Ronny Kellner

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4606433/publications.pdf Version: 2024-02-01

		759055	887953
16	1,182	12	17
papers	citations	h-index	g-index
23	23	23	1939
all docs	docs citations	times ranked	citing authors

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#	Article	IF	CITATIONS
1	Emergence of wheat blast in Bangladesh was caused by a South American lineage of Magnaporthe oryzae. BMC Biology, 2016, 14, 84.	1.7	355
2	ATG8 Expansion: A Driver of Selective Autophagy Diversification?. Trends in Plant Science, 2017, 22, 204-214.	4.3	129
3	Emerging oomycete threats to plants and animals. Philosophical Transactions of the Royal Society B: Biological Sciences, 2016, 371, 20150459.	1.8	114
4	Expression Profiling of the Wheat Pathogen Zymoseptoria tritici Reveals Genomic Patterns of Transcription and Host-Specific Regulatory Programs. Genome Biology and Evolution, 2014, 6, 1353-1365.	1.1	92
5	Nine things to know about elicitins. New Phytologist, 2016, 212, 888-895.	3.5	84
6	The biotechnological use and potential of plant pathogenic smut fungi. Applied Microbiology and Biotechnology, 2013, 97, 3253-3265.	1.7	78
7	Interspecific Sex in Grass Smuts and the Genetic Diversity of Their Pheromone-Receptor System. PLoS Genetics, 2011, 7, e1002436.	1.5	70
8	A complete toolset for the study of Ustilago bromivora and Brachypodium sp. as a fungal-temperate grass pathosystem. ELife, 2016, 5, .	2.8	49
9	The RNA-Binding Protein Rrm4 is Essential for Efficient Secretion of Endochitinase Cts1. Molecular and Cellular Proteomics, 2011, 10, M111.011213.	2.5	48
10	Two NLR immune receptors acquired high-affinity binding to a fungal effector through convergent evolution of their integrated domain. ELife, 2021, 10, .	2.8	38
11	Subfamily-Specific Specialization of RGH1/MLA Immune Receptors in Wild Barley. Molecular Plant-Microbe Interactions, 2019, 32, 107-119.	1.4	29
12	Identification of a new order of root-colonising fungi in the Entorrhizomycota: Talbotiomycetales ord. nov. on eudicotyledons. IMA Fungus, 2015, 6, 129-133.	1.7	14
13	The Plant-Dependent Life Cycle of <i>Thecaphora thlaspeos</i> : A Smut Fungus Adapted to Brassicaceae. Molecular Plant-Microbe Interactions, 2017, 30, 271-282.	1.4	13
14	Smut infection of perennial hosts: the genome and the transcriptome of the Brassicaceae smut fungus <i>Thecaphora thlaspeos</i> reveal functionally conserved and novel effectors. New Phytologist, 2019, 222, 1474-1492.	3.5	11
15	Patterns of Variation at Ustilago maydis Virulence Clusters 2A and 19A Largely Reflect the Demographic History of Its Populations. PLoS ONE, 2014, 9, e98837.	1.1	8
16	<i>Thecaphora dahuangis</i> , a new species causing leaf smut disease of the traditional medicinal plant <i>dahuang</i> (<i>Rheum palmatum</i>) in China. Plant Pathology, 2021, 70, 1292-1299.	1.2	2