

Sarika Mehra

List of Publications by Year in descending order

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papers

667
citations

567144

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33
docs citations

33
times ranked

969
citing authors

#	ARTICLE	IF	CITATIONS
1	Potentiating the Anti-Tuberculosis Efficacy of Peptide Nucleic Acids through Combinations with Permeabilizing Drugs. <i>Microbiology Spectrum</i> , 2022, 10, e0126221.	1.2	5
2	Involvement of the SCO3366 efflux pump from <i>S. coelicolor</i> in rifampicin resistance and its regulation by a TetR regulator. <i>Applied Microbiology and Biotechnology</i> , 2022, 106, 2175-2190.	1.7	4
3	PNA-mediated efflux inhibition as a therapeutic strategy towards overcoming drug resistance in <i>Mycobacterium smegmatis</i> . <i>Microbial Pathogenesis</i> , 2021, 151, 104737.	1.3	2
4	A Major Facilitator Superfamily (MFS) Efflux Pump, SCO4121, from <i>Streptomyces coelicolor</i> with Roles in Multidrug Resistance and Oxidative Stress Tolerance and Its Regulation by a MarR Regulator. <i>Applied and Environmental Microbiology</i> , 2021, 87, .	1.4	19
5	The Mycobacterial Efflux Pump EfpA Can Induce High Drug Tolerance to Many Antituberculosis Drugs, Including Moxifloxacin, in <i>Mycobacterium smegmatis</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2021, 65, e0026221.	1.4	4
6	pH-driven enhancement of anti-tubercular drug loading on iron oxide nanoparticles for drug delivery in macrophages. <i>Beilstein Journal of Nanotechnology</i> , 2021, 12, 1127-1139.	1.5	0
7	Ethanol in Combination with Oxidative Stress Significantly Impacts Mycobacterial Physiology. <i>Journal of Bacteriology</i> , 2020, 202, .	1.0	1
8	Enhancing titers and productivity of rCHO clones with a combination of an optimized fed-batch process and ER-stress adaptation. <i>Journal of Biotechnology</i> , 2020, 311, 49-58.	1.9	6
9	ZnO Nanoparticles and Rifampicin Synergistically Damage the Membrane of Mycobacteria. <i>ACS Applied Nano Materials</i> , 2020, 3, 3174-3184.	2.4	10
10	Repurposing artemisinin as an anti-mycobacterial agent in synergy with rifampicin. <i>Tuberculosis</i> , 2019, 115, 146-153.	0.8	20
11	Targeting wild-type and drug-resistant mycobacteria in infected macrophages using drug-coated nanoparticles. <i>Journal of Chemical Technology and Biotechnology</i> , 2019, 94, 768-776.	1.6	6
12	Distinct transcriptomic response of <i>S. coelicolor</i> to ciprofloxacin in a nutrient-rich environment. <i>Applied Microbiology and Biotechnology</i> , 2018, 102, 10623-10643.	1.7	4
13	The Familial Î±-Synuclein A53E Mutation Enhances Cell Death in Response to Environmental Toxins Due to a Larger Population of Oligomers. <i>Biochemistry</i> , 2018, 57, 5014-5028.	1.2	19
14	Activation of unfolded protein response pathway is important for valproic acid mediated increase in immunoglobulin G productivity in recombinant Chinese hamster ovary cells. <i>Journal of Bioscience and Bioengineering</i> , 2017, 124, 459-468.	1.1	10
15	Synergistic Response of Rifampicin with Hydroperoxides on <i>Mycobacterium</i> : A Mechanistic Study. <i>Frontiers in Microbiology</i> , 2017, 8, 2075.	1.5	15
16	Biocompatible citric acid-coated iron oxide nanoparticles to enhance the activity of first-line anti-TB drugs in <i>Mycobacterium smegmatis</i> . <i>Journal of Chemical Technology and Biotechnology</i> , 2015, 90, 1773-1781.	1.6	25
17	Dynamics of unfolded protein response in recombinant CHO cells. <i>Cytotechnology</i> , 2015, 67, 237-254.	0.7	44
18	Proteomic analysis of <i>Streptomyces coelicolor</i> in response to Ciprofloxacin challenge. <i>Journal of Proteomics</i> , 2014, 97, 222-234.	1.2	10

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19	Polyacrylic Acid-Coated Iron Oxide Nanoparticles for Targeting Drug Resistance in Mycobacteria. <i>Langmuir</i> , 2014, 30, 15266-15276.	1.6	76
20	Transcriptomic study of ciprofloxacin resistance in <i>Streptomyces coelicolor</i> A3(2). <i>Molecular BioSystems</i> , 2013, 9, 3101.	2.9	30
21	Comprehensive Phylogenetic Analysis of Mycobacteria. IFAC Postprint Volumes IPPV / International Federation of Automatic Control, 2013, 46, 101-106.	0.4	0
22	Comparative Phylogenomics of Pathogenic and Non-Pathogenic Mycobacterium. <i>PLoS ONE</i> , 2013, 8, e71248.	1.1	41
23	All-trans retinoic acid loaded block copolymer nanoparticles efficiently induce cellular differentiation in HL-60 cells. <i>European Journal of Pharmaceutical Sciences</i> , 2011, 44, 643-652.	1.9	11
24	Convergent Transcription in the Butyrolactone Regulon in <i>Streptomyces coelicolor</i> Confers a Bistable Genetic Switch for Antibiotic Biosynthesis. <i>PLoS ONE</i> , 2011, 6, e21974.	1.1	20
25	Genome-wide transcriptome analysis reveals that a pleiotropic antibiotic regulator, AfsS, modulates nutritional stress response in <i>Streptomyces coelicolor</i> A3(2). <i>BMC Genomics</i> , 2008, 9, 56.	1.2	48
26	A Bistable Gene Switch for Antibiotic Biosynthesis: The Butyrolactone Regulon in <i>Streptomyces coelicolor</i> . <i>PLoS ONE</i> , 2008, 3, e2724.	1.1	47
27	Transcriptome dynamics-based operon prediction and verification in <i>Streptomyces coelicolor</i> . <i>Nucleic Acids Research</i> , 2007, 35, 7222-7236.	6.5	30
28	TimeView. <i>Applied Bioinformatics</i> , 2006, 5, 41-44.	1.7	3
29	A framework to analyze multiple time series data: A case study with <i>Streptomyces coelicolor</i> . <i>Journal of Industrial Microbiology and Biotechnology</i> , 2006, 33, 159-172.	1.4	36
30	On-line adaptation of neural networks for bioprocess control. <i>Computers and Chemical Engineering</i> , 2005, 29, 1047-1057.	2.0	46
31	A kinetic model of quantitative real-time polymerase chain reaction. <i>Biotechnology and Bioengineering</i> , 2005, 91, 848-860.	1.7	40
32	A Boolean algorithm for reconstructing the structure of regulatory networks. <i>Metabolic Engineering</i> , 2004, 6, 326-339.	3.6	34