

Lifei Huang

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3404642/publications.pdf>

Version: 2024-02-01

11
papers

96
citations

1937685

4
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1372567

10
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all docs

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docs citations

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#	ARTICLE	IF	CITATIONS
1	Genome Resource for <i>Elsinoë batatas</i> , the Causal Agent of Stem and Foliage Scab Disease of Sweet Potato. <i>Phytopathology</i> , 2022, 112, 973-975.	2.2	4
2	Exploration of molecular mechanism of intraspecific cross-incompatibility in sweetpotato by transcriptome and metabolome analysis. <i>Plant Molecular Biology</i> , 2022, 109, 115-133.	3.9	3
3	First Report of White Rust on Sweetpotato Caused by <i>Albugo ipomoeae-panduratae</i> in Guangdong, China. <i>Plant Disease</i> , 2022, , .	1.4	0
4	The mitochondrial genome of the endophyte <i>Edenia gomezpompae</i> CRI Eg3 isolated from sweet potato. <i>Mitochondrial DNA Part B: Resources</i> , 2022, 7, 454-455.	0.4	1
5	Whole-Genome Sequencing and Comparative Genome Analysis of <i>Fusarium solani-melongenae</i> Causing Fusarium Root and Stem Rot in Sweetpotatoes. <i>Microbiology Spectrum</i> , 2022, 10, .	3.0	7
6	Complete chloroplast genome of a novel chlorophyll-deficient mutant (<i>clm</i>) in sweetpotato (<i>Ipomoea batatas</i> L.). <i>Mitochondrial DNA Part B: Resources</i> , 2021, 6, 968-969.	0.4	4
7	High-Quality Genome Resource of <i>Diaporthe destruens</i> Causing Foot Rot Disease of Sweet Potato. <i>Plant Disease</i> , 2021, 105, 3279-3281.	1.4	4
8	<i>Burkholderia gladioli</i> causes bacterial internal browning in sweetpotato of China. <i>Australasian Plant Pathology</i> , 2020, 49, 191-199.	1.0	1
9	Effect of domestic cooking methods on the anthocyanins and antioxidant activity of deeply purple-fleshed sweetpotato GZ9. <i>Heliyon</i> , 2019, 5, e01515.	3.2	24
10	Temporal patterns of gene expression associated with tuberous root formation and development in sweetpotato (<i>Ipomoea batatas</i>). <i>BMC Plant Biology</i> , 2015, 15, 180.	3.6	31
11	First Report of Bacterial Stem and Root Rot of Sweetpotato Caused by a <i>Dickeya</i> sp. (<i>Erwinia</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 10 T	1.4	17