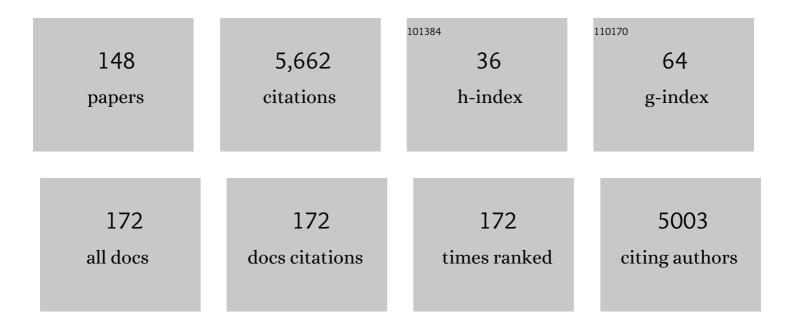
## Ji-Dong Gu

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9576870/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Microbiological deterioration and degradation of synthetic polymeric materials: recent research advances. International Biodeterioration and Biodegradation, 2003, 52, 69-91.	1.9	557
2	Bathyarchaeota: globally distributed metabolic generalists in anoxic environments. FEMS Microbiology Reviews, 2018, 42, 639-655.	3.9	206
3	Biodeterioration of concrete by the fungus Fusarium. International Biodeterioration and Biodegradation, 1998, 41, 101-109.	1.9	193
4	Microbial electrocatalysis: Redox mediators responsible for extracellular electron transfer. Biotechnology Advances, 2018, 36, 1815-1827.	6.0	183
5	Insights into the ecology, evolution, and metabolism of the widespread Woesearchaeotal lineages. Microbiome, 2018, 6, 102.	4.9	181
6	Microbial extracellular enzymes in biogeochemical cycling of ecosystems. Journal of Environmental Management, 2017, 197, 539-549.	3.8	170
7	Metagenomic and metatranscriptomic analyses reveal activity and hosts of antibiotic resistance genes in activated sludge. Environment International, 2019, 129, 208-220.	4.8	163
8	Microbial deterioration and sustainable conservation of stone monuments and buildings. Nature Sustainability, 2020, 3, 991-1004.	11.5	136
9	Complex microbial nitrogen-cycling networks in three distinct anammox-inoculated wastewater treatment systems. Water Research, 2020, 168, 115142.	5.3	109
10	More than a decade of experience of landfill leachate treatment with a full-scale anammox plant combining activated sludge and activated carbon biofilm. Chemosphere, 2017, 174, 117-126.	4.2	93
11	Genomic and transcriptomic insights into the ecology and metabolism of benthic archaeal cosmopolitan, Thermoprofundales (MBG-D archaea). ISME Journal, 2019, 13, 885-901.	4.4	92
12	Stratified Bacterial and Archaeal Community in Mangrove and Intertidal Wetland Mudflats Revealed by High Throughput 16S rRNA Gene Sequencing. Frontiers in Microbiology, 2017, 8, 2148.	1.5	91
13	Ecological responses, adaptation and mechanisms of mangrove wetland ecosystem to global climate change and anthropogenic activities. International Biodeterioration and Biodegradation, 2021, 162, 105248.	1.9	89
14	Comparative genomic inference suggests mixotrophic lifestyle for Thorarchaeota. ISME Journal, 2018, 12, 1021-1031.	4.4	86
15	Water is a critical factor in evaluating and assessing microbial colonization and destruction of Angkor sandstone monuments. International Biodeterioration and Biodegradation, 2018, 133, 9-16.	1.9	79
16	Changes in the biofilm microflora of limestone caused by atmospheric pollutants. International Biodeterioration and Biodegradation, 2000, 46, 299-303.	1.9	74
17	The community distribution of bacteria and fungi on ancient wall paintings of the Mogao Grottoes. Scientific Reports, 2015, 5, 7752.	1.6	70
18	Microbial Community Analysis of Fresh and Old Microbial Biofilms on Bayon Temple Sandstone of Angkor Thom, Cambodia. Microbial Ecology, 2010, 60, 105-115.	1.4	68

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19	Biochemical reactions and mechanisms involved in the biodeterioration of stone world cultural heritage under the tropical climate conditions. International Biodeterioration and Biodegradation, 2019, 143, 104723.	1.9	67
20	More wide occurrence and dominance of ammonia-oxidizing archaea than bacteria at three Angkor sandstone temples of Bayon, Phnom Krom and Wat Athvea in Cambodia. International Biodeterioration and Biodegradation, 2017, 117, 78-88.	1.9	66
21	Activity and Metabolic Versatility of Complete Ammonia Oxidizers in Full-Scale Wastewater Treatment Systems. MBio, 2020, 11, .	1.8	65
22	Future directions and challenges in biodeterioration research on historic materials and cultural properties. International Biodeterioration and Biodegradation, 2018, 129, 10-12.	1.9	63
