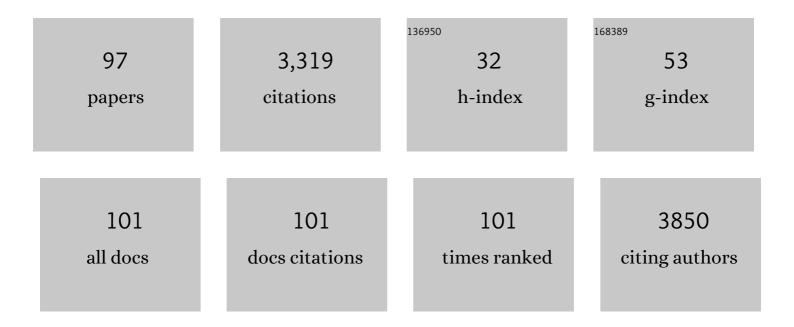


## List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Seasonal variation in the occurrence and removal of pharmaceuticals and personal care products in a wastewater treatment plant in Xiamen, China. Journal of Hazardous Materials, 2014, 277, 69-75.	12.4	223
2	Strong impact of anthropogenic contamination on the coâ€occurrence patterns of a riverine microbial community. Environmental Microbiology, 2017, 19, 4993-5009.	3.8	213
3	Influence of pretreated activated sludge for electricity generation in microbial fuel cell application. Bioresource Technology, 2013, 145, 90-96.	9.6	136
4	PPCPs in Jiulong River estuary (China): Spatiotemporal distributions, fate, and their use as chemical markers of wastewater. Chemosphere, 2016, 150, 596-604.	8.2	127
5	Biodegradation of sulfamethoxazole in bacteria from three different origins. Journal of Environmental Management, 2018, 206, 93-102.	7.8	121
6	Monitoring, mass balance and fate of pharmaceuticals and personal care products in seven wastewater treatment plants in Xiamen City, China. Journal of Hazardous Materials, 2018, 354, 81-90.	12.4	98
7	Pharmaceuticals and personal care products in a mesoscale subtropical watershed and their application as sewage markers. Journal of Hazardous Materials, 2014, 280, 696-705.	12.4	91
8	Response of bacterial communities to environmental changes in a mesoscale subtropical watershed, Southeast China. Science of the Total Environment, 2014, 472, 746-756.	8.0	88
9	Wetland plant microbial fuel cells for remediation of hexavalent chromium contaminated soils and electricity production. Journal of Hazardous Materials, 2019, 365, 137-145.	12.4	86
10	Zero-valent iron-based technologies for removal of heavy metal(loid)s and organic pollutants from the aquatic environment: Recent advances and perspectives. Journal of Cleaner Production, 2020, 277, 123478.	9.3	82
11	Characterization of triclosan metabolism in Sphingomonas sp. strain YL-JM2C. Scientific Reports, 2016, 6, 21965.	3.3	73
12	Community Structure and Function of Planktonic Crenarchaeota: Changes with Depth in the South China Sea. Microbial Ecology, 2011, 62, 549-563.	2.8	72
13	Diversity of endophytic and rhizoplane bacterial communities associated with exotic <i>Spartina alterniflora</i> and native mangrove using Illumina amplicon sequencing. Canadian Journal of Microbiology, 2015, 61, 723-733.	1.7	67
14	Community structures of ammoniaâ€oxidising archaea and bacteria in highâ€altitude lakes on the Tibetan Plateau. Freshwater Biology, 2010, 55, 2375-2390.	2.4	65
15	CO2 sequestration by methanogens in activated sludge for methane production. Applied Energy, 2015, 142, 426-434.	10.1	58
16	Bacterial community colonization on tire microplastics in typical urban water environments and associated impacting factors. Environmental Pollution, 2020, 265, 114922.	7.5	58
17	Niche Partitioning of Marine Group I Crenarchaeota in the Euphotic and Upper Mesopelagic Zones of the East China Sea. Applied and Environmental Microbiology, 2011, 77, 7469-7478.	3.1	53
18	Evaluation of Sulfadiazine Degradation in Three Newly Isolated Pure Bacterial Cultures. PLoS ONE, 2016, 11, e0165013.	2.5	52

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19	Characterization of electricity production and microbial community of food waste-fed microbial fuel cells. Chemical Engineering Research and Design, 2019, 125, 83-91.	5.6	52
20	Distinct mechanisms underlying the assembly of microeukaryotic generalists and specialists in an an anthropogenically impacted river. Science of the Total Environment, 2020, 748, 141434.	8.0	49
21	Degradation of triclocarban by a triclosan-degrading Sphingomonas sp. strain YL-JM2C. Chemosphere, 2016, 144, 292-296.	8.2	48
22	Urban ponds as hotspots of antibiotic resistome in the urban environment. Journal of Hazardous Materials, 2021, 403, 124008.	12.4	48
23	Stratified chemical and microbial characteristics between anode and cathode after long-term operation of plant microbial fuel cells for remediation of metal contaminated soils. Science of the Total Environment, 2019, 670, 585-594.	8.0	46
24	Homogeneous selection drives antibiotic resistome in two adjacent sub-watersheds, China. Journal of Hazardous Materials, 2020, 398, 122820.	12.4	46
