Raksha Singh

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6385998/publications.pdf

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20 624 12 20 papers citations h-index 944

times ranked

citing authors

docs citations

all docs

#	Article	IF	CITATIONS
1	The Pseudomonas syringae type III effector HopG1 triggers necrotic cell death that is attenuated by AtNHR2B. Scientific Reports, 2022, 12, 5388.	3.3	7
2	Exploring the Corn Microbiome: A Detailed Review on Current Knowledge, Techniques, and Future Directions. PhytoFrontiers, 2022, 2, 158-175.	1.6	11
3	Recovery Plan for Tar Spot of Corn, Caused by <i>Phyllachora maydis</i> . Plant Health Progress, 2021, 22, 596-616.	1.4	22
4	Mitogen-Activated Protein Kinase OsMEK2 and OsMPK1 Signaling Is Required for Ferroptotic Cell Death in Rice–Magnaporthe oryzae Interactions. Frontiers in Plant Science, 2021, 12, 710794.	3 . 6	14
5	Draft Genome Sequence Resource for <i>Phyllachora maydis</i> ê"An Obligate Pathogen That Causes Tar Spot of Corn with Recent Economic Impacts in the United States. Molecular Plant-Microbe Interactions, 2020, 33, 884-887.	2.6	11
6	The Arabidopsis Proteins AtNHR2A and AtNHR2B Are Multi-Functional Proteins Integrating Plant Immunity With Other Biological Processes. Frontiers in Plant Science, 2020, 11, 232.	3.6	9
7	Exploring DNA variant segregation types in pooled genome sequencing enables effective mapping of weeping trait in Malus. Journal of Experimental Botany, 2018, 69, 1499-1516.	4.8	33
8	Dissecting the functional domains of the Arabidopsis thaliana nonhost resistance 2B (AtNHR2B) protein. Plant Signaling and Behavior, 2018, 13, e1530024.	2.4	8
9	Two Chloroplast-Localized Proteins: AtNHR2A and AtNHR2B, Contribute to Callose Deposition During Nonhost Disease Resistance in <i>Arabidopsis</i> Molecular Plant-Microbe Interactions, 2018, 31, 1280-1290.	2.6	22
10	Visualization of Multicolored in vivo Organelle Markers for Co-Localization Studies in Oryza sativa. Molecules and Cells, 2017, 40, 828-836.	2.6	26
11	Magnaporthe oryzae Effector AVR-Pii Helps to Establish Compatibility by Inhibition of the Rice NADP-Malic Enzyme Resulting in Disruption of Oxidative Burst and Host Innate Immunity. Molecules and Cells, 2016, 39, 426-438.	2.6	67
12	Protein interactome analysis of 12 mitogenâ€activated protein kinase kinase kinase in rice using a yeast twoâ€hybrid system. Proteomics, 2014, 14, 105-115.	2.2	14
13	Yeast Two-Hybrid System for Dissecting the Rice MAPK Interactome. Methods in Molecular Biology, 2014, 1171, 195-216.	0.9	5
14	Understanding the Responses of Rice to Environmental Stress Using Proteomics. Journal of Proteome Research, 2013, 12, 4652-4669.	3.7	63
15	Plant proteomics in India and Nepal: current status and challenges ahead. Physiology and Molecular Biology of Plants, 2013, 19, 461-477.	3.1	7
16	The rice MAPKK–MAPK interactome: the biological significance of MAPK components in hormone signal transduction. Plant Cell Reports, 2013, 32, 923-931.	5 . 6	68
17	Rice Mitogen-Activated Protein Kinase Interactome Analysis Using the Yeast Two-Hybrid System Â. Plant Physiology, 2012, 160, 477-487.	4.8	81
18	Secretome analysis of <i>Magnaporthe oryzae</i> using in vitro systems. Proteomics, 2012, 12, 878-900.	2.2	30

#	Article	IF	CITATIONS
19	Rice OsACDR1 (Oryza sativa Accelerated Cell Death and Resistance 1) Is a Potential Positive Regulator of Fungal Disease Resistance. Molecules and Cells, 2009, 28, 431-440.	2.6	67
20	Systematic Secretome Analyses of Rice Leaf and Seed Callus Suspension-Cultured Cells: Workflow Development and Establishment of High-Density Two-Dimensional Gel Reference Maps. Journal of Proteome Research, 2008, 7, 5187-5210.	3.7	58