23	Diversity and distribution of Archaea in global estuarine ecosystems. Science of the Total Environment, 2018, 637-638, 349-358.	3.9	62
24	Diverse Asgard archaea including the novel phylum Gerdarchaeota participate in organic matter degradation. Science China Life Sciences, 2020, 63, 886-897.	2.3	61
25	Oxidation of Elemental Sulfur by Fusarium solani Strain THIF01 Harboring Endobacterium Bradyrhizobium sp Microbial Ecology, 2010, 60, 96-104.	1.4	56
26	Alteration of extracellular enzyme activity and microbial abundance by biochar addition: Implication for carbon sequestration in subtropical mangrove sediment. Journal of Environmental Management, 2016, 182, 29-36.	3.8	53
27	Anaerobic Degradation of Paraffins by Thermophilic Actinobacteria under Methanogenic Conditions. Environmental Science & Technology, 2020, 54, 10610-10620.	4.6	53
28	Higher diversity and abundance of ammonia-oxidizing archaea than bacteria detected at the Bayon Temple of Angkor Thom in Cambodia. International Biodeterioration and Biodegradation, 2016, 115, 234-243.	1.9	52
29	Biodegradability of plastics: the issues, recent advances, and future perspectives. Environmental Science and Pollution Research, 2021, 28, 1278-1282.	2.7	49
30	Microbiological community of the Royal Palace in Angkor Thom and Beng Mealea of Cambodia by Illumina sequencing based on 16S rRNA gene. International Biodeterioration and Biodegradation, 2018, 134, 127-135.	1.9	47
31	Occurrence of Aspergillus allahabadii on sandstone at Bayon temple, Angkor Thom, Cambodia. International Biodeterioration and Biodegradation, 2013, 76, 112-117.	1.9	44
32	Occurrence of anammox bacteria in a traditional full-scale wastewater treatment plant and successful inoculation for new establishment. International Biodeterioration and Biodegradation, 2017, 120, 224-231.	1.9	44
33	Faunal Burrows Alter the Diversity, Abundance, and Structure of AOA, AOB, Anammox and n-Damo Communities in Coastal Mangrove Sediments. Microbial Ecology, 2017, 74, 140-156.	1.4	42
34	Lithoautotrophical oxidation of elemental sulfur by fungi including Fusarium solani isolated from sandstone Angkor temples. International Biodeterioration and Biodegradation, 2018, 126, 95-102.	1.9	42
35	Activities and metabolic versatility of distinct anammox bacteria in a full-scale wastewater treatment system. Water Research, 2021, 206, 117763.	5.3	42
36	Specific and effective detection of anammox bacteria using PCR primers targeting the 16S rRNA gene and functional genes. Science of the Total Environment, 2020, 734, 139387.	3.9	41

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37	Direct microbial transformation of carbon dioxide to value-added chemicals: A comprehensive analysis and application potentials. Bioresource Technology, 2019, 288, 121401.	4.8	40
38	Improved anaerobic co-digestion of food waste and domestic wastewater by copper supplementation – Microbial community change and enhanced effluent quality. Science of the Total Environment, 2019, 670, 337-344.	3.9	40
39	Survival strategies of ammonia-oxidizing archaea (AOA) in a full-scale WWTP treating mixed landfill leachate containing copper ions and operating at low-intensity of aeration. Water Research, 2021, 191, 116798.	5.3	39
40	Abundance and niche specificity of different types of complete ammonia oxidizers (comammox) in salt marshes covered by different plants. Science of the Total Environment, 2021, 768, 144993.	3.9	39
41	Biofilm control by interfering with c-di-GMP metabolism and signaling. Biotechnology Advances, 2022, 56, 107915.	6.0	39
42	Existence of Novel Phylotypes of Nitrite-Dependent Anaerobic Methane-Oxidizing Bacteria in Surface and Subsurface Sediments of the South China Sea. Geomicrobiology Journal, 2015, 32, 1-10.	1.0	38
43	Impact of nitrogen pollution/deposition on extracellular enzyme activity, microbial abundance and carbon storage in coastal mangrove sediment. Chemosphere, 2017, 177, 275-283.	4.2	36
