

Yong-Guan Zhu

List of Publications by Year in descending order

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Version: 2024-02-01

657
papers

63,892
citations

576

129
h-index

1801

217
g-index

666
all docs

666
docs citations

666
times ranked

42112
citing authors

#	ARTICLE	IF	CITATIONS
1	Tire wear particles: An emerging threat to soil health. <i>Critical Reviews in Environmental Science and Technology</i> , 2023, 53, 239-257.	6.6	37
2	Soil metaphenomics: a step forward in metagenomics. <i>Archives of Agronomy and Soil Science</i> , 2022, 68, 1645-1663.	1.3	5
3	The enigma of environmental organoarsenicals: Insights and implications. <i>Critical Reviews in Environmental Science and Technology</i> , 2022, 52, 3835-3862.	6.6	20
4	Soil plastispheres as hotspots of antibiotic resistance genes and potential pathogens. <i>ISME Journal</i> , 2022, 16, 521-532.	4.4	148
5	Profiling the antibiotic resistome in soils between pristine and human-affected sites on the Tibetan Plateau. <i>Journal of Environmental Sciences</i> , 2022, 111, 442-451.	3.2	16
6	The chemical-microbial release and transformation of arsenic induced by citric acid in paddy soil. <i>Journal of Hazardous Materials</i> , 2022, 421, 126731.	6.5	14
7	Metabolic responses of indigenous bacteria in chicken faeces and maggots to multiple antibiotics via heavy water labeled single-cell Raman spectroscopy. <i>Journal of Environmental Sciences</i> , 2022, 113, 394-402.	3.2	5
8	Variations of earthworm gut bacterial community composition and metabolic functions in coastal upland soil along a 700-year reclamation chronosequence. <i>Science of the Total Environment</i> , 2022, 804, 149994.	3.9	27
9	Fluoroquinolone antibiotics disturb the defense system, gut microbiome, and antibiotic resistance genes of <i>Enchytraeus crypticus</i> . <i>Journal of Hazardous Materials</i> , 2022, 424, 127509.	6.5	24
10	Similar heterotrophic communities but distinct interactions supported by red and green snow algae in the Antarctic Peninsula. <i>New Phytologist</i> , 2022, 233, 1358-1368.	3.5	7
11	Distribution, transfer, ecological and human health risks of antibiotics in bay ecosystems. <i>Environment International</i> , 2022, 158, 106949.	4.8	24
12	Host age increased conjugal plasmid transfer in gut microbiota of the soil invertebrate <i>Caenorhabditis elegans</i> . <i>Journal of Hazardous Materials</i> , 2022, 424, 127525.	6.5	6
13	Long-term combined application of chemical fertilizers and organic manure shapes the gut microbial diversity and functional community structures of earthworms. <i>Applied Soil Ecology</i> , 2022, 170, 104250.	2.1	10
14	Landscape of genes in hospital wastewater breaking through the defense line of last-resort antibiotics. <i>Water Research</i> , 2022, 209, 117907.	5.3	13
15	Discarded masks as hotspots of antibiotic resistance genes during COVID-19 pandemic. <i>Journal of Hazardous Materials</i> , 2022, 425, 127774.	6.5	22
16	Powering biological nitrogen removal from the environment by geobatteries. <i>Trends in Biotechnology</i> , 2022, 40, 377-380.	4.9	10
17	Microbial communities on biodegradable plastics under different fertilization practices in farmland soil microcosms. <i>Science of the Total Environment</i> , 2022, 809, 152184.	3.9	22
18	Antibiotic resistance genes and antibiotic sensitivity in bacterial aerosols and their comparisons with known respiratory pathogens. <i>Journal of Aerosol Science</i> , 2022, 161, 105931.	1.8	11

