

Zheng Zheng

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	Historical Perspectives, Management, and Current Research of Citrus HLB in Guangdong Province of China, Where the Disease has been Endemic for Over a Hundred Years. <i>Phytopathology</i> , 2018, 108, 1224-1236.	2.2	56
2	Detection of Citrus Huanglongbing Based on Multi-Input Neural Network Model of UAV Hyperspectral Remote Sensing. <i>Remote Sensing</i> , 2020, 12, 2678.	4.0	47
3	A Type 3 Prophage of <i>Candidatus Liberibacter asiaticus</i> ™ Carrying a Restriction-Modification System. <i>Phytopathology</i> , 2018, 108, 454-461.	2.2	46
4	Predominance of Single Prophage Carrying a CRISPR/cas System in <i>Candidatus Liberibacter asiaticus</i> Strains in Southern China. <i>PLoS ONE</i> , 2016, 11, e0146422.	2.5	38
5	Enhancing PCR Capacity To Detect <i>Candidatus Liberibacter asiaticus</i> ™ Utilizing Whole Genome Sequence Information. <i>Plant Disease</i> , 2020, 104, 527-532.	1.4	37
6	Draft Genome Sequence of Rice Orange Leaf Phytoplasma from Guangdong, China. <i>Genome Announcements</i> , 2017, 5, .	0.8	22
7	Draft Whole-Genome Sequence of <i>Candidatus Liberibacter asiaticus</i> Strain TX2351 Isolated from Asian Citrus Psyllids in Texas, USA. <i>Genome Announcements</i> , 2017, 5, .	0.8	16
8	Analyses of Mitogenome Sequences Revealed that Asian Citrus Psyllids (<i>Diaphorina citri</i>) from California Were Related to Those from Florida. <i>Scientific Reports</i> , 2017, 7, 10154.	3.3	13
9	Establishment of a <i>Cuscuta campestris</i> -mediated enrichment system for genomic and transcriptomic analyses of <i>Candidatus Liberibacter asiaticus</i> ™. <i>Microbial Biotechnology</i> , 2021, 14, 737-751.	4.2	13
10	Prophage region and short tandem repeats of <i>Candidatus Liberibacter asiaticus</i> reveal significant population structure in China. <i>Plant Pathology</i> , 2021, 70, 959-969.	2.4	9
11	A Novel Microviridae Phage (CLasMV1) From <i>Candidatus Liberibacter asiaticus</i> Frontiers in Microbiology, 2021, 12, 754245.	3.5	9
12	Genome Sequence Resource of <i>Candidatus Liberibacter asiaticus</i> ™ from Thailand. <i>Plant Disease</i> , 2020, 104, 624-626.	1.4	7
13	The Complete Mitochondrial Genome Sequence of <i>Bactericera cockerelli</i> and Comparison with Three Other Psylloidea Species. <i>PLoS ONE</i> , 2016, 11, e0155318.	2.5	7
14	Detection of a Single-Copy Plasmid, pXFSL21, in <i>Xylella fastidiosa</i> Strain Stag's Leap with Two Toxin-Antitoxin Systems Using Next-Generation Sequencing. <i>Phytopathology</i> , 2019, 109, 240-247.	2.2	5
15	Draft Whole-Genome Sequence of a <i>Candidatus Liberibacter asiaticus</i> Strain from Yunnan, China. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.6	2
16	Genome Sequence Resource of <i>Candidatus Liberibacter asiaticus</i> ™ Strain Myan16 from Myanmar. <i>Plant Disease</i> , 2021, 105, 1162-1164.	1.4	2
17	Genome Sequence Resource of <i>Xanthomonas citri</i> pv. <i>citri</i> from Formalin-Fixed Citrus Leaves Specimen Showing Canker Lesions Collected in 1982. <i>Plant Disease</i> , 2022, 106, 304-306.	1.4	0