44	Diazotrophic microbial community and abundance in acidic subtropical natural and re-vegetated forest soils revealed by high-throughput sequencing of nifH gene. Applied Microbiology and Biotechnology, 2019, 103, 995-1005.	1.7	35
45	Community structures of bacteria and archaea associated with the biodeterioration of sandstone sculptures at the Beishiku Temple. International Biodeterioration and Biodegradation, 2021, 164, 105290.	1.9	34
46	Biotransformation of lincomycin and fluoroquinolone antibiotics by the ammonia oxidizers AOA, AOB and comammox: A comparison of removal, pathways, and mechanisms. Water Research, 2021, 196, 117003.	5.3	33
47	Differences of Microbial Community on the wall paintings preserved in situ and ex situ of the Tiantishan Grottoes, China. International Biodeterioration and Biodegradation, 2018, 132, 102-113.	1.9	32
48	Ecological distribution and potential roles of Woesearchaeota in anaerobic biogeochemical cycling unveiled by genomic analysis. Computational and Structural Biotechnology Journal, 2021, 19, 794-800.	1.9	32
49	Genomic and transcriptomic evidence of light-sensing, porphyrin biosynthesis, Calvin-Benson-Bassham cycle, and urea production in Bathyarchaeota. Microbiome, 2020, 8, 43.	4.9	31
50	Further Analysis of Anammox Bacterial Community Structures Along an Anthropogenic Nitrogen-Input Gradient from the Riparian Sediments of the Pearl River Delta to the Deep-Ocean Sediments of the South China Sea. Geomicrobiology Journal, 2015, 32, 789-798.	1.0	30
51	Co-occurrence of nitrite-dependent anaerobic ammonium and methane oxidation processes in subtropical acidic forest soils. Applied Microbiology and Biotechnology, 2016, 100, 7727-7739.	1.7	30
52	Mining, pollution and site remediation. International Biodeterioration and Biodegradation, 2018, 128, 1-2.	1.9	30
53	Dredging alleviates cyanobacterial blooms by weakening diversity maintenance of bacterioplankton community. Water Research, 2021, 202, 117449.	5.3	29
54	Lower Abundance of Ammonia-Oxidizing Archaea Than Ammonia-Oxidizing Bacteria Detected in the Subsurface Sediments of the Northern South China Sea. Geomicrobiology Journal, 2012, 29, 332-339.	1.0	27

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55	Analysis of methane-producing and metabolizing archaeal and bacterial communities in sediments of the northern South China Sea and coastal Mai Po Nature Reserve revealed by PCR amplification of mcrA and pmoA genes. Frontiers in Microbiology, 2014, 5, 789.	1.5	27
56	Exploring possible associations of the intestine bacterial microbiome with the pre-weaned weight gaining performance of piglets in intensive pig production. Scientific Reports, 2019, 9, 15534.	1.6	27
57	Microbiome characteristics and the key biochemical reactions identified on stone world cultural heritage under different climate conditions. Journal of Environmental Management, 2022, 302, 114041.	3.8	27
58	A more accurate definition of water characteristics in stone materials for an improved understanding and effective protection of cultural heritage from biodeterioration. International Biodeterioration and Biodegradation, 2022, 166, 105338.	1.9	27
59	The active microbes and biochemical processes contributing to deterioration of Angkor sandstone monuments under the tropical climate in Cambodia – A review. Journal of Cultural Heritage, 2021, 47, 218-226.	1.5	26
60	The microbial community characteristics of ancient painted sculptures in Maijishan Grottoes, China. PLoS ONE, 2017, 12, e0179718.	1.1	25
61	Current advances in molecular methods for detection of nitrite-dependent anaerobic methane oxidizing bacteria in natural environments. Applied Microbiology and Biotechnology, 2016, 100, 9845-9860.	1.7	24
62	A comparison of denitrifying bacterial community structures and abundance in acidic soils between natural forest and re-vegetated forest of Nanling Nature Reserve in southern China. Journal of Environmental Management, 2017, 198, 41-49.	3.8	24
