Jun Li

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8919752/publications.pdf

Version: 2024-02-01

11	109	7	10
papers	citations	h-index	g-index
11	11	11	111
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Combined genomic and transcriptomic analysis of the dibutyl phthalate metabolic pathway in <i>Arthrobacter</i> sp. ZJUTW. Biotechnology and Bioengineering, 2020, 117, 3712-3726.	3.3	21
2	Dissemination of Genetic Acquisition/Loss Provides a Variety of Quorum Sensing Regulatory Properties in Pseudoalteromonas. International Journal of Molecular Sciences, 2018, 19, 3636.	4.1	16
3	Characterization of <scp>NMCR</scp> â€2, a new <scp>nonâ€mobile</scp> colistin resistance enzyme: implications for an <scp>MCR</scp> â€8 ancestor. Environmental Microbiology, 2021, 23, 844-860.	3.8	12
4	Metabolic Engineering of <i>Saccharomyces cerevisiae</i> for High-Level Production of Chlorogenic Acid from Glucose. ACS Synthetic Biology, 2022, 11, 800-811.	3.8	12
5	Co-occurrence of functional modules derived from nicotine-degrading gene clusters confers additive effects in Pseudomonas sp. JY-Q. Applied Microbiology and Biotechnology, 2019, 103, 4499-4510.	3.6	11
6	Coordinated regulation of anthranilate metabolism and bacterial virulence by the GntR family regulator MpaR in <i>Pseudomonas aeruginosa</i> . Molecular Microbiology, 2020, 114, 857-869.	2.5	10
7	Whole-Genome Comparative and Pathogenicity Analysis of <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Rissen. G3: Genes, Genomes, Genetics, 2020, 10, 2159-2170.	1.8	8
8	A Type VI Secretion System Facilitates Fitness, Homeostasis, and Competitive Advantages for Environmental Adaptability and Efficient Nicotine Biodegradation. Applied and Environmental Microbiology, 2021, 87, .	3.1	7
9	Expression and functional identification of two homologous nicotine dehydrogenases, NicA2 and Nox, from Pseudomonas sp. JY-Q. Protein Expression and Purification, 2021, 178, 105767.	1.3	6
10	Impacts of horizontal gene transfer on the compact genome of the clavulanic acid-producing Streptomyces strain F613-1. 3 Biotech, 2018, 8, 472.	2.2	3
11	Comparative insights into multiple drug resistance determinants in Stenotrophomonas maltophilia MER1. Journal of Global Antimicrobial Resistance, 2021, 27, 20-25.	2.2	3