Qiang He

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5820135/publications.pdf

Version: 2024-02-01

		76326	102487
100	4,838	40	66
papers	citations	h-index	g-index
102	102	102	5281
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Global diversity and biogeography of bacterial communities in wastewater treatment plants. Nature Microbiology, 2019, 4, 1183-1195.	13.3	491
2	Strength, microstructure, efflorescence behavior and environmental impacts of waste glass geopolymers cured at ambient temperature. Journal of Cleaner Production, 2020, 252, 119610.	9.3	225
3	Mechanical and microstructural characterization of geopolymers derived from red mud and fly ashes. Journal of Cleaner Production, 2018, 186, 799-806.	9.3	180
4	Long-term successional dynamics of microbial association networks in anaerobic digestion processes. Water Research, 2016, 104, 1-10.	11.3	177
5	How sulphate-reducing microorganisms cope with stress: lessons from systems biology. Nature Reviews Microbiology, 2011, 9, 452-466.	28.6	169
6	Salt Stress in Desulfovibrio vulgaris Hildenborough: an Integrated Genomics Approach. Journal of Bacteriology, 2006, 188, 4068-4078.	2.2	155
7	Linkages of <i>Firmicutes</i> and <i>Bacteroidetes</i> populations to methanogenic process performance. Journal of Industrial Microbiology and Biotechnology, 2016, 43, 771-781.	3.0	140
8	The Electron Transfer System of Syntrophically Grown <i>Desulfovibrio vulgaris</i> . Journal of Bacteriology, 2009, 191, 5793-5801.	2.2	133
9	Characterization of Fe(III) Reduction by Chlororespiring Anaeromxyobacter dehalogenans. Applied and Environmental Microbiology, 2003, 69, 2712-2718.	3.1	126
10	Synergistic utilization of red mud for flue-gas desulfurization and fly ash-based geopolymer preparation. Journal of Hazardous Materials, 2019, 369, 503-511.	12.4	119
11	Global Analysis of Heat Shock Response in <i>Desulfovibrio vulgaris</i> Hildenborough. Journal of Bacteriology, 2006, 188, 1817-1828.	2.2	106
12	Strength properties of geopolymers derived from original and desulfurized red mud cured at ambient temperature. Construction and Building Materials, 2016, 125, 905-911.	7.2	106
13	Cell-Wide Responses to Low-Oxygen Exposure in <i>Desulfovibrio vulgaris</i> Hildenborough. Journal of Bacteriology, 2007, 189, 5996-6010.	2.2	94
14	Energetic Consequences of Nitrite Stress in Desulfovibrio vulgaris Hildenborough, Inferred from Global Transcriptional Analysis. Applied and Environmental Microbiology, 2006, 72, 4370-4381.	3.1	92
15	Genomic and microarray analysis of aromatics degradation in Geobacter metallireducens and comparison to a Geobacter isolate from a contaminated field site. BMC Genomics, 2007, 8, 180.	2.8	87
16	Sequencing of Multiple Clostridial Genomes Related to Biomass Conversion and Biofuel Production. Journal of Bacteriology, 2010, 192, 6494-6496.	2.2	81
17	Persistence of <i>Methanosaeta</i> populations in anaerobic digestion during process instability. Journal of Industrial Microbiology and Biotechnology, 2015, 42, 1129-1137.	3.0	71
18	Analysis of a Ferric Uptake Regulator (Fur) Mutant of <i>Desulfovibrio vulgaris</i> Hildenborough. Applied and Environmental Microbiology, 2007, 73, 5389-5400.	3.1	70