63	A More Comprehensive Community ofÂAmmonia-Oxidizing Archaea (AOA) Revealed by Genomic DNA and RNA Analyses of amoA Gene in Subtropical Acidic Forest Soils. Microbial Ecology, 2017, 74, 910-922.	1.4	24
64	An internal recycling mechanism between ammonia/ammonium and nitrate driven by ammonia-oxidizing archaea and bacteria (AOA, AOB, and Comammox) and DNRA on Angkor sandstone monuments. International Biodeterioration and Biodegradation, 2021, 165, 105328.	1.9	24
65	Modification of cyanobacterial bloom-derived biomass using potassium permanganate enhanced the removal of microcystins and adsorption capacity toward cadmium (II). Journal of Hazardous Materials, 2014, 272, 83-88.	6.5	23
66	Salinity-driven heterogeneity toward anammox distribution and growth kinetics. Applied Microbiology and Biotechnology, 2019, 103, 1953-1960.	1.7	23
67	Manganese enhances the immobilization of trace cadmium from irrigation water in biological soil crust. Ecotoxicology and Environmental Safety, 2019, 168, 369-377.	2.9	23
68	Establishing practical strategies to run high loading corn stover anaerobic digestion: Methane production performance and microbial responses. Bioresource Technology, 2020, 310, 123364.	4.8	23
69	Subgroup level differences of physiological activities in marine Lokiarchaeota. ISME Journal, 2021, 15, 848-861.	4.4	23
70	The dynamics of phosphorus fractions and the factors driving phosphorus cycle in Zoige Plateau peatland soil. Chemosphere, 2021, 278, 130501.	4.2	23
71	Biodegradability of plastics: the pitfalls. Applied Environmental Biotechnology, 2017, 2, 59-61.	1.0	23
72	Abundance of ammonia-oxidizing bacteria and archaea under different ventilation strategies during cattle manure composting. Journal of Environmental Management, 2018, 212, 375-383.	3.8	22

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73	Cu-bearing high-entropy alloys with excellent antiviral properties. Journal of Materials Science and Technology, 2021, 84, 59-64.	5.6	22
74	Salinity gradients shape the nitrifier community composition in Nanliu River Estuary sediments and the ecophysiology of comammox Nitrospira inopinata. Science of the Total Environment, 2021, 795, 148768.	3.9	22
75	The diversity and distribution of anammox bacteria in the marine aquaculture zones. Applied Microbiology and Biotechnology, 2016, 100, 8943-8953.	1.7	21
76	Assessment of molecular detection of anaerobic ammonium-oxidizing (anammox) bacteria in different environmental samples using PCR primers based on 16S rRNA and functional genes. Applied Microbiology and Biotechnology, 2017, 101, 7689-7702.	1.7	21
77	Practical applications of PCR primers in detection of anammox bacteria effectively from different types of samples. Applied Microbiology and Biotechnology, 2018, 102, 5859-5871.	1.7	21
78	Microbiome and nitrate removal processes by microorganisms on the ancient Preah Vihear temple of Cambodia revealed by metagenomics and N-15 isotope analyses. Applied Microbiology and Biotechnology, 2020, 104, 9823-9837.	1.7	21
79	Spatial and temporal distributions of microbial diversity under natural conditions on the sandstone stelae of the Beishiku Temple in China. International Biodeterioration and Biodegradation, 2021, 163, 105279.	1.9	21
80	A mixed blessing of viruses in wastewater treatment plants. Water Research, 2022, 215, 118237.	5.3	21
81	Successive transitory distribution of Thaumarchaeota and partitioned distribution of Bathyarchaeota from the Pearl River estuary to the northern South China Sea. Applied Microbiology and Biotechnology, 2018, 102, 8035-8048.	1.7	20
82	Metal distribution and biological diversity of crusts in paddy fields polluted with different levels of cadmium. Ecotoxicology and Environmental Safety, 2019, 184, 109620.	2.9	20
83	A Review on Sampling Techniques and Analytical Methods for Microbiota of Cultural Properties and Historical Architecture. Applied Sciences (Switzerland), 2020, 10, 8099.	1.3	20
84	Two or three domains: a new view of tree of life in the genomics era. Applied Microbiology and Biotechnology, 2018, 102, 3049-3058.	1.7	19
