Feng He

List of Publications by Year in descending order

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

Version: 2024-02-01

		1307594	1588992	
8	178	7	8	
papers	citations	h-index	g-index	
9	9	9	159	
all docs	docs citations	times ranked	citing authors	

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