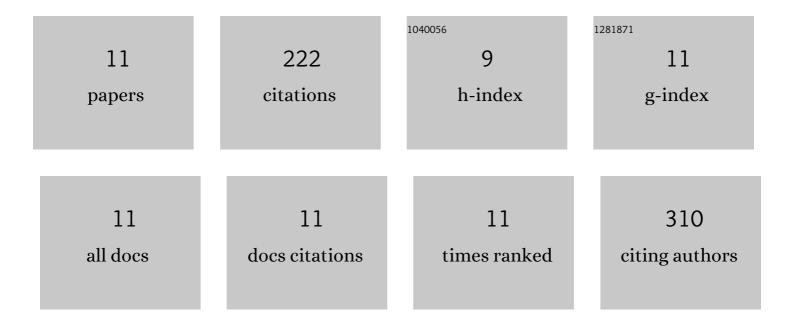
## Zhuangzhi Zhou

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4478159/publications.pdf Version: 2024-02-01



<u> 7нилистні 7ноц</u>

#	Article	IF	CITATIONS
1	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
2	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
3	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
4	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
5	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
6	Allelic variation of the rice blast resistance gene Pid3 in cultivated rice worldwide. Scientific Reports, 2017, 7, 10362.	3.3	19
7	Expression Profiles, Characterization and Function of HbTCTP in Rubber Tree (Hevea brasiliensis). Frontiers in Plant Science, 2016, 7, 789.	3.6	21
8	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
9	Excavation of Pid3 Orthologs with Differential Resistance Spectra to Magnaporthe oryzae in Rice Resource. PLoS ONE, 2014, 9, e93275.	2.5	23
10	Transgenic rice plants overexpressing BBTI4 confer partial but broad-spectrum bacterial blight resistance. Journal of Plant Biology, 2013, 56, 383-390.	2.1	15
11	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