Zhenmei Lu

List of Publications by Year in descending order

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53	1,654	19	39
papers	citations	h-index	g-index
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58	58	58	2034
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Novel pathway and acetate-facilitated complete atenolol degradation by Hydrogenophaga sp. YM1 isolated from activated sludge. Science of the Total Environment, 2022, 810, 152218.	8.0	15
2	A Multicomponent THF Hydroxylase Initiates Tetrahydrofuran Degradation in <i>Cupriavidus metallidurans</i> ZM02. Applied and Environmental Microbiology, 2022, 88, AEM0188021.	3.1	1
3	Network and meta-omics reveal the cooperation patterns and mechanisms in an efficient 1,4-dioxane-degrading microbial consortium. Chemosphere, 2022, 301, 134723.	8.2	9
4	Microbial interactions enhanced environmental fitness and expanded ecological niches under dibutyl phthalate and cadmium co-contamination. Environmental Pollution, 2022, 306, 119362.	7.5	11
5	Network-directed isolation of the cooperator Pseudomonas aeruginosa ZM03 enhanced the dibutyl phthalate degradation capacity of Arthrobacter nicotianae ZM05 under pH stress. Journal of Hazardous Materials, 2021, 410, 124667.	12.4	19
6	Identification and characterization of Nornicotine degrading strain Arthrobacter sp. NOR5. Science of the Total Environment, 2021, 764, 142894.	8.0	5
7	Efficient electrotransformation of Rhodococcus ruber YYL with abundant extracellular polymeric substances via a cell wall-weakening strategy. FEMS Microbiology Letters, 2021, 368, .	1.8	1
8	Acinetobacter tandoii ZM06 Assists Glutamicibacter nicotianae ZM05 in Resisting Cadmium Pressure to Preserve Dipropyl Phthalate Biodegradation. Microorganisms, 2021, 9, 1417.	3.6	12
9	Impact of exogenous nitrogen on the cyanobacterial abundance and community in oil-contaminated sediment: A microcosm study. Science of the Total Environment, 2020, 710, 136296.	8.0	4
10	Microplastics in the soil environment: Occurrence, risks, interactions and fate – A review. Critical Reviews in Environmental Science and Technology, 2020, 50, 2175-2222.	12.8	324
11	Biodegradation of Tetrahydrofuran by the Newly Isolated Filamentous Fungus Pseudallescheria boydii ZM01. Microorganisms, 2020, 8, 1190.	3.6	9
12	3-Hydroxypyridine Dehydrogenase HpdA Is Encoded by a Novel Four-Component Gene Cluster and Catalyzes the First Step of 3-Hydroxypyridine Catabolism in Ensifer adhaerens HP1. Applied and Environmental Microbiology, 2020, 86, .	3.1	5
13	Thiamine-Mediated Cooperation Between Auxotrophic Rhodococcus ruber ZM07 and Escherichia coli K12 Drives Efficient Tetrahydrofuran Degradation. Frontiers in Microbiology, 2020, 11, 594052.	3. 5	4
14	Bacterial catabolism of nicotine: Catabolic strains, pathways and modules. Environmental Research, 2020, 183, 109258.	7.5	24
15	Metabolite Cross-Feeding between Rhodococcus ruber YYL and Bacillus cereus MLY1 in the Biodegradation of Tetrahydrofuran under pH Stress. Applied and Environmental Microbiology, 2019, 85, .	3.1	16
16	Complex Oxidation of Apocytochromes <i>c</i> during Bacterial Cytochrome <i>c</i> Maturation. Applied and Environmental Microbiology, 2019, 85, .	3.1	16
17	Enrichment and characterization of a highly efficient tetrahydrofuran-degrading bacterial culture. Biodegradation, 2019, 30, 467-479.	3.0	13
18	Novel 3,6-Dihydroxypicolinic Acid Decarboxylase-Mediated Picolinic Acid Catabolism in <i>Alcaligenes faecalis</i> JQ135. Journal of Bacteriology, 2019, 201, .	2.2	8

