadmium in soil-rice systems: A review on the environmental implications and remediation approaches for food safety. Environment International, 2021, 156, 106749.	10.0	116
43	Soil bacterial taxonomic diversity is critical to maintaining the plant productivity. Environment International, 2020, 140, 105766.	10.0	114
44	Soil pH determines the alpha diversity but not beta diversity of soil fungal community along altitude in a typical Tibetan forest ecosystem. Journal of Soils and Sediments, 2015, 15, 1224-1232.	3.0	112
45	Water addition regulates the metabolic activity of ammonia oxidizers responding to environmental perturbations in dry subhumid ecosystems. Environmental Microbiology, 2015, 17, 444-461.	3.8	111
46	Ammonia-Oxidizing Archaea Play a Predominant Role in Acid Soil Nitrification. Advances in Agronomy, 2014, , 261-302.	5.2	109
47	Consistent responses of soil microbial taxonomic and functional attributes to mercury pollution across China. Microbiome, 2018, 6, 183.	11.1	109
48	Temporal changes of antibiotic-resistance genes and bacterial communities in two contrasting soils treated with cattle manure. FEMS Microbiology Ecology, 2016, 92, fiv169.	2.7	108
49	Fungal richness contributes to multifunctionality in boreal forest soil. Soil Biology and Biochemistry, 2019, 136, 107526.	8.8	108
50	Unraveling Microbial Communities Associated with Methylmercury Production in Paddy Soils. Environmental Science & Technology, 2018, 52, 13110-13118.	10.0	106
51	The effects of short term, long term and reapplication of biochar on soil bacteria. Science of the Total Environment, 2018, 636, 142-151.	8.0	105
52	Abundance and community structure of ammonia-oxidizing archaea and bacteria in an acid paddy soil. Biology and Fertility of Soils, 2011, 47, 323-331.	4.3	102
53	Nitrogen loading levels affect abundance and composition of soil ammonia oxidizing prokaryotes in semiarid temperate grassland. Journal of Soils and Sediments, 2011, 11, 1243-1252.	3.0	100
54	Putative ammoniaâ€oxidizing bacteria and archaea in an acidic red soil with different land utilization patterns. Environmental Microbiology Reports, 2010, 2, 304-312.	2.4	92

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55	Contrasting Euryarchaeota communities between upland and paddy soils exhibited similar pH-impacted biogeographic patterns. Soil Biology and Biochemistry, 2013, 64, 18-27.	8.8	92
56	Effects of climate warming and elevated CO 2 on autotrophic nitrification and nitrifiers in dryland ecosystems. Soil Biology and Biochemistry, 2016, 92, 1-15.	8.8	92
57	Distinct microbial communities in the active and permafrost layers on the Tibetan Plateau. Molecular Ecology, 2017, 26, 6608-6620.	3.9	92
58	Distribution and diversity of archaeal communities in selected Chinese soils. FEMS Microbiology Ecology, 2012, 80, 146-158.	2.7	91
59	Abundance and community composition of methanotrophs in a Chinese paddy soil under long-term fertilization practices. Journal of Soils and Sediments, 2008, 8, 406-414.	3.0	90
60	Effects of the Nitrification Inhibitor 3,4-Dimethylpyrazole Phosphate on Nitrification and Nitrifiers in Two Contrasting Agricultural Soils. Applied and Environmental Microbiology, 2016, 82, 5236-5248.	3.1	90
61	Influence of nitrogen fertilization on soil ammonia oxidizer and denitrifier abundance, microbial biomass, and enzyme activities in an alpine meadow. Biology and Fertility of Soils, 2014, 50, 703-713.	4.3	84
62	Long-term manure application increased the levels of antibiotics and antibiotic resistance genes in a greenhouse soil. Applied Soil Ecology, 2017, 121, 193-200.	4.3	84
63	Effects of nitrogen application rate and a nitrification inhibitor dicyandiamide on ammonia oxidizers and N2O emissions in a grazed pasture soil. Science of the Total Environment, 2013, 465, 125-135.	8.0	83
