

Zhisheng Yu

List of Publications by Year in descending order

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31
papers

829
citations

516561

16
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501076

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all docs

33
docs citations

33
times ranked

806
citing authors

#	ARTICLE	IF	CITATIONS
1	Pyrosequencing reveals the dominance of methylotrophic methanogenesis in a coal bed methane reservoir associated with Eastern Ordos Basin in China. <i>International Journal of Coal Geology</i> , 2012, 93, 56-61.	1.9	95
2	Production and characterization of surfactin-like biosurfactant produced by novel strain <i>Bacillus nealsonii</i> S2MT and its potential for oil contaminated soil remediation. <i>Microbial Cell Factories</i> , 2020, 19, 145.	1.9	79
3	Microbial diversity and biogenic methane potential of a thermogenic-gas coal mine. <i>International Journal of Coal Geology</i> , 2014, 134-135, 96-107.	1.9	51
4	Methylotrophic methanogenesis governs the biogenic coal bed methane formation in Eastern Ordos Basin, China. <i>Applied Microbiology and Biotechnology</i> , 2012, 96, 1587-1597.	1.7	49
5	Effect of natural microbiome and culturable biosurfactants-producing bacterial consortia of freshwater lake on petroleum-hydrocarbon degradation. <i>Science of the Total Environment</i> , 2021, 751, 141720.	3.9	47
6	Ethanol fermentation of acid-hydrolyzed cellulosic pyrolysate with <i>Saccharomyces cerevisiae</i> . <i>Bioresource Technology</i> , 2003, 90, 95-100.	4.8	43
7	Phylogenetic diversity of microbial communities associated with coalbed methane gas from Eastern Ordos Basin, China. <i>International Journal of Coal Geology</i> , 2015, 150-151, 120-126.	1.9	36
8	Microbial Diversity and Abundance in a Representative Small-Production Coal Mine of Central China. <i>Energy & Fuels</i> , 2013, 27, 3821-3829.	2.5	33
9	Bioconversion of coal to methane by microbial communities from soil and from an opencast mine in the Xilingol grassland of northeast China. <i>Biotechnology for Biofuels</i> , 2019, 12, 236.	6.2	33
10	Cloning of a novel levoglucosan kinase gene from <i>Lipomyces starkeyi</i> and its expression in <i>Escherichia coli</i> . <i>World Journal of Microbiology and Biotechnology</i> , 2009, 25, 1589-1595.	1.7	31
11	Microbial communities from the Huaibei Coalfield alter the physicochemical properties of coal in methanogenic bioconversion. <i>International Journal of Coal Geology</i> , 2019, 202, 85-94.	1.9	26
12	How does biochar amendment affect soil methane oxidation? A review. <i>Journal of Soils and Sediments</i> , 2021, 21, 1575-1586.	1.5	25
13	Microbial distribution and variation in produced water from separators to storage tanks of shale gas wells in Sichuan Basin, China. <i>Environmental Science: Water Research and Technology</i> , 2017, 3, 340-351.	1.2	24
14	Seasonal Changes in Bacterial Communities Cause Foaming in a Wastewater Treatment Plant. <i>Microbial Ecology</i> , 2016, 71, 660-671.	1.4	21
15	Mathematical modeling of the fermentation of acid-hydrolyzed pyrolytic sugars to ethanol by the engineered strain <i>Escherichia coli</i> ATCC 11177. <i>Applied Microbiology and Biotechnology</i> , 2015, 99, 4093-4105.	1.7	19
16	Purification and characterization of levoglucosan kinase from <i>Lipomyces starkeyi</i> YZ-215. <i>World Journal of Microbiology and Biotechnology</i> , 2008, 24, 15-22.	1.7	16
17	Proteomic and metabolomic analysis of the cellular biomarkers related to inhibitors tolerance in <i>Zymomonas mobilis</i> ZM4. <i>Biotechnology for Biofuels</i> , 2018, 11, 283.	6.2	14
18	Habitat filtering shapes the differential structure of microbial communities in the Xilingol grassland. <i>Scientific Reports</i> , 2019, 9, 19326.	1.6	14

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19	Evaluation of di-rhamnolipid biosurfactants production by a novel <i>Pseudomonas</i> sp. S1WB: Optimization, characterization and effect on petroleum-hydrocarbon degradation. <i>Ecotoxicology and Environmental Safety</i> , 2022, 242, 113892.	2.9	13
20	Biomass pyrolysis liquid to citric acid via 2-step bioconversion. <i>Microbial Cell Factories</i> , 2014, 13, 182.	1.9	11
21	Gut region induces gastrointestinal microbiota community shift in Ujimqin sheep (<i>Ovis</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 10	1.8	10
22	Conversion efficiency of bioethanol from levoglucosan was improved by the newly engineered <i>Escherichia coli</i> . <i>Environmental Progress and Sustainable Energy</i> , 2021, 40, e13687.	1.3	8
23	Response of soil bacterial communities to moisture and grazing in the Tibetan alpine steppes on a small spatial scale. <i>Geomicrobiology Journal</i> , 2019, 36, 559-569.	1.0	6
24	Assessing the Effect of Physicochemical Properties of Saline and Sodic Soil on Soil Microbial Communities. <i>Agriculture (Switzerland)</i> , 2022, 12, 782.	1.4	6
25	Regeneration of unconventional natural gas by methanogens co-existing with sulfate-reducing prokaryotes in deep shale wells in China. <i>Scientific Reports</i> , 2020, 10, 16042.	1.6	5
26	Deciphering the initial products of coal during methanogenic bioconversion: Based on an untargeted metabolomics approach. <i>GCB Bioenergy</i> , 2021, 13, 967-978.	2.5	5
27	Inhibitor tolerance and bioethanol fermentability of levoglucosan-utilizing <i>Escherichia coli</i> were enhanced by overexpression of stress-responsive gene <i>ycfR</i> : The proteomics-guided metabolic engineering. <i>Synthetic and Systems Biotechnology</i> , 2021, 6, 384-395.	1.8	5
28	Enhanced straw fermentation process based on microbial electrolysis cell coupled anaerobic digestion. <i>Chinese Journal of Chemical Engineering</i> , 2022, 44, 239-245.	1.7	4
29	Study on Adsorption of As(III) by a New Bio-Material from Chitin Pyrolysis. <i>Water (Switzerland)</i> , 2021, 13, 2944.	1.2	4
30	Only mass migration of fungi runs through the biotopes of soil, phyllosphere, and feces. <i>Journal of Soils and Sediments</i> , 2021, 21, 1151-1164.	1.5	2
31	Omics analysis coupled with gene editing revealed potential transporters and regulators related to levoglucosan metabolism efficiency of the engineered <i>Escherichia coli</i> . , 2022, 15, 2.		0