85	Optogenetic Modulation of a Catalytic Biofilm for the Biotransformation of Indole into Tryptophan. ChemSusChem, 2019, 12, 5142-5148.	3.6	19
86	Influence of critical factors on nitrogen removal contribution by anammox and denitrification in an anammox-inoculated wastewater treatment system. Journal of Water Process Engineering, 2021, 40, 101868.	2.6	19
87	Innovative approaches for the processes involved in microbial biodeterioration of cultural heritage materials. Current Opinion in Biotechnology, 2022, 75, 102716.	3.3	19
88	Community assembly, potential functions and interactions between fungi and microalgae associated with biodeterioration of sandstone at the Beishiku Temple in Northwest China. Science of the Total Environment, 2022, 835, 155372.	3.9	19
89	New PCR primers targeting hydrazine synthase and cytochrome c biogenesis proteins in anammox bacteria. Applied Microbiology and Biotechnology, 2017, 101, 1267-1287.	1.7	18
90	Biosorption of diethyl phthalate ester by living and nonliving Burkholderia cepacia and the role of its cell surface components. Chemosphere, 2017, 178, 187-196.	4.2	18

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91	Assessing ecological health of mangrove ecosystems along South China Coast by the pressure–state–response (PSR) model. Ecotoxicology, 2021, 30, 622-631.	1.1	18
92	Effects of bloom-forming cyanobacterial extracellular polymeric substances on the adsorption of cadmium onto kaolinite: behaviors and possible mechanisms. SpringerPlus, 2016, 5, 542.	1.2	17
93	Two identical copies of the hydrazine synthase gene clusters found in the genomes of anammox bacteria. International Biodeterioration and Biodegradation, 2018, 132, 236-240.	1.9	17
94	Linkages between anammox and denitrifying bacterial communities and nitrogen loss rates in highâ€elevation rivers. Limnology and Oceanography, 2021, 66, 765-778.	1.6	17
95	Bacterial and fungal communities in the sandstone biofilms of two famous Buddhist grottoes in China. International Biodeterioration and Biodegradation, 2021, 163, 105267.	1.9	17
96	Proteomic analysis of hepatic tissue of ciguatoxin (CTX) contaminated coral reef fish Cephalopholis argus and moray eel Gymnothorax undulatus. Harmful Algae, 2012, 13, 65-71.	2.2	16
97	Differential distribution patterns of ammonia-oxidizing archaea and bacteria in acidic soils of Nanling National Nature Reserve forests in subtropical China. Antonie Van Leeuwenhoek, 2016, 109, 237-251.	0.7	16
98	Nutrient limitation status in a subtropical mangrove ecosystem revealed by analysis of enzymatic stoichiometry and microbial abundance for sediment carbon cycling. International Biodeterioration and Biodegradation, 2018, 128, 3-10.	1.9	16
99	Redirecting marine antibiofouling innovations from sustainable horizons. Trends in Ecology and Evolution, 2022, 37, 469-472.	4.2	15
100	Simulation of in situ oil reservoir conditions in a laboratory bioreactor testing for methanogenic conversion of crude oil and analysis of the microbial community. International Biodeterioration and Biodegradation, 2019, 136, 24-33.	1.9	14
101	Seasonal Variability of Extracellular Enzymes Involved in Carbon Mineralization in Sediment of a Subtropical Mangrove Wetland. Geomicrobiology Journal, 2015, 32, 68-76.	1.0	13
102	Influence of mangrove roots on microbial abundance and ecoenzyme activity in sediments of a subtropical coastal mangrove ecosystem. International Biodeterioration and Biodegradation, 2018, 132, 10-17.	1.9	13
103	Cyanobacterial bloom mitigation by sanguinarine and its effects on aquatic microbial community structure. Environmental Pollution, 2019, 253, 497-506.	3.7	13
104	Nano-TiO2 enhances the adsorption of Cd(II) on biological soil crusts under mildly acidic conditions. Journal of Contaminant Hydrology, 2020, 229, 103583.	1.6	13
105	Relationship of proteomic variation and toxin synthesis in the dinoflagellate Alexandrium tamarense Cl01 under phosphorus and inorganic nitrogen limitation. Ecotoxicology, 2015, 24, 1744-1753.	1.1	12