64	Global homogenization of the structure and function in the soil microbiome of urban greenspaces. Science Advances, 2021, 7, .	10.3	83
65	Soil type determines the abundance and community structure of ammonia-oxidizing bacteria and archaea in flooded paddy soils. Journal of Soils and Sediments, 2010, 10, 1510-1516.	3.0	82
66	Activity, abundance and community structure of anammox bacteria along depth profiles in three different paddy soils. Soil Biology and Biochemistry, 2015, 91, 212-221.	8.8	82
67	Abundance and community structure of ammonia-oxidizing bacteria and archaea in a temperate forest ecosystem under ten-years elevated CO2. Soil Biology and Biochemistry, 2012, 46, 163-171.	8.8	81
68	Microbial communities in crop phyllosphere and root endosphere are more resistant than soil microbiota to fertilization. Soil Biology and Biochemistry, 2021, 153, 108113.	8.8	81
69	Adaptive responses of comammox Nitrospira and canonical ammonia oxidizers to long-term fertilizations: Implications for the relative contributions of different ammonia oxidizers to soil nitrogen cycling. Science of the Total Environment, 2019, 668, 224-233.	8.0	79
70	Effects of Cellular Sorption on Mercury Bioavailability and Methylmercury Production by Desulfovibrio desulfuricans ND132. Environmental Science & Technology, 2016, 50, 13335-13341.	10.0	78
71	Impacts of reclaimed water irrigation on soil antibiotic resistome in urban parks of Victoria, Australia. Environmental Pollution, 2016, 211, 48-57.	7.5	78
72	Identity of biocrust species and microbial communities drive the response of soil multifunctionality to simulated global change. Soil Biology and Biochemistry, 2017, 107, 208-217.	8.8	78

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73	Fertilization changes soil microbiome functioning, especially phagotrophic protists. Soil Biology and Biochemistry, 2020, 148, 107863.	8.8	78
74	Arsenic and cadmium as predominant factors shaping the distribution patterns of antibiotic resistance genes in polluted paddy soils. Journal of Hazardous Materials, 2020, 389, 121838.	12.4	77
75	Microbial Community and Functional Structure Significantly Varied among Distinct Types of Paddy Soils But Responded Differently along Gradients of Soil Depth Layers. Frontiers in Microbiology, 2017, 8, 945.	3.5	76
76	Nitrifierâ€induced denitrification is an important source of soil nitrous oxide and can be inhibited by a nitrification inhibitor 3,4â€dimethylpyrazole phosphate. Environmental Microbiology, 2017, 19, 4851-4865.	3.8	75
77	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
78	Response of denitrification genes nirS, nirK, and nosZ to irrigation water quality in a Chinese agricultural soil. Environmental Science and Pollution Research, 2011, 18, 1644-1652.	5.3	70
79	The large-scale distribution of ammonia oxidizers in paddy soils is driven by soil pH, geographic distance, and climatic factors. Frontiers in Microbiology, 2015, 6, 938.	3.5	70
80	Palaeoclimate explains a unique proportion of the global variation in soil bacterial communities. Nature Ecology and Evolution, 2017, 1, 1339-1347.	7.8	70
81	The effect of temperature and moisture on the source of N2O and contributions from ammonia oxidizers in an agricultural soil. Biology and Fertility of Soils, 2017, 53, 141-152.	4.3	69
82	Nitrogen fertiliser-induced changes in N2O emissions are attributed more to ammonia-oxidising bacteria rather than archaea as revealed using 1-octyne and acetylene inhibitors in two arable soils. Biology and Fertility of Soils, 2016, 52, 1163-1171.	4.3	65
83	Dissimilatory nitrate reduction to ammonium dominates nitrate reduction in long-term low nitrogen fertilized rice paddies. Soil Biology and Biochemistry, 2019, 131, 149-156.	8.8	64
84	Microbial regulation of natural antibiotic resistance: Understanding the protist-bacteria interactions for evolution of soil resistome. Science of the Total Environment, 2020, 705, 135882.	8.0	63