#	Article	IF	Citations
19	Microbial functional trait of rRNA operon copy numbers increases with organic levels in anaerobic digesters. ISME Journal, 2017, 11, 2874-2878.	9.8	70
20	Diversity, function and assembly of mangrove root-associated microbial communities at a continuous fine-scale. Npj Biofilms and Microbiomes, 2020, 6, 52.	6.4	68
21	Impact of elevated nitrate on sulfate-reducing bacteria: a comparative Study of <i>Desulfovibrio vulgaris</i> . ISME Journal, 2010, 4, 1386-1397.	9.8	67
22	Energy metabolism in Desulfovibrio vulgaris Hildenborough: insights from transcriptome analysis. Antonie Van Leeuwenhoek, 2008, 93, 347-362.	1.7	66
23	Mechanisms of enhanced cellulosic bioethanol fermentation by co-cultivation of Clostridium and Thermoanaerobacter spp Bioresource Technology, 2011, 102, 9586-9592.	9.6	66
24	Global Transcriptional, Physiological, and Metabolite Analyses of the Responses of <i>Desulfovibrio vulgaris</i> Hildenborough to Salt Adaptation. Applied and Environmental Microbiology, 2010, 76, 1574-1586.	3.1	64
25	Mechanical property and microstructure characteristics of geopolymer stabilized aggregate base. Construction and Building Materials, 2018, 191, 1120-1127.	7.2	64
26	Response of <i>Desulfovibrio vulgaris</i> to Alkaline Stress. Journal of Bacteriology, 2007, 189, 8944-8952.	2.2	62
27	Temporal Transcriptomic Analysis as Desulfovibrio vulgaris Hildenborough Transitions into Stationary Phase during Electron Donor Depletion. Applied and Environmental Microbiology, 2006, 72, 5578-5588.	3.1	57
28	Robustness of archaeal populations in anaerobic co-digestion of dairy and poultry wastes. Bioresource Technology, 2011, 102, 779-785.	9.6	57
29	Characterization of the impact of acetate and lactate on ethanolic fermentation by Thermoanaerobacter ethanolicus. Bioresource Technology, 2009, 100, 5955-5965.	9.6	55
30	SCycDB: A curated functional gene database for metagenomic profiling of sulphur cycling pathways. Molecular Ecology Resources, 2021, 21, 924-940.	4.8	52
31	Structure and regulation of the cellulose degradome in Clostridium cellulolyticum. Biotechnology for Biofuels, 2013, 6, 73.	6.2	49
32	Unexpected competitiveness of Methanosaeta populations at elevated acetate concentrations in methanogenic treatment of animal wastewater. Applied Microbiology and Biotechnology, 2017, 101, 1729-1738.	3.6	49
33	Chemical, Mechanical, and Durability Properties of Concrete with Local Mineral Admixtures under Sulfate Environment in Northwest China. Materials, 2014, 7, 3772-3785.	2.9	48
34	Microbial reduction of selenium oxyanions by Anaeromyxobacter dehalogenans. Bioresource Technology, 2010, 101, 3760-3764.	9.6	47
35	Enhancing biomethanation of municipal waste sludge with grease trap waste as a co-substrate. Renewable Energy, 2011, 36, 1802-1807.	8.9	47
36	Hydrogen peroxideâ€induced oxidative stress responses in <i>Desulfovibrio vulgaris</i> Hildenborough. Environmental Microbiology, 2010, 12, 2645-2657.	3.8	46

#	Article	IF	Citations
37	Enhancing Cd(II) sorption by red mud with heat treatment: Performance and mechanisms of sorption. Journal of Environmental Management, 2020, 255, 109866.	7.8	44
38	Impact of substrate overloading on archaeal populations in anaerobic digestion of animal waste. Journal of Applied Microbiology, 2012, 113, 1371-1379.	3.1	43
39	Correlation of Genomic and Physiological Traits of Thermoanaerobacter Species with Biofuel Yields. Applied and Environmental Microbiology, 2011, 77, 7998-8008.	3.1	42
40	Sonneratia apetala introduction alters methane cycling microbial communities and increases methane emissions in mangrove ecosystems. Soil Biology and Biochemistry, 2020, 144, 107775.	8.8	42
41	Attenuation of veterinary antibiotics in full-scale vermicomposting of swine manure via the housefly larvae (Musca domestica). Scientific Reports, 2014, 4, 6844.	3.3	41
42	Segmenting areas of potential contamination for adaptive robotic disinfection in built environments. Building and Environment, 2020, 184, 107226.	6.9	40
43	Insight into the effect of hydrogenation on efficiency of hydrothermal liquefaction and physico-chemical properties of biocrude oil. Bioresource Technology, 2014, 163, 143-151.	9.6	39
44	Experimental and Thermodynamic Study of Alkali-Activated Waste Glass and Calcium Sulfoaluminate Cement Blends: Shrinkage, Efflorescence Potential, and Phase Assemblages. Journal of Materials in Civil Engineering, 2021, 33, .	2.9	39
45	Transcriptional response of Desulfovibrio vulgaris Hildenborough to oxidative stress mimicking environmental conditions. Archives of Microbiology, 2008, 189, 451-461.	2.2	37
46	Driving forces of effluent nutrient variability in field scale bioretention. Ecological Engineering, 2016, 94, 622-628.	3.6	36
47	Toward broader applications of iron ore waste in pollution control: Adsorption of norfloxacin. Journal of Hazardous Materials, 2021, 418, 126273.	12.4	36
48	Gene Turnover Contributes to the Evolutionary Adaptation of Acidithiobacillus caldus: Insights from Comparative Genomics. Frontiers in Microbiology, 2016, 7, 1960.	3.5	35
49	Fate of free chlorine in drinking water during distribution in premise plumbing. Ecotoxicology, 2015, 24, 2151-2155.	2.4	34
50	Responses of Soil Bacteria to Long-Term and Short-Term Cadmium Stress as Revealed by Microbial Community Analysis. Bulletin of Environmental Contamination and Toxicology, 2009, 82, 367-372.	2.7	32
51	Functional Characterization of Crp/Fnr-Type Global Transcriptional Regulators in Desulfovibrio vulgaris Hildenborough. Applied and Environmental Microbiology, 2012, 78, 1168-1177.	3.1	32
52	Induction characteristics of reductive dehalogenation in the ortho-halophenol-respiring bacterium, Anaeromyxobacter dehalogenans. Biodegradation, 2002, 13, 307-316.	3.0	31
53	Removal of ciprofloxacin as an emerging pollutant: A novel application for bauxite residue reuse. Journal of Cleaner Production, 2020, 253, 120049.	9.3	28
54	Impact of building closures during the COVID-19 pandemic on Legionella infection risks. American Journal of Infection Control, 2021, 49, 1564-1566.	2.3	26