106	Microbial reduction of CO2 from injected NaH13CO3 with degradation of n-hexadecane in the enrichment culture derived from a petroleum reservoir. International Biodeterioration and Biodegradation, 2018, 127, 192-200.	1.9	12
107	Stochastic assembly process dominates bacterial succession during a long-term microbial enhanced oil recovery. Science of the Total Environment, 2021, 790, 148203.	3.9	12
108	Coastal and marine pollution and ecotoxicology. Ecotoxicology, 2015, 24, 1407-1410.	1.1	11

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109	Abundance and Diversity of Aerobic/Anaerobic Ammonia/Ammonium-Oxidizing Microorganisms in an Ammonium-Rich Aquitard in the Pearl River Delta of South China. Microbial Ecology, 2018, 76, 81-91.	1.4	11
110	Assessment of Five Electron‣huttling Molecules in the Extracellular Electron Transfer of Electromethanogenesis by using <i>Methanosarcina barkeri</i> . ChemElectroChem, 2020, 7, 3783-3789.	1.7	11
111	Highly efficient removal of phosphorus from agricultural runoff by a new akadama clay barrier-vegetated drainage ditch system (VDD) and its mechanism. Journal of Environmental Management, 2021, 290, 112575.	3.8	11
112	Dominance of ammonia-oxidizing archaea community induced by land use change from Masson pine to eucalypt plantation in subtropical China. Applied Microbiology and Biotechnology, 2016, 100, 6859-6869.	1.7	10
113	Seasonal and spatial variations in diversity and abundance of bacterial laccase-like genes in sediments of a subtropical mangrove ecosystem. International Biodeterioration and Biodegradation, 2016, 114, 260-267.	1.9	10
114	Realization of biodeterioration to cultural heritage protection in China. International Biodeterioration and Biodegradation, 2017, 117, 128-130.	1.9	10
115	A xylan-degrading thermophilic and obligate anaerobe Xylanivirga thermophila gen. nov., sp. nov., isolated from an anammox dominant wastewater treatment plant, and proposal of Xylanivirgaceae fam. nov Anaerobe, 2020, 61, 102075.	1.0	10
116	Diversity, Abundance, and Distribution of Wood-Decay Fungi in Major Parks of Hong Kong. Forests, 2020, 11, 1030.	0.9	10
117	New evidence for a hydroxylation pathway for anaerobic alkane degradation supported by analyses of functional genes and signature metabolites in oil reservoirs. AMB Express, 2021, 11, 18.	1.4	10
118	Characterization of Aerobic Bacteria Involved in Degrading Polyethylene Glycol (PEG)-3400 Obtained by Plating and Enrichment Culture Techniques. Journal of Polymers and the Environment, 2007, 15, 57-65.	2.4	9
119	A global analysis on the distribution pattern of the bacteria coupling simultaneous methane oxidation to nitrite reduction. International Biodeterioration and Biodegradation, 2018, 129, 123-132.	1.9	9
120	Molecular Existence and Diversity of Nitrite-Dependent Anaerobic Methane Oxidizing (n-Damo) Bacteria in the Lakes of Badain of the Gobi Desert. Geomicrobiology Journal, 2019, 36, 522-532.	1.0	9
121	Diversity and spatial–temporal distribution of airborne fungi at the world culture heritage site Maijishan Grottoes in China. Aerobiologia, 2021, 37, 681-694.	0.7	9
122	Influence of Macrofaunal Burrows on Extracellular Enzyme Activity and Microbial Abundance in Subtropical Mangrove Sediment. Microbial Ecology, 2018, 76, 92-101.	1.4	7
123	Nitrification mainly driven by ammonia-oxidizing bacteria and nitrite-oxidizing bacteria in an an an anammox-inoculated wastewater treatment system. AMB Express, 2021, 11, 158.	1.4	7
124	The environmental factors used in correlation analysis with microbial community of environmental and cultural heritage samples. International Biodeterioration and Biodegradation, 2022, 173, 105460.	1.9	7
125	Distribution, diversity and abundance of bacterial laccase-like genes in different particle size fractions of sediments in a subtropical mangrove ecosystem. Ecotoxicology, 2015, 24, 1508-1516.	1.1	6
