

ZhenChen Zhang

List of Publications by Year in descending order

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docs citations

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57
citing authors

#	ARTICLE	IF	CITATIONS
1	Calystegia hederacea: a new natural host of Sweet potato latent virus in China. Plant Disease, 2022, , .	1.4	0
2	Complete genome sequence of a divergent sweet potato chlorotic stunt virus isolate infecting Calystegia hederacea in China. Archives of Virology, 2021, 166, 2037-2040.	2.1	3
3	Species and genetic variability of sweet potato viruses in China. Phytopathology Research, 2021, 3, .	2.4	8
4	Complete genome sequence of a novel varicosavirus infecting tall morning glory (Ipomoea purpurea). Archives of Virology, 2021, 166, 3225-3228.	2.1	3
5	Evidence for seed transmission of sweet potato symptomless virus 1 in sweet potato (Ipomoea batatas). Journal of Plant Pathology, 2020, 102, 299-303.	1.2	6
6	Seed transmission of sweet potato pakakuy virus in sweet potato (Ipomoea batatas). Journal of General Plant Pathology, 2020, 86, 205-210.	1.0	2
7	First report of sweet potato feathery mottle virus infecting Amaranthus blitum in China. Journal of Plant Pathology, 2020, 102, 965-965.	1.2	3
8	Diversity of Sweepoviruses Infecting Sweet Potato in China. Plant Disease, 2017, 101, 2098-2103.	1.4	7
9	Complete genome sequence of a novel monopartite begomovirus infecting sweet potato in China. Archives of Virology, 2014, 159, 1537-1540.	2.1	6
10	Complete Genomic Sequence and Comparative Analysis of the Genome Segments of Sweet Potato Chlorotic Stunt Virus in China. PLoS ONE, 2014, 9, e106323.	2.5	15
11	Complete genome sequence of a novel monopartite begomovirus infecting sweet potato in China. Virus Genes, 2013, 47, 591-594.	1.6	5
12	Molecular variability of sweet potato chlorotic stunt virus (SPCSV) and five potyviruses infecting sweet potato in China. Archives of Virology, 2013, 158, 491-495.	2.1	13
13	Complete Genome Sequences of Two Sweet Potato Chlorotic Stunt Virus Isolates from China. Genome Announcements, 2013, 1, .	0.8	4