

Xiao-Ren Chen

List of Publications by Year in descending order

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papers

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#	ARTICLE	IF	CITATIONS
1	A Small Cysteine-Rich Phytotoxic Protein of <i>Phytophthora capsici</i> Functions as Both Plant Defense Elicitor and Virulence Factor. <i>Molecular Plant-Microbe Interactions</i> , 2021, 34, 891-903.	1.4	14
2	Expression of the small cysteine-rich protein SCR96 from <i>Phytophthora cactorum</i> in mammalian cells: phytotoxicity and exploitation of its polyclonal antibody. <i>Biotechnology Letters</i> , 2020, 42, 125-133.	1.1	6
3	Identification and characterization of <i>Diaporthe</i> species causing leaf blight disease on the medicinal herb <i>Polygonatum sibiricum</i> . <i>Journal of General Plant Pathology</i> , 2020, 86, 468-476.	0.6	4
4	Investigation of the genetic diversity of <i>Phytophthora capsici</i> in China using a universal fluorescent labelling method. <i>Journal of Phytopathology</i> , 2019, 167, 111-122.	0.5	4
5	The RXLR Effector PcAvh1 Is Required for Full Virulence of <i>Phytophthora capsici</i> . <i>Molecular Plant-Microbe Interactions</i> , 2019, 32, 986-1000.	1.4	39
6	First Report of <i>Botrytis cinerea</i> Causing Leaf Spot of Chinese Quince in China. <i>Plant Disease</i> , 2019, 103, 1027.	0.7	2
7	Transcription profiling and identification of infection-related genes in <i>Phytophthora cactorum</i> . <i>Molecular Genetics and Genomics</i> , 2018, 293, 541-555.	1.0	23
8	Identification and functional analysis of the NLP-encoding genes from the phytopathogenic oomycete <i>Phytophthora capsici</i> . <i>Molecular Genetics and Genomics</i> , 2018, 293, 931-943.	1.0	35
9	The Plant Ribosome-Inactivating Proteins Play Important Roles in Defense against Pathogens and Insect Pest Attacks. <i>Frontiers in Plant Science</i> , 2018, 9, 146.	1.7	83
10	<i>Phytophthora sojae</i> . , 2017, , 199-223.		5
11	SCR96, a small cysteine-rich secretory protein of <i>Phytophthora cactorum</i> , can trigger cell death in the Solanaceae and is important for pathogenicity and oxidative stress tolerance. <i>Molecular Plant Pathology</i> , 2016, 17, 577-587.	2.0	42
12	Expression of resistance gene analogs in woodland strawberry (<i>Fragaria vesca</i>) during infection with <i>Phytophthora cactorum</i> . <i>Molecular Genetics and Genomics</i> , 2016, 291, 1967-1978.	1.0	23
13	Identification and characterization of <i>Phytophthora helicoides</i> causing stem rot of Shatangju mandarin seedlings in China. <i>European Journal of Plant Pathology</i> , 2016, 146, 715-727.	0.8	17
14	The Top 10 oomycete pathogens in molecular plant pathology. <i>Molecular Plant Pathology</i> , 2015, 16, 413-434.	2.0	695
15	Transcriptomic analysis of the phytopathogenic oomycete <i>Phytophthora cactorum</i> provides insights into infection-related effectors. <i>BMC Genomics</i> , 2014, 15, 980.	1.2	33
16	RNA-Seq Reveals Infection-Related Gene Expression Changes in <i>Phytophthora capsici</i> . <i>PLoS ONE</i> , 2013, 8, e74588.	1.1	49
17	First Report of <i>Pestalotiopsis sydowiana</i> Causing Leaf Necrosis of <i>Myrica rubra</i> in China. <i>Plant Disease</i> , 2012, 96, 764-764.	0.7	2
18	Identification and analysis of <i>Phytophthora cactorum</i> genes up-regulated during cyst germination and strawberry infection. <i>Current Genetics</i> , 2011, 57, 297-315.	0.8	12

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19	Green fluorescent protein (GFP) as a vital marker for studying the interaction of <i>Phytophthora sojae</i> and soybean. <i>Science Bulletin</i> , 2009, 54, 2822-2829.	4.3	2
20	The PsCZF1 gene encoding a C2H2 zinc finger protein is required for growth, development and pathogenesis in <i>Phytophthora sojae</i> . <i>Microbial Pathogenesis</i> , 2009, 47, 78-86.	1.3	40
21	Differences in the induction of the oxidative burst in compatible and incompatible interactions of soybean and <i>Phytophthora sojae</i> . <i>Physiological and Molecular Plant Pathology</i> , 2008, 73, 16-24.	1.3	18
22	Identification of <i>Phytophthora sojae</i> genes upregulated during the early stage of soybean infection. <i>FEMS Microbiology Letters</i> , 2007, 269, 280-288.	0.7	36
23	Differential screening reveals genes differentially expressed in low- and high-virulence near-isogenic <i>Phytophthora sojae</i> lines. <i>Fungal Genetics and Biology</i> , 2006, 43, 826-839.	0.9	13