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19	How to build Urbanome, the genome of the city?. <i>Science of the Total Environment</i> , 2022, 810, 152310.	3.9	2
20	Removal of potentially toxic elements from contaminated soil and water using bone char compared to plant- and bone-derived biochars: A review. <i>Journal of Hazardous Materials</i> , 2022, 427, 128131.	6.5	31
21	Organochlorine contamination enriches virus-encoded metabolism and pesticide degradation associated auxiliary genes in soil microbiomes. <i>ISME Journal</i> , 2022, 16, 1397-1408.	4.4	45
22	Biosafety of human environments can be supported by effective use of renewable biomass. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	3.3	5
23	Impacts of global change on the phyllosphere microbiome. <i>New Phytologist</i> , 2022, 234, 1977-1986.	3.5	75
24	Identification of the rhizosphere microbes that actively consume plant-derived carbon. <i>Soil Biology and Biochemistry</i> , 2022, 166, 108577.	4.2	14
25	Nanopore sequencing analysis of integron gene cassettes in sewages and soils. <i>Science of the Total Environment</i> , 2022, 817, 152766.	3.9	9
26	The ecological clusters of soil organisms drive the ecosystem multifunctionality under long-term fertilization. <i>Environment International</i> , 2022, 161, 107133.	4.8	53
27	å™→è€ä½“å¼“ç”ÿæ€ç—æ³•äŽš€ä½“å€—å¥å²¼¼šçŽ“çŠ¶ã€€æ’æ~äŽæœ°é‡. <i>Scientia Sinica Vitae</i> , 2022, . . 0.1		0
28	Viral diversity and potential environmental risk in microplastic at watershed scale: Evidence from metagenomic analysis of plastisphere. <i>Environment International</i> , 2022, 161, 107146.	4.8	23
29	Cross-biome antibiotic resistance decays after millions of years of soil development. <i>ISME Journal</i> , 2022, 16, 1864-1867.	4.4	8
30	Calling for comprehensive explorations between soil invertebrates and arbuscular mycorrhizas. <i>Trends in Plant Science</i> , 2022, 27, 793-801.	4.3	10
31	Abundance cannot represent antibiotic resistance risk. <i>Soil Ecology Letters</i> , 2022, 4, 291-292.	2.4	5
32	Soil inorganic carbon sequestration through alkalinity regeneration using biologically induced weathering of rock powder and biochar. <i>Soil Ecology Letters</i> , 2022, 4, 293-306.	2.4	9
33	Patterns and drivers of the degradability of dissolved organic matter in dryland soils on the Tibetan Plateau. <i>Journal of Applied Ecology</i> , 2022, 59, 884-894.	1.9	5
34	Widespread of Potential Pathogen-Derived Extracellular Vesicles Carrying Antibiotic Resistance Genes in Indoor Dust. <i>Environmental Science & Technology</i> , 2022, 56, 5653-5663.	4.6	12
35	Influences of arsenate and/or phosphate adsorption to ferrihydrite on iron-reducing and arsenic-reducing microbial communities in paddy soil revealed by rRNA-13C-acetate probing. <i>Soil Biology and Biochemistry</i> , 2022, 169, 108679.	4.2	5
36	Continentalâ€scale plant invasions reshuffle the soil microbiome of blue carbon ecosystems. <i>Global Change Biology</i> , 2022, 28, 4423-4438.	4.2	14