25	Long-term impacts of silver nanoparticles in an anaerobic–anoxic–oxic membrane bioreactor system. Chemical Engineering Journal, 2015, 276, 83-90.	12.7	45
26	Horizontal and vertical gene transfer drive sediment antibiotic resistome in an urban lagoon system. Journal of Environmental Sciences, 2021, 102, 11-23.	6.1	45
27	Microbial diversity in the snow, a moraine lake and a stream in Himalayan glacier. Extremophiles, 2011, 15, 411-421.	2.3	44
28	Different community assembly mechanisms underlie similar biogeography of bacteria and microeukaryotes in Tibetan lakes. FEMS Microbiology Ecology, 2020, 96, .	2.7	43
29	A decentralized wastewater treatment system using microbial fuel cell techniques and its response to a copper shock load. Bioresource Technology, 2013, 143, 76-82.	9.6	38
30	Salinity Impact on Bacterial Community Composition in Five High-Altitude Lakes from the Tibetan Plateau, Western China. Geomicrobiology Journal, 2013, 30, 462-469.	2.0	36
31	Prokaryotic footprints in urban water ecosystems: A case study of urban landscape ponds in a coastal city, China. Environmental Pollution, 2018, 242, 1729-1739.	7.5	35
32	Biogeography of the free-living and particle-attached bacteria in Tibetan lakes. FEMS Microbiology Ecology, 2019, 95, .	2.7	35
33	Enhanced production of secondary biogenic coalbed natural gas from a subbituminous coal treated by hydrogen peroxide and its geochemical and microbiological analyses. Fuel, 2019, 236, 1345-1355.	6.4	35
34	Impacts of human disturbance on the biogeochemical nitrogen cycle in a subtropical river system revealed by nitrifier and denitrifier genes. Science of the Total Environment, 2020, 746, 141139.	8.0	35
35	Seeking key microorganisms for enhancing methane production in anaerobic digestion of waste sewage sludge. Applied Microbiology and Biotechnology, 2018, 102, 5323-5334.	3.6	34
36	Biogeography of Planktonic and Benthic Archaeal Communities in a Subtropical Eutrophic Estuary of China. Microbial Ecology, 2015, 70, 322-335.	2.8	31

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37	Domestic wastewater causes nitrate pollution in an agricultural watershed, China. Science of the Total Environment, 2022, 823, 153680.	8.0	30
38	Understanding gaseous nitrogen removal through direct measurement of dissolved N2 and N2O in a subtropical river-reservoir system. Ecological Engineering, 2014, 70, 56-67.	3.6	29
39	Altererythrobacter estronivorus sp. nov., an Estrogen-Degrading Strain Isolated from Yundang Lagoon of Xiamen City in China. Current Microbiology, 2016, 72, 634-640.	2.2	28
40	Bisphenol A attenuation in natural microcosm: Contribution of ecological components and identification of transformation pathways through stable isotope tracing. Journal of Hazardous Materials, 2020, 385, 121584.	12.4	28
41	Environmental Filtering Drives the Assembly of Habitat Generalists and Specialists in the Coastal Sand Microbial Communities of Southern China. Microorganisms, 2019, 7, 598.	3.6	27
42	Phylogenetic diversity of bacterial communities in South China Sea mesoscale cyclonic eddy perturbations. Research in Microbiology, 2011, 162, 320-329.	2.1	26
43	Community Structure of Archaea from Deep-Sea Sediments of the South China Sea. Microbial Ecology, 2010, 60, 796-806.	2.8	25
44	Response of microbial communities to bioturbation by artificially introducing macrobenthos to mudflat sediments for in situ bioremediation in a typical semi-enclosed bay, southeast China. Marine Pollution Bulletin, 2015, 94, 114-122.	5.0	24
45	Effect of a weak magnetic field on triclosan removal using zero-valent iron under aerobic and anaerobic conditions. Chemical Engineering Journal, 2018, 346, 24-33.	12.7	24
46	Archaeal community in a human-disturbed watershed in southeast China: diversity, distribution, and responses to environmental changes. Applied Microbiology and Biotechnology, 2016, 100, 4685-4698.	3.6	23
47	Contribution of biotic and abiotic factors in the natural attenuation of sulfamethoxazole: A path analysis approach. Science of the Total Environment, 2018, 633, 1217-1226.	8.0	23
48	A comparison of pelagic, littoral, and riverine bacterial assemblages in Lake Bangongco, Tibetan Plateau. FEMS Microbiology Ecology, 2014, 89, 211-221.	2.7	22
