Yuxiu Zhang

List of Publications by Year in descending order

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414414 394421 1,288 32 19 32 citations h-index g-index papers 35 35 35 1659 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Bioaugmentation of quinoline-degrading bacteria for coking wastewater treatment: performance and microbial community analysis. Journal of Environmental Science and Health - Part A Toxic/Hazardous Substances and Environmental Engineering, 2022, 57, 601-619.	1.7	1
2	Performance prediction of ZVI-based anaerobic digestion reactor using machine learning algorithms. Waste Management, 2021, 121, 59-66.	7.4	56
3	Vegetation dynamics of coal mining city in an arid desert region of Northwest China from 2000 to 2019. Journal of Arid Land, 2021, 13, 534-547.	2.3	8
4	Deciphering soil bacterial community structure in subsidence area caused by underground coal mining in arid and semiarid area. Applied Soil Ecology, 2021, 163, 103916.	4.3	20
5	Characteristics of biochars prepared by co-pyrolysis of sewage sludge and cotton stalk intended for use as soil amendments. Environmental Technology (United Kingdom), 2020, 41, 1347-1357.	2.2	35
6	New insights of enhanced anaerobic degradation of refractory pollutants in coking wastewater: Role of zero-valent iron in metagenomic functions. Bioresource Technology, 2020, 300, 122667.	9.6	36
7	Rhizosphere Soil Microbial Properties on Tetraena mongolica in the Arid and Semi-Arid Regions, China. International Journal of Environmental Research and Public Health, 2020, 17, 5142.	2.6	9
8	The Response of Arbuscular Mycorrhizal Fungal Communities to the Soil Environment of Underground Mining Subsidence Area in Northwest China. International Journal of Environmental Research and Public Health, 2020, 17, 9157.	2.6	4
9	Land Subsidence in a Coal Mining Area Reduced Soil Fertility and Led to Soil Degradation in Arid and Semi-Arid Regions. International Journal of Environmental Research and Public Health, 2019, 16, 3929.	2.6	66
10	Occurrence and Succession of Bacterial Community in O3/BAC Process of Drinking Water Treatment. International Journal of Environmental Research and Public Health, 2019, 16, 3112.	2.6	4
11	A novel phenol and ammonia recovery process for coal gasification wastewater altering the bacterial community and increasing pollutants removal in anaerobic/anoxic/aerobic system. Science of the Total Environment, 2019, 661, 203-211.	8.0	33
12	The enhancement of pyridine degradation by < i>Rhodococcus < /i>KDPy1 in coking wastewater. FEMS Microbiology Letters, 2019, 366, .	1.8	14
13	Metagenomic insights into the microbiota profiles and bioaugmentation mechanism of organics removal in coal gasification wastewater in an anaerobic/anoxic/oxic system by methanol. Bioresource Technology, 2018, 264, 106-115.	9.6	53
14	The response of soil bacterial communities to mining subsidence in the west China aeolian sand area. Applied Soil Ecology, 2017, 121, 1-10.	4.3	61
15	Improvement of oxytetracycline production mediated via cooperation of resistance genes in Streptomyces rimosus. Science China Life Sciences, 2017, 60, 992-999.	4.9	32
16	Isolation and characterization of a novel cadmium-regulated Yellow Stripe-Like transporter (SnYSL3) in Solanum nigrum. Plant Cell Reports, 2017, 36, 281-296.	5.6	74
17	Growth and Cadmium Accumulation of Solanum nigrum L. Seedling were Enhanced by Heavy Metal-Tolerant Strains of Pseudomonas aeruginosa. Water, Air, and Soil Pollution, 2016, 227, 1.	2.4	26
18	Analysis of polycyclic aromatic hydrocarbons (PAHs) and their adsorption characteristics on activated sludge during biological treatment of coking wastewater. Desalination and Water Treatment, 2016, 57, 23633-23643.	1.0	6

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19	Quinoline-degrading strain <i>Pseudomonas aeruginosa</i> KDQ4 isolated from coking activated sludge is capable of the simultaneous removal of phenol in a dual substrate system. Journal of Environmental Science and Health - Part A Toxic/Hazardous Substances and Environmental Engineering, 2016, 51, 1139-1148.	1.7	14
20	Impacts of addition of natural zeolite or a nitrification inhibitor on antibiotic resistance genes during sludge composting. Water Research, 2016, 91, 339-349.	11.3	255
21	Analysis of a diverse bacterial community and degradation of organic compounds in a bioprocess for coking wastewater treatment. Desalination and Water Treatment, 2016, 57, 19096-19105.	1.0	11
22	Identification of a cluster-situated activator of oxytetracycline biosynthesis and manipulation of its expression for improved oxytetracycline production in Streptomyces rimosus. Microbial Cell Factories, 2015, 14, 46.	4.0	50
23	A copper-responsive gene cluster is required for copper homeostasis and contributes to oxidative resistance in Deinococcus radiodurans R1. Molecular BioSystems, 2014, 10, 2607-2616.	2.9	9
24	Silicon attenuates cadmium toxicity in Solanum nigrum L. by reducing cadmium uptake and oxidative stress. Plant Physiology and Biochemistry, 2013, 68, 1-7.	5.8	108
25	Cloning, characterization, and expression of the BjEXPA1 gene and its promoter region from Brassica juncea L Plant Growth Regulation, 2011, 64, 39-51.	3.4	14
26	Silicon-induced alleviation of cadmium toxicity in hyperaccumulator Solanum nigrum L. , 2011 , , .		0
27	Antioxidative response of metal-accumulator and non-accumulator plants under cadmium stress. Plant and Soil, 2008, 310, 137-149.	3.7	121
28	Expression and function of two dehydrins under environmental stresses in Brassica juncea L Molecular Breeding, 2008, 21, 431-438.	2.1	26
29	Indian Mustard Aquaporin Improves Drought and Heavy-metal Resistance in Tobacco. Molecular Biotechnology, 2008, 40, 280-292.	2.4	61
30	Rapid in vitro multiplication and ex vitro rooting of Malus zumi (Matsumura) Rehd. Acta Physiologiae Plantarum, 2007, 30, 129-132.	2.1	38
31	Cloning and Expression Analysis of SKn-Type Dehydrin Gene From Bean in Response to Heavy Metals. Molecular Biotechnology, 2006, 32, 205-218.	2.4	38
32	PCR-enriched cDNA pool method for cloning of gene homologues. Plant Molecular Biology Reporter, 2005, 23, 219-226.	1.8	4