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37	Speciation Evolution of Phosphorus and Sulfur Derived from Sewage Sludge Biochar in Soil: Ageing Effects. <i>Environmental Science & Technology</i> , 2022, 56, 6639-6646.	4.6	13
38	Effects of Trophic Level and Land Use on the Variation of Animal Antibiotic Resistome in the Soil Food Web. <i>Environmental Science & Technology</i> , 2022, 56, 14937-14947.	4.6	19
39	How different nitrogen fertilizers affect arsenic mobility in paddy soil after straw incorporation?. <i>Journal of Hazardous Materials</i> , 2022, 436, 129135.	6.5	10
40	Embedded Health Risk from Arsenic in Globally Traded Rice. <i>Environmental Science & Technology</i> , 2022, 56, 6415-6425.	4.6	10
41	Unveiling the role of dissolved organic matter on phosphorus sorption and availability in a 5-year manure amended paddy soil. <i>Science of the Total Environment</i> , 2022, 838, 155892.	3.9	8
42	Organic fertilizer potentiates the transfer of typical antibiotic resistance gene among special bacterial species. <i>Journal of Hazardous Materials</i> , 2022, 435, 128985.	6.5	15
43	Globally distributed mining-impacted environments are underexplored hotspots of multidrug resistance genes. <i>ISME Journal</i> , 2022, 16, 2099-2113.	4.4	35
44	Root stoichiometry explains wheat endophytes and their link with crop production after four decades of fertilization. <i>Science of the Total Environment</i> , 2022, 846, 157407.	3.9	4
45	Estuarine plastisphere as an overlooked source of N ₂ O production. <i>Nature Communications</i> , 2022, 13, .	5.8	63
46	Sustainable removal of soil arsenic by naturally-formed iron oxides on plastic tubes. <i>Journal of Hazardous Materials</i> , 2022, 439, 129626.	6.5	3
47	A critical review of microplastic pollution in urban freshwater environments and legislative progress in China: Recommendations and insights. <i>Critical Reviews in Environmental Science and Technology</i> , 2021, 51, 2637-2680.	6.6	34
48	The co-evolution of life and organics on earth: Expansions of energy harnessing. <i>Critical Reviews in Environmental Science and Technology</i> , 2021, 51, 603-625.	6.6	2
49	Antibiotic resistome in the livestock and aquaculture industries: Status and solutions. <i>Critical Reviews in Environmental Science and Technology</i> , 2021, 51, 2159-2196.	6.6	109
50	Earthworm gut: An overlooked niche for anaerobic ammonium oxidation in agricultural soil. <i>Science of the Total Environment</i> , 2021, 752, 141874.	3.9	6
51	Co-selection of antibiotic resistance genes, and mobile genetic elements in the presence of heavy metals in poultry farm environments. <i>Science of the Total Environment</i> , 2021, 755, 142702.	3.9	122
52	Air pollution could drive global dissemination of antibiotic resistance genes. <i>ISME Journal</i> , 2021, 15, 270-281.	4.4	95
53	Rare taxa maintain the stability of crop microbiomes and ecosystem functions. <i>Environmental Microbiology</i> , 2021, 23, 1907-1924.	1.8	132
54	Biodiversity of key-stone phylotypes determines crop production in a 4-decade fertilization experiment. <i>ISME Journal</i> , 2021, 15, 550-561.	4.4	208

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55	Lessons learned from COVID-19 on potentially pathogenic soil microorganisms. <i>Soil Ecology Letters</i> , 2021, 3, 1-5.	2.4	18
56	Deterministic selection dominates microbial community assembly in termite mounds. <i>Soil Biology and Biochemistry</i> , 2021, 152, 108073.	4.2	60
57	Fates of Antibiotic Resistance Genes in the Gut Microbiome from Different Soil Fauna under Long-Term Fertilization. <i>Environmental Science & Technology</i> , 2021, 55, 423-432.	4.6	26
58	Metagenomic and ^{14}C tracing evidence for autotrophic microbial CO_2 fixation in paddy soils. <i>Environmental Microbiology</i> , 2021, 23, 924-933.	1.8	13
59	Host selection shapes crop microbiome assembly and network complexity. <i>New Phytologist</i> , 2021, 229, 1091-1104.	3.5	349
60	Evaluation of Microbe-Driven Soil Organic Matter Quantity and Quality by Thermodynamic Theory. <i>MBio</i> , 2021, 12, .	1.8	7
61	Herbicide Selection Promotes Antibiotic Resistance in Soil Microbiomes. <i>Molecular Biology and Evolution</i> , 2021, 38, 2337-2350.	3.5	68
62	Long-Term Fertilization Shapes the Putative Electrotrophic Microbial Community in Paddy Soils Revealed by Microbial Electrosynthesis Systems. <i>Environmental Science & Technology</i> , 2021, 55, 3430-3441.	4.6	17
63	Bacterial communities are more sensitive to ocean acidification than fungal communities in estuarine sediments. <i>FEMS Microbiology Ecology</i> , 2021, 97, .	1.3	7
64	Termite mounds reduce soil microbial diversity by filtering rare microbial taxa. <i>Environmental Microbiology</i> , 2021, 23, 2659-2668.	1.8	8
65	Potential of indigenous crop microbiomes for sustainable agriculture. <i>Nature Food</i> , 2021, 2, 233-240.	6.2	51
66	Biotic and abiotic factors distinctly drive contrasting biogeographic patterns between phyllosphere and soil resistomes in natural ecosystems. <i>ISME Communications</i> , 2021, 1, .	1.7	23
67	Soil-Food-Environment-Health Nexus for Sustainable Development. <i>Research</i> , 2021, 2021, 9804807.	2.8	15
68	Antibiotic resistance in the soil ecosystem: A One Health perspective. <i>Current Opinion in Environmental Science and Health</i> , 2021, 20, 100230.	2.1	43
69	Deciphering Potential Roles of Earthworms in Mitigation of Antibiotic Resistance in the Soils from Diverse Ecosystems. <i>Environmental Science & Technology</i> , 2021, 55, 7445-7455.	4.6	49
70	Developing Surrogate Markers for Predicting Antibiotic Resistance "Hot Spots" in Rivers Where Limited Data Are Available. <i>Environmental Science & Technology</i> , 2021, 55, 7466-7478.	4.6	21
71	Seasonal change is a major driver of soil resistomes at a watershed scale. <i>ISME Communications</i> , 2021, 1, .	1.7	20
72	Termite mound formation reduces the abundance and diversity of soil resistomes. <i>Environmental Microbiology</i> , 2021, 23, 7661-7670.	1.8	7