85	Effects of mercury on the activity and community composition of soil ammonia oxidizers. Environmental Science and Pollution Research, 2010, 17, 1237-1244.	5.3	62
86	Nitrification Is a Primary Driver of Nitrous Oxide Production in Laboratory Microcosms from Different Land-Use Soils. Frontiers in Microbiology, 2016, 7, 1373.	3.5	62
87	Diversity and potential biogeochemical impacts of viruses in bulk and rhizosphere soils. Environmental Microbiology, 2021, 23, 588-599.	3.8	62
88	Abundance and community structure of ammonia-oxidizingArchaeaandBacteriain response to fertilization and mowing in a temperate steppe in Inner Mongolia. FEMS Microbiology Ecology, 2014, 89, 67-79.	2.7	61
89	Responses of soil nitrous oxide production and abundances and composition of associated microbial communities to nitrogen and water amendment. Biology and Fertility of Soils, 2017, 53, 601-611.	4.3	61
90	Deterministic selection dominates microbial community assembly in termite mounds. Soil Biology and Biochemistry, 2021, 152, 108073.	8.8	60

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91	Niche differentiation of clade A comammox Nitrospira and canonical ammonia oxidizers in selected forest soils. Soil Biology and Biochemistry, 2020, 149, 107925.	8.8	59
92	Effects of 3,4-dimethylpyrazole phosphate (DMPP) on nitrification and the abundance and community composition of soil ammonia oxidizers in three land uses. Biology and Fertility of Soils, 2016, 52, 927-939.	4.3	56
93	Influence of rice straw amendment on mercury methylation and nitrification in paddy soils. Environmental Pollution, 2016, 209, 53-59.	7.5	56
94	Microbial functional attributes, rather than taxonomic attributes, drive top soil respiration, nitrification and denitrification processes. Science of the Total Environment, 2020, 734, 139479.	8.0	56
95	Diversity of herbaceous plants and bacterial communities regulates soil resistome across forest biomes. Environmental Microbiology, 2018, 20, 3186-3200.	3.8	55
96	Linking soil bacterial diversity to ecosystem multifunctionality using backward-elimination boosted trees analysis. Journal of Soils and Sediments, 2009, 9, 547-554.	3.0	54
97	Multiple factors drive the abundance and diversity of the diazotrophic community in typical farmland soils of China. FEMS Microbiology Ecology, 2019, 95, .	2.7	54
98	Salinity as a predominant factor modulating the distribution patterns of antibiotic resistance genes in ocean and river beach soils. Science of the Total Environment, 2019, 668, 193-203.	8.0	54
99	Manure application increases microbiome complexity in soil aggregate fractions: Results of an 18-year field experiment. Agriculture, Ecosystems and Environment, 2021, 307, 107249.	5.3	54
100	Large-scale patterns of soil antibiotic resistome in Chinese croplands. Science of the Total Environment, 2020, 712, 136418.	8.0	53
101	Fertilization alters protistan consumers and parasites in cropâ€associated microbiomes. Environmental Microbiology, 2021, 23, 2169-2183.	3.8	52
102	Ensuring planetary survival: the centrality of organic carbon in balancing the multifunctional nature of soils. Critical Reviews in Environmental Science and Technology, 2022, 52, 4308-4324.	12.8	52
103	Oxytetracycline and Ciprofloxacin Exposure Altered the Composition of Protistan Consumers in an Agricultural Soil. Environmental Science & Technology, 2020, 54, 9556-9563.	10.0	51
104	Potential of indigenous crop microbiomes for sustainable agriculture. Nature Food, 2021, 2, 233-240.	14.0	51
105	Coupling of soil prokaryotic diversity and plant diversity across latitudinal forest ecosystems. Scientific Reports, 2016, 6, 19561.	3.3	50
106	Frontiers in the microbial processes of ammonia oxidation in soils and sediments. Journal of Soils and Sediments, 2014, 14, 1023-1029.	3.0	49
