## Ming-Rong Deng

List of Publications by Year in descending order

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759233 794594 29 455 12 19 h-index g-index citations papers 29 29 29 297 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Flavobacterium proteolyticum sp. nov., isolated from aquaculture water. Archives of Microbiology, 2022, 204, 146.	2.2	4
2	Upstream Activation Sequence Can Function as an Insulator for Chromosomal Regulation of Heterologous Pathways Against Position Effects in Saccharomyces cerevisiae. Applied Biochemistry and Biotechnology, 2022, , 1.	2.9	1
3	Engineering a Balanced Acetyl Coenzyme A Metabolism in <i>Saccharomyces cerevisiae</i> for Lycopene Production through Rational and Evolutionary Engineering. Journal of Agricultural and Food Chemistry, 2022, 70, 4019-4029.	5.2	14
4	Croceicoccus gelatinilyticus sp. nov., isolated from a tidal flat sediment. Archives of Microbiology, 2022, 204, 93.	2.2	10
5	Comparative Genomics Reveals Genetic Diversity and Metabolic Potentials of the Genus <i>Qipengyuania</i> and Suggests Fifteen Novel Species. Microbiology Spectrum, 2022, 10, e0126421.	3.0	55
6	Transcriptome Analysis Reveals a Promotion of Carotenoid Production by Copper Ions in Recombinant Saccharomyces cerevisiae. Microorganisms, 2021, 9, 233.	3.6	9
7	Roseibium litorale sp. nov., isolated from a tidal flat sediment and proposal for the reclassification of Labrenzia polysiphoniae as Roseibium polysiphoniae comb. nov International Journal of Systematic and Evolutionary Microbiology, 2021, 71, .	1.7	15
8	Inhella proteolytica sp. nov. and Inhella gelatinilytica sp. nov., two novel species of the genus Inhella isolated from aquaculture water. Archives of Microbiology, 2021, 203, 3191-3200.	2.2	12
9	Qipengyuania soli sp. nov., Isolated from Mangrove Soil. Current Microbiology, 2021, 78, 2806-2814.	2.2	11
10	Comparative genomic analysis of the genus Novosphingobium and the description of two novel species Novosphingobium aerophilum sp. nov. and Novosphingobium jiangmenense sp. nov. Systematic and Applied Microbiology, 2021, 44, 126202.	2.8	31
11	Identification of a novel metabolic engineering target for carotenoid production in Saccharomyces cerevisiae via ethanol-induced adaptive laboratory evolution. Bioresources and Bioprocessing, 2021, 8,	4.2	6
12	Chitinilyticum piscinae sp. nov., isolated from aquaculture water. International Journal of Systematic and Evolutionary Microbiology, 2021, 71, .	1.7	6
13	Salipiger mangrovisoli sp. nov., isolated from mangrove soil and the proposal for the reclassification of Paraphaeobacter pallidus as Salipiger pallidus comb. nov International Journal of Systematic and Evolutionary Microbiology, 2021, 71, .	1.7	15
14	Phylogenomic Analysis Substantiates the gyrB Gene as a Powerful Molecular Marker to Efficiently Differentiate the Most Closely Related Genera Myxococcus, Corallococcus, and Pyxidicoccus. Frontiers in Microbiology, 2021, 12, 763359.	<b>3.</b> 5	8
15	Discovery of Mycothiogranaticins from Streptomyces vietnamensis GIMV4.0001 and the Regulatory Effect of Mycothiol on the Granaticin Biosynthesis. Frontiers in Chemistry, 2021, 9, 802279.	3.6	9
16	An aberrant metabolic flow toward early shunt products in the granaticin biosynthetic machinery of Streptomyces vietnamensis GIMV4.0001. Journal of Antibiotics, 2020, 73, 260-264.	2.0	7
17	A myxobacterial LPMO10 has oxidizing cellulose activity for promoting biomass enzymatic saccharification of agricultural crop straws. Bioresource Technology, 2020, 318, 124217.	9.6	21
18	Characterization and Crystal Structure of a Nonheme Diiron Monooxygenase Involved in Platensimycin and Platencin Biosynthesis. Journal of the American Chemical Society, 2019, 141, 12406-12412.	13.7	23

#	Article	IF	CITATION
19	Cryptic and Stereospecific Hydroxylation, Oxidation, and Reduction in Platensimycin and Platencin Biosynthesis. Journal of the American Chemical Society, 2019, 141, 4043-4050.	13.7	25
20	Discovery of the Tiancilactone Antibiotics by Genome Mining of Atypical Bacterial Typeâ€II Diterpene Synthases. ChemBioChem, 2018, 19, 1727-1733.	2.6	18
21	Complete genome sequence of Streptomyces vietnamensis GIMV4.0001 T, a genetically manipulable producer of the benzoisochromanequinone antibiotic granaticin. Journal of Biotechnology, 2015, 200, 6-7.	3.8	7
22	Description of a Gram-negative bacterium, Sphingomonas guangdongensis sp. nov International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 1697-1702.	1.7	26
23	Sphingomonas gimensis sp. nov., a novel Gram-negative bacterium isolated from abandoned lead–zinc ore mine. Antonie Van Leeuwenhoek, 2014, 105, 1091-1097.	1.7	23
24	Acinetobacter guangdongensis sp. nov., isolated from abandoned lead–zinc ore. International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 3417-3421.	1.7	14
25	Streptomyces vietnamensis GIMV4.0001: a granaticin-producing strain that can be readily genetically manipulated. Journal of Antibiotics, 2011, 64, 345-347.	2.0	12
26	Granaticins and their biosynthetic gene cluster from Streptomyces vietnamensis: evidence of horizontal gene transfer. Antonie Van Leeuwenhoek, 2011, 100, 607-617.	1.7	17
27	Streptomyces caeruleatus sp. nov., with dark blue diffusible pigment. International Journal of Systematic and Evolutionary Microbiology, 2011, 61, 507-511.	1.7	16
28	Detection of a novel bacterium associated with spores of the arbuscular mycorrhizal fungus <i>Gigaspora margarita</i> . Canadian Journal of Microbiology, 2009, 55, 771-775.	1.7	6
29	Streptomyces vietnamensis sp. nov., a streptomycete with violet–blue diffusible pigment isolated from soil in Vietnam. International Journal of Systematic and Evolutionary Microbiology, 2007, 57, 1770-1774.	1.7	34