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73	Impact of Urbanization on Antibiotic Resistome in Different Microplastics: Evidence from a Large-Scale Whole River Analysis. <i>Environmental Science & Technology</i> , 2021, 55, 8760-8770.	4.6	57
74	Super pathogens from environmental biotechnologies threaten global health. <i>National Science Review</i> , 2021, 8, nwab110.	4.6	4
75	Novel clades of soil biphenyl degraders revealed by integrating isotope probing, multi-omics, and single-cell analyses. <i>ISME Journal</i> , 2021, 15, 3508-3521.	4.4	14
76	Antibiotic exposure decreases soil arsenic oral bioavailability in mice by disrupting ileal microbiota and metabolic profile. <i>Environment International</i> , 2021, 151, 106444.	4.8	26
77	Agricultural land-use change and rotation system exert considerable influences on the soil antibiotic resistome in Lake Tai Basin. <i>Science of the Total Environment</i> , 2021, 771, 144848.	3.9	27
78	Vertical distribution of antibiotic resistance genes in an urban green facade. <i>Environment International</i> , 2021, 152, 106502.	4.8	24
79	Mycorrhiza and Iron Tailings Synergistically Enhance Maize Resistance to Arsenic on Medium Arsenic-Polluted Soils Through Increasing Phosphorus and Iron Uptake. <i>Bulletin of Environmental Contamination and Toxicology</i> , 2021, 107, 1155-1160.	1.3	5
80	Arbuscular mycorrhizal fungi and plant diversity drive restoration of nitrogen cycling microbial communities. <i>Molecular Ecology</i> , 2021, 30, 4133-4146.	2.0	12
81	Spatial patterns of urban green space and its actual utilization status in China based on big data analysis. <i>Big Earth Data</i> , 2021, 5, 391-409.	2.0	11
82	Arsenic transformation and volatilization by arbuscular mycorrhizal symbiosis under axenic conditions. <i>Journal of Hazardous Materials</i> , 2021, 413, 125390.	6.5	14
83	High-Throughput Single-Cell Technology Reveals the Contribution of Horizontal Gene Transfer to Typical Antibiotic Resistance Gene Dissemination in Wastewater Treatment Plants. <i>Environmental Science & Technology</i> , 2021, 55, 11824-11834.	4.6	33
84	Paper-Based Devices As a New Tool for Rapid and on-Site Monitoring of "Superbugs". <i>Environmental Science & Technology</i> , 2021, 55, 12133-12135.	4.6	2
85	Environmental antimicrobial resistance is associated with faecal pollution in Central Thailand's coastal aquaculture region. <i>Journal of Hazardous Materials</i> , 2021, 416, 125718.	6.5	25
86	Longitudinal study on the effects of growth-promoting and therapeutic antibiotics on the dynamics of chicken cloacal and litter microbiomes and resistomes. <i>Microbiome</i> , 2021, 9, 178.	4.9	30
87	Precipitation increases the abundance of fungal plant pathogens in <i>Eucalyptus</i> phyllosphere. <i>Environmental Microbiology</i> , 2021, 23, 7688-7700.	1.8	20
88	Stimulation of N ₂ O emission via bacterial denitrification driven by acidification in estuarine sediments. <i>Global Change Biology</i> , 2021, 27, 5564-5579.	4.2	34
89	High Arsenic Levels Increase Activity Rather than Diversity or Abundance of Arsenic Metabolism Genes in Paddy Soils. <i>Applied and Environmental Microbiology</i> , 2021, 87, e0138321.	1.4	9
90	Continental-Scale Paddy Soil Bacterial Community Structure, Function, and Biotic Interaction. <i>MSystems</i> , 2021, 6, e0136820.	1.7	6