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19	Novel tetrahydrofuran (THF) degradation-associated genes and cooperation patterns of a THF-degrading microbial community as revealed by metagenomic. Chemosphere, 2019, 231, 173-183.	8.2	10
20	Identification and Characterization of a Novel $\langle i \rangle$ pic $\langle i \rangle$ Gene Cluster Responsible for Picolinic Acid Degradation in Alcaligenes faecalis JQ135. Journal of Bacteriology, 2019, 201, .	2.2	18
21	Bacterial degradation of anthraquinone dyes. Journal of Zhejiang University: Science B, 2019, 20, 528-540.	2.8	83
22	ndpT encodes a new protein involved in nicotine catabolism by Sphingomonas melonis TY. Applied Microbiology and Biotechnology, 2018, 102, 10171-10181.	3.6	8
23	Cyanobacterial blooms in oilâ€contaminated subtidal sediments revealed by integrated approaches. Environmental Microbiology Reports, 2018, 10, 444-452.	2.4	5
24	Characterization of a Novel Nicotine Degradation Gene Cluster ndp in Sphingomonas melonis TY and Its Evolutionary Analysis. Frontiers in Microbiology, 2017, 8, 337.	3.5	16
25	pH Stress-Induced Cooperation between Rhodococcus ruber YYL and Bacillus cereus MLY1 in Biodegradation of Tetrahydrofuran. Frontiers in Microbiology, 2017, 8, 2297.	3.5	25
26	The Complete Genome Sequence of the Nicotine-Degrading Bacterium Shinella sp. HZN7. Frontiers in Microbiology, 2016, 7, 1348.	3.5	24
27	Conversion of nornicotine to 6-hydroxy-nornicotine and 6-hydroxy-myosmine by Shinella sp. strain HZN7. Applied Microbiology and Biotechnology, 2016, 100, 10019-10029.	3.6	14
28	Impact of 4-epi-oxytetracycline on the gut microbiota and blood metabolomics of Wistar rats. Scientific Reports, 2016, 6, 23141.	3.3	12
29	Short-term toxicity assessments of an antibiotic metabolite in Wistar rats and its metabonomics analysis by ultra-high performance liquid chromatography coupled to quadrupole time-of-flight mass spectrometry. Toxicology and Applied Pharmacology, 2016, 293, 1-9.	2.8	23
30	Two Novel Sets of Genes Essential for Nicotine Degradation by Sphingomonas melonis TY. Frontiers in Microbiology, 2016, 7, 2060.	3.5	12
31	Trehalose promotes Rhodococcus sp. strain YYL colonization in activated sludge under tetrahydrofuran (THF) stress. Frontiers in Microbiology, 2015, 6, 438.	3.5	7
32	Individual or synchronous biodegradation of di-n-butyl phthalate and phenol by Rhodococcus ruber strain DP-2. Journal of Hazardous Materials, 2014, 273, 104-109.	12.4	36
33	Bioremediation of the tobacco waste-contaminated soil by Pseudomonas sp. HF-1: nicotine degradation and microbial community analysis. Applied Microbiology and Biotechnology, 2013, 97, 6077-6088.	3.6	14
34	Successful bioaugmentation of an activated sludge reactor with Rhodococcus sp. YYL for efficient tetrahydrofuran degradation. Journal of Hazardous Materials, 2013, 261, 550-558.	12.4	22
35	Biodegradation of di-n-butyl phthalate by a stable bacterial consortium, HD-1, enriched from activated sludge. Bioresource Technology, 2013, 128, 526-532.	9.6	75
36	Correction to GeoChip-Based Analysis of Microbial Functional Gene Diversity in a Landfill Leachate-Contaminated Aquifer. Environmental Science & Envir	10.0	1

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37	GeoChip-Based Analysis of Microbial Functional Gene Diversity in a Landfill Leachate-Contaminated Aquifer. Environmental Science & Environmental Scien	10.0	64
38	Microbial gene functions enriched in the Deepwater Horizon deep-sea oil plume. ISME Journal, 2012, 6, 451-460.	9.8	240
39	Unraveling the concentration-dependent metabolic response of Pseudomonas sp. HF-1 to nicotine stress by 1H NMR-based metabolomics. Ecotoxicology, 2012, 21, 1314-1324.	2.4	22
40	The effect of tetrahydrofuran on the enzymatic activity and microbial community in activated sludge from a sequencing batch reactor. Ecotoxicology, 2012, 21, 56-65.	2.4	15
41	Nicotine degradation by two novel bacterial isolates of Acinetobacter sp. TW and Sphingomonas sp. TY and their responses in the presence of neonicotinoid insecticides. World Journal of Microbiology and Biotechnology, 2011, 27, 1633-1640.	3.6	47
42	Reduction of oxidative stress by bioaugmented strain Pseudomonas sp. HF-1 and selection of potential biomarkers in sequencing batch reactor treating tobacco wastewater. Ecotoxicology, 2010, 19, 1117-1123.	2.4	13
43	Isolation and characterization of a new strain of Methanothermobacter marburgensis DX01 from hot springs in China. Anaerobe, 2010, 16, 54-59.	2.1	10
44	Assessment of toxicity of tetrahydrofuran on the microbial community in activated sludge. Bioresource Technology, 2010, 101, 5213-5221.	9.6	52
45	Antioxidant Enzyme Activity in Bacterial Resistance to Nicotine Toxicity by Reactive Oxygen Species. Archives of Environmental Contamination and Toxicology, 2009, 57, 456-462.	4.1	10
46	Studies on biosorption equilibrium and kinetics of Cd2+ by Streptomyces sp. K33 and HL-12. Journal of Hazardous Materials, 2009, 164, 423-431.	12.4	41
47	Isolation, identification and characterization of a novel Rhodococcus sp. strain in biodegradation of tetrahydrofuran and its medium optimization using sequential statistics-based experimental designs. Bioresource Technology, 2009, 100, 2762-2769.	9.6	60
48	Catalase and superoxide dismutase activities in a Stenotrophomonas maltophilia WZ2 resistant to herbicide pollution. Ecotoxicology and Environmental Safety, 2009, 72, 136-143.	6.0	44
49	Bioaugmentation with the nicotine-degrading bacterium Pseudomonas sp. HF-1 in a sequencing batch reactor treating tobacco wastewater: Degradation study and analysis of its mechanisms. Water Research, 2009, 43, 4187-4196.	11.3	74
50	Variations of Bacterial Community Structure in Flooded Paddy Soil Contaminated with Herbicide Quinclorac. Journal of Environmental Science and Health - Part B Pesticides, Food Contaminants, and Agricultural Wastes, 2006, 41, 821-832.	1.5	14
51	The Response ofEscherichia coli,Bacillus subtilis, andBurkholderia cepaciaWZ1 to Oxidative Stress of Exposure to Quinclorac. Journal of Environmental Science and Health - Part B Pesticides, Food Contaminants, and Agricultural Wastes, 2004, 39, 431-441.	1.5	17
52	Phylogenetic and Degradation Characterization of Burkholderia cepacia WZ1 Degrading Herbicide Quinclorac. Journal of Environmental Science and Health - Part B Pesticides, Food Contaminants, and Agricultural Wastes, 2003, 38, 771-782.	1.5	21
53	Influences of quinclorac on culturable microorganisms and soil respiration in flooded paddy soil. Biomedical and Environmental Sciences, 2003, 16, 314-22.	0.2	4