Yufan Chen

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9722281/publications.pdf

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11	202	7	11
papers	citations	h-index	g-index
11	11	11	146
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	The complete genome sequence of Dickeya zeae EC1 reveals substantial divergence from other Dickeya strains and species. BMC Genomics, 2015, 16, 571.	2.8	47
2	Fis is a global regulator critical for modulation of virulence factor production and pathogenicity of Dickeya zeae. Scientific Reports, 2018, 8, 341.	3.3	38
3	Production of Novel Antibiotics Zeamines through Optimizing Dickeya zeae Fermentation Conditions. PLoS ONE, 2014, 9, e116047.	2.5	24
4	Genetic Modulation of c-di-GMP Turnover Affects Multiple Virulence Traits and Bacterial Virulence in Rice Pathogen Dickeya zeae. PLoS ONE, 2016, 11, e0165979.	2.5	19
5	Systematic Analysis of c-di-GMP Signaling Mechanisms and Biological Functions in Dickeya zeae EC1. MBio, 2020, 11, .	4.1	18
6	Pseudomonas sp. ST 4 produces variety of active compounds to interfere fungal sexual mating and hyphal growth. Microbial Biotechnology, 2020, 13, 107-117.	4.2	14
7	The Roles of Microbial Cell-Cell Chemical Communication Systems in the Modulation of Antimicrobial Resistance. Antibiotics, 2020, 9, 779.	3.7	14
8	Isolation, Characterization, and Genomic Investigation of a Phytopathogenic Strain of <i>Stenotrophomonas maltophilia </i> Phytopathology, 2021, 111, 2088-2099.	2.2	8
9	Cyclic diâ€GMP modulates sessileâ€motile phenotypes and virulence in <i>Dickeya oryzae</i> via two PilZ domain receptors. Molecular Plant Pathology, 2022, 23, 870-884.	4.2	8
10	OhrR is a central transcriptional regulator of virulence in <i>Dickeya zeae</i> . Molecular Plant Pathology, 2022, 23, 45-59.	4.2	7
11	The GacA-GacS Type Two-Component System Modulates the Pathogenicity of <i>Dickeya oryzae</i> EC1 Mainly by Regulating the Production of Zeamines. Molecular Plant-Microbe Interactions, 2022, 35, 369-379.	2.6	5