126	Effects of reforestation on ammonia-oxidizing microbial community composition and abundance in subtropical acidic forest soils. Applied Microbiology and Biotechnology, 2018, 102, 5309-5322.	1.7	6

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127	Functional dominance and community compositions of ammonia-oxidizing archaea in extremely acidic soils of natural forests. Applied Microbiology and Biotechnology, 2019, 103, 4229-4240.	1.7	6
128	Targeted assemblies of <i>cas1</i> suggest CRISPR-Cas's response to soil warming. ISME Journal, 2020, 14, 1651-1662.	4.4	6
129	High-throughput sequencing reveals the main drivers of niche-differentiation of bacterial community in the surface sediments of the northern South China sea. Marine Environmental Research, 2022, 178, 105641.	1.1	6
130	Archaeal Communities of South China Mangroves and Their Potential Roles in the Nitrogen Cycle. Geomicrobiology Journal, 2022, 39, 697-704.	1.0	5
131	Simultaneous detection of transcribed functional assA gene and the corresponding metabolites of linear alkanes (C4, C5, and C7) in production water of a low-temperature oil reservoir. Science of the Total Environment, 2020, 746, 141290.	3.9	4
132	Dominant and Active Methanogens in the Production Waters From a High-Temperature Petroleum Reservoir by DNA- and RNA-Based Analysis. Geomicrobiology Journal, 2021, 38, 191-198.	1.0	4
133	An Fe(II)-oxidizing consortium from Wudalianchi volcano spring in Northeast China for bioleaching of Cu and Ni from printed circuit boards (PCBs) with the dominance of Acidithiobacillus spp International Biodeterioration and Biodegradation, 2022, 167, 105355.	1.9	4
134	Biogeographic pattern of the nirS gene-targeted anammox bacterial community and composition in the northern South China Sea and a coastal Mai Po mangrove wetland. Applied Microbiology and Biotechnology, 2020, 104, 3167-3181.	1.7	3
135	Microbiota and Biochemical Processes Involved in Biodeterioration of Cultural Heritage and Protection. , 2021, , 37-58.		3
136	Discovery of the nonâ€cosmopolitan lineages in <i>Candidatus</i> Thermoprofundales. Environmental Microbiology, 2022, 24, 3063-3080.	1.8	3
137	Diversity, abundance, and distribution of anammox bacteria in shipping channel sediment of Hong Kong by analysis of DNA and RNA. Ecotoxicology, 2021, 30, 1705-1718.	1.1	2
138	Distribution of ammonia-oxidizing archaea and bacteria along an engineered coastal ecosystem in subtropical China. Ecotoxicology, 2021, 30, 1769-1779.	1.1	2
139	Shu-Pei Cheng: A life-long pursuit for Environmental Science and Pollution Control. Ecotoxicology, 2021, 30, 1284-1286.	1.1	1
140	To remember a passionate environmentalist. Ecotoxicology, 2021, 30, 1287-1289.	1.1	1
141	A brief introduction on the life of Shu-Pei Cheng. Ecotoxicology, 2021, 30, 1281-1283.	1.1	1
142	An international workshop on Conservation of Cultural Heritage held at the City University of Macau, China. International Biodeterioration and Biodegradation, 2021, 164, 105297.	1.9	1
143	Aerobic degradation and metabolite identification of the N-heterocyclic indole by the Pseudomonas putida strain mpky-1 isolated from subtropical mangrove sediment. Applied Environmental Biotechnology, 2017, 2, 1-10.	1.0	1
144	Microbial biomass C and N dynamics, and 15N incorporation into microbial biomass under faba bean, canola, barley, and summer fal- low in a Gray Luvisol. Applied Environmental Biotechnology, 2017, 2, 47-58.	1.0	1

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145	Bacterial and Archaeal Community Distribution in Oilfield Water Re-injection Facilities and the Influences from Microorganisms in Injected Water. Microbial Ecology, 2021, , 1.	1.4	1
146	Perspectives on Microbial Electron Transfer Networks for Environmental Biotechnology. Frontiers in Microbiology, 2022, 13, 845796.	1.5	1
147	Preface. Ecotoxicology, 2021, 30, 1279-1280.	1.1	Ο
148	Biodegradability of chemically synthesized syndiotactic poly(β-[R]- hydroxybutyrate) in soil of Northeast China. Applied Environmental Biotechnology, 2017, 2, 43-46.	1.0	0