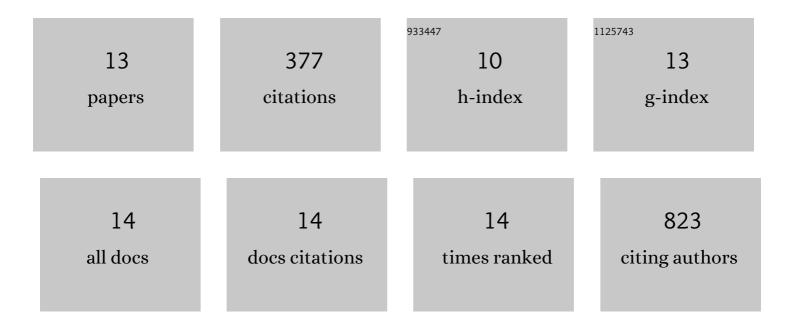
Kaili Zhong

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Genome-wide identification of the histone acetyltransferase gene family in Triticum aestivum. BMC Genomics, 2021, 22, 49.	2.8	22
2	Construction and biological characterization of an infectious full-length cDNA clone of a Chinese isolate of Wheat yellow mosaic virus. Virology, 2021, 556, 101-109.	2.4	22
3	Comprehensive Proteomic Analysis of Lysine Acetylation in Nicotiana benthamiana After Sensing CWMV Infection. Frontiers in Microbiology, 2021, 12, 672559.	3.5	7
4	A virus-derived siRNA activates plant immunity by interfering with ROS scavenging. Molecular Plant, 2021, 14, 1088-1103.	8.3	33
5	Genome-Wide Identification and Characterization of the Cystatin Gene Family in Bread Wheat (Triticum) Tj ETQq1	1 0.7843 4.1	14 rgBT /O
6	<i>Rice blackâ€streaked dwarf virus</i> â€encoded P5â€1 regulates the ubiquitination activity of SCF E3 ligases and inhibits jasmonate signaling to benefit its infection in rice. New Phytologist, 2020, 225, 896-912.	7.3	59
7	NbWRKY40 Positively Regulates the Response of Nicotiana benthamiana to Tomato Mosaic Virus via Salicylic Acid Signaling. Frontiers in Plant Science, 2020, 11, 603518.	3.6	18
8	Wheat Yellow Mosaic Virus NIb Interacting with Host Light Induced Protein (LIP) Facilitates Its Infection through Perturbing the Abscisic Acid Pathway in Wheat. Biology, 2019, 8, 80.	2.8	28
9	The inhibitor of apoptosis protein MoBir1 is involved in the suppression of hydrogen peroxide-induced fungal cell death, reactive oxygen species generation, and pathogenicity of rice blast fungus. Applied Microbiology and Biotechnology, 2019, 103, 6617-6627.	3.6	8
10	The seven transmembrane domain protein MoRgs7 functions in surface perception and undergoes coronin MoCrn1-dependent endocytosis in complex with Gα subunit MoMagA to promote cAMP signaling and appressorium formation in Magnaporthe oryzae. PLoS Pathogens, 2019, 15, e1007382.	4.7	28
11	Histone acetyltransferase MoHat1 acetylates autophagy-related proteins MoAtg3 and MoAtg9 to orchestrate functional appressorium formation and pathogenicity in <i>Magnaporthe oryzae</i> . Autophagy, 2019, 15, 1234-1257.	9.1	69
12	System-Wide Characterization of MoArf GTPase Family Proteins and Adaptor Protein MoGga1 Involved in the Development and Pathogenicity of Magnaporthe oryzae. MBio, 2019, 10, .	4.1	14
13	MoDnm1 Dynamin Mediating Peroxisomal and Mitochondrial Fission in Complex with MoFis1 and MoMdv1 Is Important for Development of Functional Appressorium in Magnaporthe oryzae. PLoS Pathogens, 2016, 12, e1005823.	4.7	62