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91	Future research needs for environmental science in China. <i>Geography and Sustainability</i> , 2021, , .	1.9	3
92	Gammaproteobacteria, a core taxon in the guts of soil fauna, are potential responders to environmental concentrations of soil pollutants. <i>Microbiome</i> , 2021, 9, 196.	4.9	46
93	MoS ₂ Nanosheetsâ€™Cyanobacteria Interaction: Reprogrammed Carbon and Nitrogen Metabolism. <i>ACS Nano</i> , 2021, 15, 16344-16356.	7.3	28
94	Trophic level drives the host microbiome of soil invertebrates at a continental scale. <i>Microbiome</i> , 2021, 9, 189.	4.9	18
95	Raman biosensor and molecular tools for integrated monitoring of pathogens and antimicrobial resistance in wastewater. <i>TrAC - Trends in Analytical Chemistry</i> , 2021, 143, 116415.	5.8	13
96	Insights into the roles of fungi and protist in the giant panda gut microbiome and antibiotic resistome. <i>Environment International</i> , 2021, 155, 106703.	4.8	26
97	Characterization of tetracycline-resistant microbiome in soil-plant systems by combination of H218O-based DNA-Stable isotope probing and metagenomics. <i>Journal of Hazardous Materials</i> , 2021, 420, 126440.	6.5	10
98	Viral Community and Virus-Associated Antibiotic Resistance Genes in Soils Amended with Organic Fertilizers. <i>Environmental Science & Technology</i> , 2021, 55, 13881-13890.	4.6	49
99	Warming-driven migration of core microbiota indicates soil property changes at continental scale. <i>Science Bulletin</i> , 2021, 66, 2025-2035.	4.3	12
100	How can fertilization regimes and durations shape earthworm gut microbiota in a long-term field experiment?. <i>Ecotoxicology and Environmental Safety</i> , 2021, 224, 112643.	2.9	9
101	Combined pollution of arsenic and Polymyxin B enhanced arsenic toxicity and enriched ARG abundance in soil and earthworm gut microbiotas. <i>Journal of Environmental Sciences</i> , 2021, 109, 171-180.	3.2	17
102	Will a Non-antibiotic Metalloid Enhance the Spread of Antibiotic Resistance Genes: The Selenate Story. <i>Environmental Science & Technology</i> , 2021, 55, 1004-1014.	4.6	42
103	Influence of Legacy Mercury on Antibiotic Resistomes: Evidence from Agricultural Soils with Different Cropping Systems. <i>Environmental Science & Technology</i> , 2021, 55, 13913-13922.	4.6	19
104	Does biological rhythm transmit from plants to rhizosphere microbes?. <i>Environmental Microbiology</i> , 2021, 23, 6895-6906.	1.8	8
105	Technologies and perspectives for achieving carbon neutrality. <i>Innovation(China)</i> , 2021, 2, 100180.	5.2	306
106	Global meta-analysis of microplastic contamination in reservoirs with a novel framework. <i>Water Research</i> , 2021, 207, 117828.	5.3	68
107	Build in prevention and preparedness to improve climate resilience in coastal cities: Lessons from Chinaâ€™s GBA. <i>One Earth</i> , 2021, 4, 1356-1360.	3.6	13
108	Controlling pathogenic risks of water treatment biotechnologies at the source by genetic editing means. <i>Environmental Microbiology</i> , 2021, 23, 7578-7590.	1.8	9

