

Feng He

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/485980/publications.pdf>

Version: 2024-02-01

8
papers

178
citations

1307594
7
h-index

1588992
8
g-index

9
all docs

9
docs citations

9
times ranked

159
citing authors

#	ARTICLE	IF	CITATIONS
1	A Virulence Essential CRN Effector of <i>Phytophthora capsici</i> Suppresses Host Defense and Induces Cell Death in Plant Nucleus. <i>PLoS ONE</i> , 2015, 10, e0127965.	2.5	45
2	Whole Genome Re-sequencing Reveals Natural Variation and Adaptive Evolution of <i>Phytophthora sojae</i> . <i>Frontiers in Microbiology</i> , 2019, 10, 2792.	3.5	39
3	The transcription factor VpCRZ1 is required for fruiting body formation and pathogenicity in <i>Valsa pyri</i> . <i>Microbial Pathogenesis</i> , 2016, 95, 101-110.	2.9	22
4	Transcriptomics Analysis of the Chinese Pear Pathotype of <i>Alternaria alternata</i> Gives Insights into Novel Mechanisms of HSAF Antifungal Activities. <i>International Journal of Molecular Sciences</i> , 2018, 19, 1841.	4.1	22
5	PsAAT3, an oomycete-specific aspartate aminotransferase, is required for full pathogenicity of the oomycete pathogen <i>Phytophthora sojae</i> . <i>Fungal Biology</i> , 2016, 120, 620-630.	2.5	20
6	The Fungal-Specific Transcription Factor VpFSTF1 Is Required for Virulence in <i>Valsa pyri</i> . <i>Frontiers in Microbiology</i> , 2019, 10, 2945.	3.5	14
7	Comparative transcriptomics of two <i>Valsa pyri</i> isolates uncover different strategies for virulence and growth. <i>Microbial Pathogenesis</i> , 2018, 123, 478-486.	2.9	13
8	Identification and characterization of a stem canker and twig dieback disease of pear caused by <i>Neofusicoccum parvum</i> in Chinese mainland. <i>Phytopathology Research</i> , 2022, 4, .	2.4	3