

Kaili Zhong

List of Publications by Year in descending order

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Version: 2024-02-01

13
papers

377
citations

933447

10
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1125743

13
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14
all docs

14
docs citations

14
times ranked

823
citing authors

#	ARTICLE	IF	CITATIONS
1	Histone acetyltransferase MoHat1 acetylates autophagy-related proteins MoAtg3 and MoAtg9 to orchestrate functional appressorium formation and pathogenicity in <i>Magnaporthe oryzae</i> . <i>Autophagy</i> , 2019, 15, 1234-1257.	9.1	69
2	MoDnm1 Dynamin Mediating Peroxisomal and Mitochondrial Fission in Complex with MoFis1 and MoMdv1 Is Important for Development of Functional Appressorium in <i>Magnaporthe oryzae</i> . <i>PLoS Pathogens</i> , 2016, 12, e1005823.	4.7	62
3	<i>Rice black-streaked dwarf virus</i> -encoded P5 ϵ 1 regulates the ubiquitination activity of SCF E3 ligases and inhibits jasmonate signaling to benefit its infection in rice. <i>New Phytologist</i> , 2020, 225, 896-912.	7.3	59
4	A virus-derived siRNA activates plant immunity by interfering with ROS scavenging. <i>Molecular Plant</i> , 2021, 14, 1088-1103.	8.3	33
5	Wheat Yellow Mosaic Virus Nib Interacting with Host Light Induced Protein (LIP) Facilitates Its Infection through Perturbing the Abscisic Acid Pathway in Wheat. <i>Biology</i> , 2019, 8, 80.	2.8	28
6	The seven transmembrane domain protein MoRgs7 functions in surface perception and undergoes coronin MoCrn1-dependent endocytosis in complex with Gl ϵ subunit MoMagA to promote cAMP signaling and appressorium formation in <i>Magnaporthe oryzae</i> . <i>PLoS Pathogens</i> , 2019, 15, e1007382.	4.7	28
7	Genome-wide identification of the histone acetyltransferase gene family in <i>Triticum aestivum</i> . <i>BMC Genomics</i> , 2021, 22, 49.	2.8	22
8	Construction and biological characterization of an infectious full-length cDNA clone of a Chinese isolate of Wheat yellow mosaic virus. <i>Virology</i> , 2021, 556, 101-109.	2.4	22
9	NbWRKY40 Positively Regulates the Response of <i>Nicotiana benthamiana</i> to Tomato Mosaic Virus via Salicylic Acid Signaling. <i>Frontiers in Plant Science</i> , 2020, 11, 603518.	3.6	18
10	System-Wide Characterization of MoArf GTPase Family Proteins and Adaptor Protein MoGga1 Involved in the Development and Pathogenicity of <i>Magnaporthe oryzae</i> . <i>MBio</i> , 2019, 10, .	4.1	14
11	The inhibitor of apoptosis protein MoBir1 is involved in the suppression of hydrogen peroxide-induced fungal cell death, reactive oxygen species generation, and pathogenicity of rice blast fungus. <i>Applied Microbiology and Biotechnology</i> , 2019, 103, 6617-6627.	3.6	8
12	Comprehensive Proteomic Analysis of Lysine Acetylation in <i>Nicotiana benthamiana</i> After Sensing CWMV Infection. <i>Frontiers in Microbiology</i> , 2021, 12, 672559.	3.5	7
13	Genome-Wide Identification and Characterization of the Cystatin Gene Family in Bread Wheat (<i>Triticum</i>) Tj ETQq1 1 0.784314 rgBT / 0	4.1	5