107	Immediate effects of nitrogen, phosphorus, and potassium amendments on the methanotrophic activity and abundance in a Chinese paddy soil under short-term incubation experiment. Journal of Soils and Sediments, 2013, 13, 189-196.	3.0	48
108	Field-based evidence for consistent responses of bacterial communities to copper contamination in two contrasting agricultural soils. Frontiers in Microbiology, 2015, 6, 31.	3.5	47

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109	Plant-driven niche differentiation of ammonia-oxidizing bacteria and archaea in global drylands. ISME Journal, 2019, 13, 2727-2736.	9.8	47
110	The influence of soil age on ecosystem structure and function across biomes. Nature Communications, 2020, 11, 4721.	12.8	47
111	Succession of plant and soil microbial communities with restoration of abandoned land in the Loess Plateau, China. Journal of Soils and Sediments, 2013, 13, 760-769.	3.0	46
112	Differentiated Mechanisms of Biochar Mitigating Straw-Induced Greenhouse Gas Emissions in Two Contrasting Paddy Soils. Frontiers in Microbiology, 2018, 9, 2566.	3.5	46
113	Niche differentiation of comammox Nitrospira and canonical ammonia oxidizers in soil aggregate fractions following 27-year fertilizations. Agriculture, Ecosystems and Environment, 2020, 304, 107147.	5.3	46
114	Initial Copper Stress Strengthens the Resistance of Soil Microorganisms to a Subsequent Copper Stress. Microbial Ecology, 2014, 67, 931-941.	2.8	44
115	Effects of different agricultural wastes on the dissipation of PAHs and the PAH-degrading genes in a PAH-contaminated soil. Chemosphere, 2017, 172, 286-293.	8.2	44
116	Time-dependent shifts in populations and activity of bacterial and archaeal ammonia oxidizers in response to liming in acidic soils. Soil Biology and Biochemistry, 2017, 112, 77-89.	8.8	44
117	Antibiotic resistance in urban green spaces mirrors the pattern of industrial distribution. Environment International, 2019, 132, 105106.	10.0	42
118	Changes of the denitrifying communities in a multi-stage free water surface constructed wetland. Science of the Total Environment, 2019, 650, 1419-1425.	8.0	41
119	Species identity of biocrust-forming lichens drives the response of soil nitrogen cycle to altered precipitation frequency and nitrogen amendment. Soil Biology and Biochemistry, 2016, 96, 128-136.	8.8	40
120	Microbial nitrous oxide emissions in dryland ecosystems: mechanisms, microbiome and mitigation. Environmental Microbiology, 2017, 19, 4808-4828.	3.8	40
121	Sorption mechanism and distribution of cadmium by different microbial species. Journal of Environmental Management, 2019, 237, 552-559.	7.8	40
122	Ecological Drivers of Biogeographic Patterns of Soil Archaeal Community. PLoS ONE, 2013, 8, e63375.	2.5	39
123	Response of ammonia oxidizing microbes to the stresses of arsenic and copper in two acidic alfisols. Applied Soil Ecology, 2014, 77, 59-67.	4.3	39
124	Dryland forest management alters fungal community composition and decouples assembly of root- and soil-associated fungal communities. Soil Biology and Biochemistry, 2017, 109, 14-22.	8.8	39
125	Nitrogen Addition Decreases Dissimilatory Nitrate Reduction to Ammonium in Rice Paddies. Applied and Environmental Microbiology, 2018, 84, .	3.1	39
126	Distributions and environmental drivers of archaea and bacteria in paddy soils. Journal of Soils and Sediments, 2019, 19, 23-37.	3.0	39

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127	Growth of comammox Nitrospira is inhibited by nitrification inhibitors in agricultural soils. Journal of Soils and Sediments, 2020, 20, 621-628.	3.0	38
128	Quantitative analyses of the abundance and composition of ammoniaâ€oxidizing bacteria and ammoniaâ€oxidizing archaea of a Chinese upland red soil under longâ€term fertilization practices. Environmental Microbiology, 2007, 9, 3152-3152.	3.8	36