#	Article	IF	Citations
55	PERFORMANCE VARIATIONS OF COD AND NITROGEN REMOVAL BY VEGETATED SUBMERGED BED WETLANDS. Journal of the American Water Resources Association, 2002, 38, 1679-1689.	2.4	24
56	Understanding building-occupant-microbiome interactions toward healthy built environments: A review. Frontiers of Environmental Science and Engineering, 2021, 15, 65.	6.0	24
57	Acetate Threshold Concentrations Suggest Varying Energy Requirements during Anaerobic Respiration by Anaeromyxobacter dehalogenans. Applied and Environmental Microbiology, 2004, 70, 6940-6943.	3.1	23
58	Continuous Cellulosic Bioethanol Fermentation by Cyclic Fed-Batch Cocultivation. Applied and Environmental Microbiology, 2013, 79, 1580-1589.	3.1	23
59	Airborne infection risks of SARS-CoV-2 in U.S. schools and impacts of different intervention strategies. Sustainable Cities and Society, 2021, 74, 103188.	10.4	23
60	Exogenous phosphorus inputs alter complexity of soil-dissolved organic carbon in agricultural riparian wetlands. Chemosphere, 2014, 95, 572-580.	8.2	22
61	Detection of African swine fever virus-like sequences in ponds in the Mississippi Delta through metagenomic sequencing. Virus Genes, 2013, 46, 441-446.	1.6	21
62	Integrated environment-occupant-pathogen information modeling to assess and communicate room-level outbreak risks of infectious diseases. Building and Environment, 2021, 187, 107394.	6.9	21
63	A single-step DNAzyme sensor for ultra-sensitive and rapid detection of Pb2+ ions. Electrochimica Acta, 2021, 368, 137551.	5.2	19
64	Do microorganism stoichiometric alterations affect carbon sequestration in paddy soil subjected to phosphorus input?. Ecological Applications, 2015, 25, 866-879.	3.8	18
65	Healthy waterways and ecologically sustainable cities in <scp>Beijingâ€Tianjinâ€Hebei</scp> urban agglomeration (northern China): Challenges and future directions. Wiley Interdisciplinary Reviews: Water, 2021, 8, e1500.	6.5	18
66	lodine-enhanced ultrasound degradation of sulfamethazine in water. Ultrasonics Sonochemistry, 2018, 42, 759-767.	8.2	17
67	Characteristics of Cadmium Sorption by Heat-Activated Red Mud in Aqueous Solution. Scientific Reports, 2018, 8, 13558.	3.3	16
68	MCycDB: A curated database for comprehensively profiling methane cycling processes of environmental microbiomes. Molecular Ecology Resources, 2022, 22, 1803-1823.	4.8	16
69	CCl ₄ â€Enhanced Ultrasonic Irradiation for Ciprofloxacin Degradation and Antibiotic Activity. Water Environment Research, 2018, 90, 579-588.	2.7	15
70	Progressive Microbial Community Networks with Incremental Organic Loading Rates Underlie Higher Anaerobic Digestion Performance. MSystems, 2020, 5, .	3.8	15
71	Impact of alternative electron acceptors on selenium(IV) reduction by Anaeromyxobacter dehalogenans. Bioresource Technology, 2011, 102, 3578-3580.	9.6	14
72	Comparative analysis of impact of human occupancy on indoor microbiomes. Frontiers of Environmental Science and Engineering, 2021, 15, 89.	6.0	14