49	Melaminivora alkalimesophila gen. nov., sp. nov., a melamine-degrading betaproteobacterium isolated from a melamine-producing factory. International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 1938-1944.	1.7	22
50	Fecal pollution mediates the dominance of stochastic assembly of antibiotic resistome in an urban lagoon (Yundang lagoon), China. Journal of Hazardous Materials, 2021, 417, 126083.	12.4	22
51	Strong impact of micropollutants on prokaryotic communities at the horizontal but not vertical scales in a subtropical reservoir, China. Science of the Total Environment, 2020, 721, 137767.	8.0	19
52	Vertical variation of bacterial community in Nam Co, a large stratified lake in central Tibetan Plateau. Antonie Van Leeuwenhoek, 2016, 109, 1323-1335.	1.7	17
53	Genetic Diversity of Picocyanobacteria in Tibetan Lakes: Assessing the Endemic and Universal Distributions. Applied and Environmental Microbiology, 2014, 80, 7640-7650.	3.1	16
54	Electrochemical Characterization of a Novel Exoelectrogenic Bacterium Strain SCS5, Isolated from a Mediator-Less Microbial Fuel Cell and Phylogenetically Related to <i>Aeromonas jandaei</i> . Microbes and Environments, 2016, 31, 213-225.	1.6	16

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55	Integrated assessment of major and trace elements in surface and core sediments from an urban lagoon, China: Potential ecological risks and influencing factors. Marine Pollution Bulletin, 2021, 170, 112651.	5.0	16
56	Genome Sequence of the 17β-Estradiol-Utilizing Bacterium Sphingomonas Strain KC8. Journal of Bacteriology, 2011, 193, 4266-4267.	2.2	15
57	Deterministic and stochastic processes driving the shift in the prokaryotic community composition in wastewater treatment plants of a coastal Chinese city. Applied Microbiology and Biotechnology, 2019, 103, 9155-9168.	3.6	15
58	Tracking microeukaryotic footprint in a peri-urban watershed, China through machine-learning approaches. Science of the Total Environment, 2022, 806, 150401.	8.0	15
59	Draft Genome Sequence of Pseudomonas nitroreducens Strain TX1, Which Degrades Nonionic Surfactants and Estrogen-Like Alkylphenols. Genome Announcements, 2014, 2, .	0.8	14
60	Response of prokaryotic communities to extreme precipitation events in an urban coastal lagoon: A case study of Yundang lagoon, China. Science of the Total Environment, 2020, 706, 135937.	8.0	14
61	Dynamics of Autotrophic Marine Planktonic Thaumarchaeota in the East China Sea. PLoS ONE, 2013, 8, e61087.	2.5	13
62	Enrichment and Characterization of a Psychrotolerant Consortium Degrading Crude Oil Alkanes Under Methanogenic Conditions. Microbial Ecology, 2015, 70, 433-444.	2.8	13
63	Deciphering the Assembly Processes of the Key Ecological Assemblages of Microbial Communities in Thirteen Full-Scale Wastewater Treatment Plants. Microbes and Environments, 2019, 34, 169-179.	1.6	13
64	How habitat heterogeneity shapes bacterial and protistan communities in temperate coastal areas near estuaries. Environmental Microbiology, 2022, 24, 1775-1789.	3.8	13
65	Seasonal and spatial variations of prokaryoplankton communities in a salinity-influenced watershed, China. FEMS Microbiology Ecology, 2017, 93, .	2.7	12
66	Fate of glacier surface snowâ€originating bacteria in the glacierâ€fed hydrologic continuums. Environmental Microbiology, 2021, 23, 6450-6462.	3.8	12
67	Distinct strategies of the habitat generalists and specialists in sediment of Tibetan lakes. Environmental Microbiology, 2022, 24, 4153-4166.	3.8	12
68	Draft Genome Sequence of the Bisphenol A-Degrading Bacterium Sphingobium sp. Strain YL23. Genome Announcements, 2013, 1, .	0.8	11
69	Characterization of a novel melamine-degrading bacterium isolated from a melamine-manufacturing factory in China. Applied Microbiology and Biotechnology, 2014, 98, 3287-3293.	3.6	10
70	A Review of Micropollutant Removal by Microalgae. , 2019, , 41-55.		10
71	Long-term operation of bio-catalyzed cathodes within continuous flow membrane-less microbial fuel cells. Chemosphere, 2021, 266, 129059.	8.2	10
72	Characterization and Performance of Lactate-Feeding Consortia for Reductive Dechlorination of Trichloroethene. Microorganisms, 2021, 9, 751.	3.6	10

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73	Microbial Degradation of Phenolic Compounds. Microorganisms for Sustainability, 2019, , 305-320.	0.7	10