129	Response of bacterial pdo1, nah, and C12O genes to aged soil PAH pollution in a coke factory area. Environmental Science and Pollution Research, 2014, 21, 9754-9763.	5.3	36
130	Response of ammonia oxidizers and denitrifiers to repeated applications of a nitrification inhibitor and a urease inhibitor in two pasture soils. Journal of Soils and Sediments, 2017, 17, 974-984.	3.0	36
131	Rare earth oxide nanoparticles promote soil microbial antibiotic resistance by selectively enriching antibiotic resistance genes. Environmental Science: Nano, 2019, 6, 456-466.	4.3	36
132	Enhanced nitrogen retention by lignite during poultry litter composting. Journal of Cleaner Production, 2020, 277, 122422.	9.3	36
133	Effects of super absorbent polymers on soil microbial properties and Chinese cabbage (Brassica) Tj ETQq1 1 0.784314 rgBT /Overlock	3.0	34
134	Microbial functional traits in phyllosphere are more sensitive to anthropogenic disturbance than in soil. Environmental Pollution, 2020, 265, 114954.	7.5	34
135	Fates and Use Efficiency of Nitrogen Fertilizer in Maize Cropping Systems and Their Responses to Technologies and Management Practices: A Global Analysis on Field ¹⁵N Tracer Studies. Earth's Future, 2021, 9, e2020EF001514.	6.3	34
136	Distinct factors drive the diversity and composition of protistan consumers and phototrophs in natural soil ecosystems. Soil Biology and Biochemistry, 2021, 160, 108317.	8.8	34
137	Niche specialization of comammox Nitrospira in terrestrial ecosystems: Oligotrophic or copiotrophic?. Critical Reviews in Environmental Science and Technology, 2023, 53, 161-176.	12.8	34
138	Temporal dynamics of fungal communities in soybean rhizosphere. Journal of Soils and Sediments, 2017, 17, 491-498.	3.0	33
139	Bacterial composition and spatiotemporal variation in sediments of Jiaozhou Bay, China. Journal of Soils and Sediments, 2015, 15, 732-744.	3.0	32
140	Climatic factors have unexpectedly strong impacts on soil bacterial Î²-diversity in 12 forest ecosystems. Soil Biology and Biochemistry, 2020, 142, 107699.	8.8	32
141	Genetic and functional diversity of ubiquitous DNA viruses in selected Chinese agricultural soils. Scientific Reports, 2017, 7, 45142.	3.3	31
142	Soil aggregate size and long-term fertilization effects on the function and community of ammonia oxidizers. Geoderma, 2019, 338, 107-117.	5.1	31
143	Fate of antibiotic resistance genes during high-solid anaerobic co-digestion of pig manure with lignite. Bioresource Technology, 2020, 303, 122906.	9.6	30
144	Candidatus Brocadia and Candidatus Kuenenia predominated in anammox bacterial community in selected Chinese paddy soils. Journal of Soils and Sediments, 2015, 15, 1977-1986.	3.0	29

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145	Sorghum rhizosphere effects reduced soil bacterial diversity by recruiting specific bacterial species under low nitrogen stress. Science of the Total Environment, 2021, 770, 144742.	8.0	29
146	Host Species and Geography Differentiate Honeybee Gut Bacterial Communities by Changing the Relative Contribution of Community Assembly Processes. MBio, 2021, 12, e0075121.	4.1	29
147	Environmental Filtering Process Has More Important Roles than Dispersal Limitation in Shaping Large-Scale Prokaryotic Beta Diversity Patterns of Grassland Soils. Microbial Ecology, 2016, 72, 221-230.	2.8	28
148	Interactive effects of multiple climate change factors on ammonia oxidizers and denitrifiers in a temperate steppe. FEMS Microbiology Ecology, 2017, 93, .	2.7	28
149	The biogeography of fungal communities in paddy soils is mainly driven by geographic distance. Journal of Soils and Sediments, 2018, 18, 1795-1805.	3.0	28
150	Canonical ammonia oxidizers, rather than comammox Nitrospira, dominated autotrophic nitrification during the mineralization of organic substances in two paddy soils. Soil Biology and Biochemistry, 2021, 156, 108192.	8.8	28
