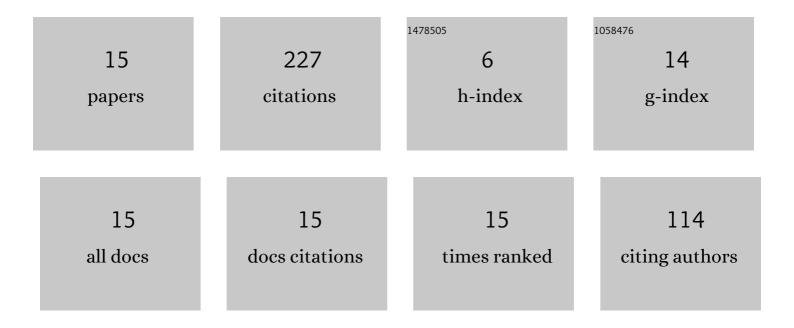
Ming Hu

List of Publications by Year in descending order

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MINC HU

#	Article	lF	CITATIONS
1	Dickeya zeae strains isolated from rice, banana and clivia rot plants show great virulence differentials. BMC Microbiology, 2018, 18, 136.	3.3	43
2	A twoâ€component regulatory system VfmIH modulates multiple virulence traits inDickeya zeae. Molecular Microbiology, 2019, 111, 1493-1509.	2.5	32
3	Screening, Identification and Efficacy Evaluation of Antagonistic Bacteria for Biocontrol of Soft Rot Disease Caused by Dickeya zeae. Microorganisms, 2020, 8, 697.	3.6	29
4	<i>Enterobacter asburiae</i> and <i>Pantoea ananatis</i> Causing Rice Bacterial Blight in China. Plant Disease, 2021, 105, 2078-2088.	1.4	23
5	Five Fungal Pathogens Are Responsible for Bayberry Twig Blight and Fungicides Were Screened for Disease Control. Microorganisms, 2020, 8, 689.	3.6	17
6	Virulence Factor Identification in the Banana Pathogen Dickeya zeae MS2. Applied and Environmental Microbiology, 2019, 85, .	3.1	15
7	Dickeya Manipulates Multiple Quorum Sensing Systems to Control Virulence and Collective Behaviors. Frontiers in Plant Science, 2022, 13, 838125.	3.6	10
8	lsolation, Characterization, and Genomic Investigation of a Phytopathogenic Strain of <i>Stenotrophomonas maltophilia</i> . Phytopathology, 2021, 111, 2088-2099.	2.2	8
9	Microbial Diversity Analysis and Genome Sequencing Identify Xanthomonas perforans as the Pathogen of Bacterial Leaf Canker of Water Spinach (Ipomoea aquatic). Frontiers in Microbiology, 2021, 12, 752760.	3.5	8
10	Genomic and Functional Dissections of Dickeya zeae Shed Light on the Role of Type III Secretion System and Cell Wall-Degrading Enzymes to Host Range and Virulence. Microbiology Spectrum, 2022, 10, e0159021.	3.0	8
11	Isolation and Genome Analysis of Pectobacterium colocasium sp. nov. and Pectobacterium aroidearum, Two New Pathogens of Taro. Frontiers in Plant Science, 2022, 13, 852750.	3.6	8
12	First Report of Bacterial Soft Rot Disease on Taro Caused by <i>Dickeya fangzhongdai</i> in China. Plant Disease, 2021, 105, 3737.	1.4	7
13	OhrR is a central transcriptional regulator of virulence in <i>Dickeya zeae</i> . Molecular Plant Pathology, 2022, 23, 45-59.	4.2	7
14	Five Plant Natural Products Are Potential Type III Secretion System Inhibitors to Effectively Control Soft-Rot Disease Caused by Dickeya. Frontiers in Microbiology, 2022, 13, 839025.	3.5	7
15	The integration host factor regulates multiple virulence pathways in bacterial pathogen <i>Dickeya zeae</i> MS2. Molecular Plant Pathology, 2022, 23, 1487-1507.	4.2	5