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109	Spatial and temporal dynamics of microbiomes and resistomes in broiler litter stockpiles. <i>Computational and Structural Biotechnology Journal</i> , 2021, 19, 6201-6211.	1.9	5
110	Bioavailable arsenic and amorphous iron oxides provide reliable predictions for arsenic transfer in soil-wheat system. <i>Journal of Hazardous Materials</i> , 2020, 383, 121160.	6.5	34
111	Identification of potential electrotrophic microbial community in paddy soils by enrichment of microbial electrolysis cell biocathodes. <i>Journal of Environmental Sciences</i> , 2020, 87, 411-420.	3.2	7
112	Phosphorus fractions and oxygen isotope composition of inorganic phosphate in typical agricultural soils. <i>Chemosphere</i> , 2020, 239, 124622.	4.2	25
113	Partial replacement of inorganic phosphorus (P) by organic manure reshapes phosphate mobilizing bacterial community and promotes P bioavailability in a paddy soil. <i>Science of the Total Environment</i> , 2020, 703, 134977.	3.9	95
114	Large-scale patterns of soil antibiotic resistome in Chinese croplands. <i>Science of the Total Environment</i> , 2020, 712, 136418.	3.9	53
115	Chronic kidney disease of unknown etiology (CKDu): Using a system dynamics model to conceptualize the multiple environmental causative pathways of the epidemic. <i>Science of the Total Environment</i> , 2020, 705, 135766.	3.9	11
116	Microbial resistance promotes plant production in a four-decade nutrient fertilization experiment. <i>Soil Biology and Biochemistry</i> , 2020, 141, 107679.	4.2	59
117	Rare microbial taxa as the major drivers of ecosystem multifunctionality in long-term fertilized soils. <i>Soil Biology and Biochemistry</i> , 2020, 141, 107686.	4.2	247
118	Host identity determines plant associated resistomes. <i>Environmental Pollution</i> , 2020, 258, 113709.	3.7	23
119	Restoring Abandoned Farmland to Mitigate Climate Change on a Full Earth. <i>One Earth</i> , 2020, 3, 176-186.	3.6	60
120	Integrating Biomedical, Ecological, and Sustainability Sciences to Manage Emerging Infectious Diseases. <i>One Earth</i> , 2020, 3, 23-26.	3.6	22
121	Mediated electrochemical analysis as emerging tool to unravel links between microbial redox cycling of natural organic matter and anoxic nitrogen cycling. <i>Earth-Science Reviews</i> , 2020, 208, 103281.	4.0	10
122	Meteorological impact on the COVID-19 pandemic: A study across eight severely affected regions in South America. <i>Science of the Total Environment</i> , 2020, 744, 140881.	3.9	56
123	Coupled anaerobic methane oxidation and reductive arsenic mobilization in wetland soils. <i>Nature Geoscience</i> , 2020, 13, 799-805.	5.4	71
124	Could Global Intensification of Nitrogen Fertilisation Increase Immunogenic Proteins and Favour the Spread of Coeliac Pathology?. <i>Foods</i> , 2020, 9, 1602.	1.9	9
125	Transboundary Environmental Footprints of the Urban Food Supply Chain and Mitigation Strategies. <i>Environmental Science & Technology</i> , 2020, 54, 10460-10471.	4.6	28
126	Crop production correlates with soil multitrophic communities at the large spatial scale. <i>Soil Biology and Biochemistry</i> , 2020, 151, 108047.	4.2	43