151	Responses of soil microbial community to nitrogen fertilizer and precipitation regimes in a semi-arid steppe. Journal of Soils and Sediments, 2018, 18, 762-774.	3.0	27
152	Plant Diversity Enhances Soil Fungal Diversity and Microbial Resistance to Plant Invasion. Applied and Environmental Microbiology, 2021, 87, .	3.1	27
153	Agricultural land-use change and rotation system exert considerable influences on the soil antibiotic resistome in Lake Tai Basin. Science of the Total Environment, 2021, 771, 144848.	8.0	27
154	Ammonia-oxidizing bacteria play an important role in nitrification of acidic soils: A meta-analysis. Geoderma, 2021, 404, 115395.	5.1	27
155	Effects of repeated applications of urea with DMPP on ammonia oxidizers, denitrifiers, and non-targeted microbial communities of an agricultural soil in Queensland, Australia. Applied Soil Ecology, 2020, 147, 103392.	4.3	26
156	Aridity decreases soil protistan network complexity and stability. Soil Biology and Biochemistry, 2022, 166, 108575.	8.8	26
157	Shifts in the abundance and community structure of soil ammonia oxidizers in a wet sclerophyll forest under long-term prescribed burning. Science of the Total Environment, 2014, 470-471, 578-586.	8.0	25
158	Manure Application Did Not Enrich Antibiotic Resistance Genes in Root Endophytic Bacterial Microbiota of Cherry Radish Plants. Applied and Environmental Microbiology, 2020, 86, .	3.1	25
159	Niche specialization of comammox Nitrospira clade A in terrestrial ecosystems. Soil Biology and Biochemistry, 2021, 156, 108231.	8.8	25
160	Long-term nickel exposure altered the bacterial community composition but not diversity in two contrasting agricultural soils. Environmental Science and Pollution Research, 2015, 22, 10496-10505.	5.3	24
161	Ectomycorrhizal fungi inoculation alleviates simulated acid rain effects on soil ammonia oxidizers and denitrifiers in Masson pine forest. Environmental Microbiology, 2019, 21, 299-313.	3.8	24
162	Dynamics of sulfate reduction and sulfate-reducing prokaryotes in anaerobic paddy soil amended with rice straw. Biology and Fertility of Soils, 2010, 46, 283-291.	4.3	23

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163	Changes in soil nematode abundance and composition under elevated [CO2] and canopy warming in a rice paddy field. Plant and Soil, 2019, 445, 425-437.	3.7	23
164	Lime and ammonium carbonate fumigation coupled with bioâ€organic fertilizer application steered banana rhizosphere to assemble a unique microbiome against Panama disease. Microbial Biotechnology, 2019, 12, 515-527.	4.2	23
165	Host identity determines plant associated resistomes. Environmental Pollution, 2020, 258, 113709.	7.5	23
166	Impact of sulfate and iron oxide on bacterial community dynamics in paddy soil under alternate watering conditions. Journal of Hazardous Materials, 2021, 408, 124417.	12.4	23
167	Biotic and abiotic factors distinctly drive contrasting biogeographic patterns between phyllosphere and soil resistomes in natural ecosystems. ISME Communications, 2021, 1, .	4.2	23
168	The overlap of soil and vegetable microbes drives the transfer of antibiotic resistance genes from manure-amended soil to vegetables. Science of the Total Environment, 2022, 828, 154463.	8.0	23
169	Effects of the nitrification inhibitor acetylene on nitrous oxide emissions and ammonia-oxidizing microorganisms of different agricultural soils under laboratory incubation conditions. Applied Soil Ecology, 2017, 119, 80-90.	4.3	22
170	Effects of the nitrification inhibitor dicyandiamide (DCD) on N2Oemissions and the abundance of nitrifiers and denitrifiers in two contrasting agricultural soils. Journal of Soils and Sediments, 2017, 17, 1635-1643.	3.0	22
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