74	Repeated introduction of micropollutants enhances microbial succession despite stable degradation patterns. ISME Communications, 2022, 2, .	4.2	10
75	Microbial community structure analysis and isolation of vanadium-resistant strains in vanadium mining–impacted soil. Journal of Soils and Water Conservation, 2019, 74, 296-308.	1.6	9
76	Hydrothermal conversion of waste cartons into a magnetic carbon-iron composite for use as an efficient and recyclable dye adsorbent. Journal of Colloid and Interface Science, 2020, 578, 717-725.	9.4	9
77	Elemental Contaminants in Surface Sediments from Jiulong River Estuary, China: Pollution Level and Ecotoxicological Risk Assessment. Water (Switzerland), 2020, 12, 1640.	2.7	9
78	A comprehensive review on the influence of light on signaling cross-talk and molecular communication against phyto-microbiome interactions. Critical Reviews in Biotechnology, 2021, 41, 370-393.	9.0	9
79	Pathogens Removal in a Sustainable and Economic High-Rate Algal Pond Wastewater Treatment System. Sustainability, 2021, 13, 13232.	3.2	9
80	Continuous antibiotic attenuation in algal membrane photobioreactor: Performance and kinetics. Journal of Hazardous Materials, 2022, 434, 128910.	12.4	9
81	The spatial distribution of archaeal lipids in a mesoscale subtropical watershed, Southeast China. Science China Earth Sciences, 2016, 59, 1317-1328.	5.2	8
82	Assessment of the fate of silver nanoparticles in the A2O-MBR system. Science of the Total Environment, 2016, 544, 901-907.	8.0	8
83	Croceicoccus bisphenolivorans sp. nov., a bisphenol A-degrading bacterium isolated from seawater. International Journal of Systematic and Evolutionary Microbiology, 2021, 71, .	1.7	8
84	Temporal variability of microbial communities during the past 600Âyears in a Tibetan lake sediment core. Palaeogeography, Palaeoclimatology, Palaeoecology, 2021, 584, 110678.	2.3	8
85	Environmental factors shaping the archaeal community structure and ether lipid distribution in a subtropic river and estuary, China. Applied Microbiology and Biotechnology, 2018, 102, 461-474.	3.6	7
86	Integration of pre-colonized and mediator immobilized mixed culture for the improvement of electricity production of microbial fuel cells. Environmental Technology and Innovation, 2021, 22, 101514.	6.1	7
87	Draft Genome Sequence of Triclosan-Degrading Bacterium <i>Sphingomonas</i> sp. Strain YL-JM2C, Isolated from a Wastewater Treatment Plant in China. Genome Announcements, 2015, 3, .	0.8	6
88	Reconciliation of Spatiotemporal Influences on Two-Dimensional Distribution and Fate of Emerging Contaminants in a Subtropical River. ACS ES&T Water, 0, , .	4.6	6
89	Predicting Microbial Species in a River Based on Physicochemical Properties by Bio-Inspired Metaheuristic Optimized Machine Learning. Sustainability, 2019, 11, 6889.	3.2	5
90	Dispersal Limitation Expands the Diversity of Coral Microbiome Metacommunity in the South China Sea. Frontiers in Marine Science, 2021, 8, .	2.5	5

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91	Comparative study on the removal of organic pollutants by magnetic composite and pre-magnetized zero-valent iron activated persulfate. Chemosphere, 2022, 286, 131722.	8.2	5
92	Draft genome sequence of Dyadobacter tibetensis type strain (Y620-1) isolated from glacial ice. Standards in Genomic Sciences, 2014, 9, 883-892.	1.5	4
93	Elevational patterns of abundant and rare bacterial diversity and composition in mountain streams in the southeast of the Tibetan Plateau. Science China Earth Sciences, 2019, 62, 853-862.	5.2	4
94	A Comprehensive Profile of Antibiotic Resistance Genes in the Water Column of a Shallow-Sea Hydrothermal Vent Ecosystem. Sustainability, 2022, 14, 1776.	3.2	3
95	Changes in Wastewater Treatment Performance and the Microbial Community during the Bioaugmentation of a Denitrifying Pseudomonas Strain in the Low Carbon–Nitrogen Ratio Sequencing Batch Reactor. Water (Switzerland), 2022, 14, 540.	2.7	2
96	Performance Assessment of Natural Wastewater Treatment Plants by Multivariate Statistical Models: A Case Study. Sustainability, 2022, 14, 7658.	3.2	1
97	Storm Promotes the Dissemination of Antibiotic Resistome in an Urban Lagoon Through Enhancing Bio-Interactions. SSRN Electronic Journal, 0, , .	0.4	Ο