## Zhuangzhi Zhou

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Functional Analysis of <i>Pid3-A4</i> , an Ortholog of Rice Blast Resistance Gene <i>Pid3</i> Revealed by Allele Mining in Common Wild Rice. Phytopathology, 2013, 103, 594-599.	2.2	59
2	Importance of OsRac1 and RAI1 in signalling of nucleotideâ€binding site leucineâ€rich repeat proteinâ€mediated resistance to rice blast disease. New Phytologist, 2019, 223, 828-838.	7.3	27
3	Excavation of Pid3 Orthologs with Differential Resistance Spectra to Magnaporthe oryzae in Rice Resource. PLoS ONE, 2014, 9, e93275.	2.5	23
4	Expression Profiles, Characterization and Function of HbTCTP in Rubber Tree (Hevea brasiliensis). Frontiers in Plant Science, 2016, 7, 789.	3.6	21
5	A Rice NBS-ARC Gene Conferring Quantitative Resistance to Bacterial Blight Is Regulated by a Pathogen Effector-Inducible miRNA. Molecular Plant, 2020, 13, 1752-1767.	8.3	20
6	Allelic variation of the rice blast resistance gene Pid3 in cultivated rice worldwide. Scientific Reports, 2017, 7, 10362.	3.3	19
7	Transgenic rice plants overexpressing BBTI4 confer partial but broad-spectrum bacterial blight resistance. Journal of Plant Biology, 2013, 56, 383-390.	2.1	15
8	Identification of a G2-like transcription factor, OsPHL3, functions as a negative regulator of flowering in rice by co-expression and reverse genetic analysis. BMC Plant Biology, 2018, 18, 157.	3.6	15
9	OsNPR3.3-dependent salicylic acid signaling is involved in recessive gene xa5-mediated immunity to rice bacterial blight. Scientific Reports, 2020, 10, 6313.	3.3	12
10	Endoplasmic reticulum membraneâ€bound MoSec62 is involved in the suppression of rice immunity and is essential for the pathogenicity of <i>Magnaporthe oryzae</i> . Molecular Plant Pathology, 2016, 17, 1211-1222.	4.2	6
11	The OsSPK1–OsRac1–RAI1 defense signaling pathway is shared by two distantly related NLR proteins in rice blast resistance. Plant Physiology, 2021, 187, 2852-2864	4.8	5