Theo Van Der Lee

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11657762/publications.pdf

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26 papers 3,368 citations

361413 20 h-index 26 g-index

26 all docs

26 docs citations

times ranked

26

2950 citing authors

#	Article	IF	CITATIONS
1	Evaluation of Fusarium Head Blight Resistance in 410 Chinese Wheat Cultivars Selected for Their Climate Conditions and Ecological Niche Using Natural Infection Across Three Distinct Experimental Sites. Frontiers in Plant Science, 2022, 13, .	3.6	4
2	At the scene of the crime: New insights into the role of weakly pathogenic members of the fusarium head blight disease complex. Molecular Plant Pathology, 2020, 21, 1559-1572.	4.2	25
3	FgPex3, a Peroxisome Biogenesis Factor, Is Involved in Regulating Vegetative Growth, Conidiation, Sexual Development, and Virulence in Fusarium graminearum. Frontiers in Microbiology, 2019, 10, 2088.	3.5	19
4	Host and Cropping System Shape the Fusarium Population: 3ADON-Producers Are Ubiquitous in Wheat Whereas NIV-Producers Are More Prevalent in Rice. Toxins, 2018, 10, 115.	3.4	31
5	Targeting Trichothecene Biosynthetic Genes. Methods in Molecular Biology, 2017, 1542, 173-189.	0.9	7
6	Biogeography of <i>Fusarium graminearum </i> species complex and chemotypes: a review. Food Additives and Contaminants - Part A Chemistry, Analysis, Control, Exposure and Risk Assessment, 2015, 32, 453-460.	2.3	202
7	Increased Difficulties to Control Late Blight in Tunisia Are Caused by a Genetically Diverse <i>Phytophthora infestans</i> Population Next to the Clonal Lineage NA-01. Plant Disease, 2014, 98, 898-908.	1.4	17
8	Simultaneous real-time PCR detection of Fusarium asiaticum, F. ussurianum and F. vorosii, representing the Asian clade of the F. graminearum species complex. International Journal of Food Microbiology, 2013, 166, 148-154.	4.7	7
9	Efficient multiplex simple sequence repeat genotyping of the oomycete plant pathogen Phytophthora infestans. Journal of Microbiological Methods, 2013, 92, 316-322.	1.6	105
10	Population Analysis of the Fusarium graminearum Species Complex from Wheat in China Show a Shift to More Aggressive Isolates. PLoS ONE, 2012, 7, e31722.	2.5	180
11	The genetic basis for 3-ADON and 15-ADON trichothecene chemotypes in Fusarium. Fungal Genetics and Biology, 2011, 48, 485-495.	2.1	180
12	Geographic substructure of Fusarium asiaticum isolates collected from barley in China. European Journal of Plant Pathology, 2010, 127, 239-248.	1.7	17
13	<i>Phytophthora infestans</i> Isolates Lacking Class I <i>ipiO</i> Variants Are Virulent on <i>Rpi-blb1</i> Potato. Molecular Plant-Microbe Interactions, 2009, 22, 1535-1545.	2.6	118
14	High-Density Genetic Linkage Maps of Phytophthora infestans Reveal Trisomic Progeny and Chromosomal Rearrangements. Genetics, 2004, 167, 1643-1661.	2.9	57
15	Quantitative Detection of Fusarium Species in Wheat Using TaqMan. European Journal of Plant Pathology, 2004, 110, 481-494.	1.7	196
16	Synteny in Toxigenic Fusarium Species: The Fumonisin Gene Cluster and the Mating Type Region as Examples. European Journal of Plant Pathology, 2004, 110, 533-544.	1.7	66
17	Major Changes in Fusarium spp. in Wheat in the Netherlands. European Journal of Plant Pathology, 2003, 109, 743-754.	1.7	277
18	Chromosomal Deletion in Isolates of Phytophthora infestans Correlates with Virulence on R3, R10, and R11 Potato Lines. Molecular Plant-Microbe Interactions, 2001, 14, 1444-1452.	2.6	33

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19	Mapping of Avirulence Genes in Phytophthora infestans With Amplified Fragment Length Polymorphism Markers Selected by Bulked Segregant Analysis. Genetics, 2001, 157, 949-956.	2.9	84
20	The Detection of Nonhybrid, Trisomic, and Triploid Offspring in Sexual Progeny of a Mating of Phytophthora infestans. Fungal Genetics and Biology, 1999, 26, 198-208.	2.1	37
21	Dissection of the Fusarium I2 Gene Cluster in Tomato Reveals Six Homologs and One Active Gene Copy. Plant Cell, 1998, 10, 1055-1068.	6.6	332
22	Dissection of the Fusarium I2 Gene Cluster in Tomato Reveals Six Homologs and One Active Gene Copy. Plant Cell, 1998, 10, 1055.	6.6	38
23	Loss of Production of the Elicitor Protein INF1 in the Clonal Lineage US-1 of Phytophthora infestans. Phytopathology, 1998, 88, 1315-1323.	2.2	35
24	AFLP Linkage Map of the OomycetePhytophthora infestans. Fungal Genetics and Biology, 1997, 21, 278-291.	2.1	147
25	AFLP-Based Fine Mapping of theMloGene to a 30-kb DNA Segment of the Barley Genome. Genomics, 1997, 44, 61-70.	2.9	88
26	The Barley Mlo Gene: A Novel Control Element of Plant Pathogen Resistance. Cell, 1997, 88, 695-705.	28.9	1,066