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127	COVID-19 reveals the systemic nature of urban health globally. <i>Cities and Health</i> , 2020, , 1-5.	1.6	12
128	Temporal Dynamics of Antibiotic Resistome in the Plastisphere during Microbial Colonization. <i>Environmental Science & Technology</i> , 2020, 54, 11322-11332.	4.6	135
129	Antibiotic Resistance in the Collembolan Gut Microbiome Accelerated by the Nonantibiotic Drug Carbamazepine. <i>Environmental Science & Technology</i> , 2020, 54, 10754-10762.	4.6	25
130	Cyanobacterial blooms contribute to the diversity of antibiotic-resistance genes in aquatic ecosystems. <i>Communications Biology</i> , 2020, 3, 737.	2.0	66
131	Soil bacterial taxonomic diversity is critical to maintaining the plant productivity. <i>Environment International</i> , 2020, 140, 105766.	4.8	114
132	High starter phosphorus fertilization facilitates soil phosphorus turnover by promoting microbial functional interaction in an arable soil. <i>Journal of Environmental Sciences</i> , 2020, 94, 179-185.	3.2	14
133	Space Is More Important than Season when Shaping Soil Microbial Communities at a Large Spatial Scale. <i>MSystems</i> , 2020, 5, .	1.7	71
134	Dysbiosis in the Gut Microbiota of Soil Fauna Explains the Toxicity of Tire Tread Particles. <i>Environmental Science & Technology</i> , 2020, 54, 7450-7460.	4.6	71
135	Response to the commentary by M.W.C. Dharma-wardana on "Chronic kidney disease of unknown etiology (CKDu): Using a system dynamics model to conceptualize the multiple environmental causative pathways of the epidemic". <i>Science of the Total Environment</i> , 2020, 721, 137591.	3.9	0
136	Dam Construction as an Important Anthropogenic Activity Disturbing Soil Organic Carbon in Affected Watersheds. <i>Environmental Science & Technology</i> , 2020, 54, 7932-7941.	4.6	6
137	Microbial functional traits in phyllosphere are more sensitive to anthropogenic disturbance than in soil. <i>Environmental Pollution</i> , 2020, 265, 114954.	3.7	34
138	Abundance, diversity, and structure of Geobacteraceae community in paddy soil under long-term fertilization practices. <i>Applied Soil Ecology</i> , 2020, 153, 103577.	2.1	16
139	Changes in the environmental microbiome in the Anthropocene. <i>Global Change Biology</i> , 2020, 26, 3175-3177.	4.2	30
140	The driving factors of nematode gut microbiota under long-term fertilization. <i>FEMS Microbiology Ecology</i> , 2020, 96, .	1.3	12
141	Abundance of kinless hubs within soil microbial networks are associated with high functional potential in agricultural ecosystems. <i>Environment International</i> , 2020, 142, 105869.	4.8	158
142	Economic Valuation of Earth's Critical Zone: A Pilot Study of the Zhangxi Catchment, China. <i>Sustainability</i> , 2020, 12, 1699.	1.6	3
143	Rice Grain Cadmium Concentrations in the Global Supply-Chain. <i>Exposure and Health</i> , 2020, 12, 869-876.	2.8	63
144	The Lancet Infectious Diseases Commission on antimicrobial resistance: 6 years later. <i>Lancet Infectious Diseases</i> , The, 2020, 20, e51-e60.	4.6	161

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145	Characterization of antibiotic resistance genes and bacterial community in selected municipal and industrial sewage treatment plants beside Poyang Lake. <i>Water Research</i> , 2020, 174, 115603.	5.3	45
146	Arsenic transformation mediated by gut microbiota affects the fecundity of <i>Caenorhabditis elegans</i> . <i>Environmental Pollution</i> , 2020, 260, 113991.	3.7	8
147	The characterization of arsenic biotransformation microbes in paddy soil after straw biochar and straw amendments. <i>Journal of Hazardous Materials</i> , 2020, 391, 122200.	6.5	29
148	The Great Oxidation Event expanded the genetic repertoire of arsenic metabolism and cycling. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 10414-10421.	3.3	96
149	Effects of Earthworms on the Microbiomes and Antibiotic Resistomes of Detritus Fauna and Phyllospheres. <i>Environmental Science & Technology</i> , 2020, 54, 6000-6008.	4.6	41
150	Microbiome and antibiotic resistome in household dust from Beijing, China. <i>Environment International</i> , 2020, 139, 105702.	4.8	32
151	Prevalence of Antibiotic Resistome in Ready-to-Eat Salad. <i>Frontiers in Public Health</i> , 2020, 8, 92.	1.3	23
152	Antimicrobial Resistance is a Health Risk in Chinese Cities—Now it Has Been Mapped. <i>Urban Health and Wellbeing</i> , 2020, , 45-48.	0.3	1
153	Soil amendment with sewage sludge affects soil prokaryotic community composition, mobilome and resistome. <i>FEMS Microbiology Ecology</i> , 2019, 95, .	1.3	12
154	High-throughput characterization of antibiotic resistome in soil amended with commercial organic fertilizers. <i>Journal of Soils and Sediments</i> , 2019, 19, 641-651.	1.5	11
155	Straw biochar increases the abundance of inorganic phosphate solubilizing bacterial community for better rape (<i>Brassica napus</i>) growth and phosphate uptake. <i>Science of the Total Environment</i> , 2019, 647, 1113-1120.	3.9	76
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