#	Article	IF	Citations
73	Seasonal Variations in Hydraulic Performance of Rock-Plant Filters. Environmental Technology (United Kingdom), 2001, 22, 991-999.	2.2	13
74	Fractional Characteristics of Coal Fly Ash for Beneficial Use. Journal of Materials in Civil Engineering, 2013, 25, 63-69.	2.9	12
75	Distinctive non-methanogen archaeal populations in anaerobic digestion. Applied Microbiology and Biotechnology, 2016, 100, 419-430.	3.6	12
76	Investigation of Sequential Dissolution of Asphalt Binder in Common Solvents by FTIR and Binder Fractionation. Journal of Materials in Civil Engineering, 2015, 27, .	2.9	11
77	Characterization of bacterial diversity in drinking water by pyrosequencing. Water Science and Technology: Water Supply, 2013, 13, 358-367.	2.1	10
78	The Genome of Nosema sp. Isolate YNPr: A Comparative Analysis of Genome Evolution within the Nosema/Vairimorpha Clade. PLoS ONE, 2016, 11, e0162336.	2.5	10
79	Patterns of syntrophic interactions in methanogenic conversion of propionate. Applied Microbiology and Biotechnology, 2021, 105, 8937-8949.	3.6	10
80	Development of Methanoculleus -specific real-time quantitative PCR assay for assessing methanogen communities in anaerobic digestion. Journal of Applied Microbiology, 2014, 116, 1474-1481.	3.1	9
81	Impact of Roadway Stormwater Runoff on Microbial Contamination in the Receiving Stream. Journal of Environmental Quality, 2017, 46, 1065-1071.	2.0	7
82	Passive Immunization of Chickens with Anti-Enterobactin Egg Yolk Powder for Campylobacter Control. Vaccines, 2021, 9, 569.	4.4	7
83	Linking Stoichiometric Homeostasis of Microorganisms with Soil Phosphorus Dynamics in Wetlands Subjected to Microcosm Warming. PLoS ONE, 2014, 9, e85575.	2.5	6
84	Assessing the Runoff Reduction Potential of Highway Swales and WinSLAMM as a Predictive Tool. Sustainability, 2018, 10, 2871.	3.2	6
85	Biodegradation of waste asphalt shingle by white rot fungi. Journal of Cleaner Production, 2021, 310, 127448.	9.3	6
86	Identification of Propionate-Degrading Microbial Populations in Methanogenic Processes for Waste Treatment: <i>Methanosaeta</i> and <i>Methanoculleus</i> Environmental Engineering Science, 2022, 39, 202-211.	1.6	6
87	The generation of high biomass from chlororespiring bacteria using a continuous fed-batch bioreactor. Applied Microbiology and Biotechnology, 2004, 65, 377-382.	3.6	5
88	Coping with the Environment: How Microbes Survive Environmental Challenges. International Journal of Microbiology, 2011, 2011, 1-2.	2.3	5
89	Dynamic formation of zeolite synthesized from fly ash by alkaline hydrothermal conversion. Waste Management and Research, 2013, 31, 1160-1169.	3.9	5
90	Enrichment and specific quantification of Methanocalculus in anaerobic digestion. Journal of Bioscience and Bioengineering, 2015, 120, 677-683.	2.2	5

#	Article	IF	Citations
91	Effect of planting and fertilization on lead partitioning in dredged sediment. Ecotoxicology, 2018, 27, 69-80.	2.4	5
92	Identification, Diversity and Evolution of MITEs in the Genomes of Microsporidian Nosema Parasites. PLoS ONE, 2015, 10, e0123170.	2.5	4
93	Microbiome Profiles of Nebulizers in Hospital Use. Journal of Aerosol Medicine and Pulmonary Drug Delivery, 2022, , .	1.4	4
94	Optimization of Medium Composition and Culture Conditions for Cell Multiplication of a High Quality Milk Beer Fermentation Yeast (<i>Kluyveromyces marxianus</i>). Food Science and Technology Research, 2020, 26, 351-361.	0.6	3
95	Microbiome-Based Source Identification of Microbial Contamination in Nebulizers Used by Inpatients. Journal of Hospital Infection, 2022, , .	2.9	2
96	Modeling of transient thermal dissipation of nanoscale phase-change memory cells in the pulse domain. International Journal of Heat and Mass Transfer, 2016, 94, 301-305.	4.8	1
97	Nationwide assessment of energy costs and policies to limit airborne infection risks in U.S. schools. Journal of Building Engineering, 2022, 45, 103533.	3.4	1
98	Temperature dependence of SET switching characteristics in phase-change memory cells. Journal Physics D: Applied Physics, 2016, 49, 385101.	2.8	0
99	Comparative analysis of the fermentation performance of highâ€quality milk beer strains () Tj ETQq1 1 0.784314 International Journal of Dairy Technology, 2020, 73, 552-562.	rgBT /Ov	erlock 10 T
100	Cover Image, Volume 8, Issue 2. Wiley Interdisciplinary Reviews: Water, 2021, 8, e1515.	6.5	0