

# Yifeng Wang

## List of Publications by Year in descending order

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33  
papers

1,350  
citations

489802

18  
h-index

466096

32  
g-index

33  
all docs

33  
docs citations

33  
times ranked

1651  
citing authors

#	ARTICLE	IF	CITATIONS
1	RLB (RICE LATERAL BRANCH) recruits PRC2-mediated H3K27 tri-methylation on <i>OsCKX4</i> to regulate lateral branching. <i>Plant Physiology</i> , 2022, 188, 460-476.	2.3	26
2	The <i>OsNAC23-Tre6P-SnRK1a</i> feed-forward loop regulates sugar homeostasis and grain yield in rice. <i>Molecular Plant</i> , 2022, 15, 706-722.	3.9	52
3	<i>WRKY72</i> Negatively Regulates Seed Germination Through Interfering Gibberellin Pathway in Rice. <i>Rice Science</i> , 2021, 28, 1-5.	1.7	7
4	Posttranslational Modification of <i>Waxy</i> to Genetically Improve Starch Quality in Rice Grain. <i>International Journal of Molecular Sciences</i> , 2021, 22, 4845.	1.8	13
5	<i>OsbZIP72</i> Is Involved in Transcriptional Gene-Regulation Pathway of Abscisic Acid Signal Transduction by Activating Rice High-Affinity Potassium Transporter <i>OsHKT1;1</i> . <i>Rice Science</i> , 2021, 28, 257-267.	1.7	16
6	<i>bZIP72</i> promotes submerged rice seed germination and coleoptile elongation by activating <i>ADH1</i> . <i>Plant Physiology and Biochemistry</i> , 2021, 169, 112-118.	2.8	8
7	<i>OsABF1</i> Represses Gibberellin Biosynthesis to Regulate Plant Height and Seed Germination in Rice ( <i>Oryza sativa</i> L.). <i>International Journal of Molecular Sciences</i> , 2021, 22, 12220.	1.8	9
8	Cyclin-Dependent Kinase Inhibitors <i>KRP1</i> and <i>KRP2</i> Are Involved in Grain Filling and Seed Germination in Rice ( <i>Oryza sativa</i> L.). <i>International Journal of Molecular Sciences</i> , 2020, 21, 245.	1.8	12
9	Genome-Wide Identification of lncRNAs During Rice Seed Development. <i>Genes</i> , 2020, 11, 243.	1.0	15
10	Abscisic acid promotes jasmonic acid biosynthesis via a <i>SAPK10</i> - <i>bZIP72</i> - <i>AOC</i> ™ pathway to synergistically inhibit seed germination in rice ( <i>Oryza sativa</i> ). <i>New Phytologist</i> , 2020, 228, 1336-1353.	3.5	93
11	<i>RMS2</i> Encoding a GDSL Lipase Mediates Lipid Homeostasis in Anthers to Determine Rice Male Fertility. <i>Plant Physiology</i> , 2020, 182, 2047-2064.	2.3	57
12	<i>SAPK10</i> -Mediated Phosphorylation on <i>WRKY72</i> Releases Its Suppression on Jasmonic Acid Biosynthesis and Bacterial Blight Resistance. <i>IScience</i> , 2019, 16, 499-510.	1.9	51
13	Protein Interactomic Analysis of SAPKs and ABA-Inducible <i>bZIPs</i> Revealed Key Roles of <i>SAPK10</i> in Rice Flowering. <i>International Journal of Molecular Sciences</i> , 2019, 20, 1427.	1.8	18
14	<i>NFYB1</i> - <i>YC12</i> - <i>bHLH144</i> complex directly activates <i>Wx</i> to regulate grain quality in rice ( <i>Oryza sativa</i> L.). <i>Plant Biotechnology Journal</i> , 2019, 17, 1222-1235.	4.1	103
15	Construction of a Quantitative Acetylomic Tissue Atlas in Rice ( <i>Oryza sativa</i> L.). <i>Molecules</i> , 2018, 23, 2843.	1.7	6
16	Notched Belly Grain 4, a Novel Allele of Dwarf 11, Regulates Grain Shape and Seed Germination in Rice ( <i>Oryza sativa</i> L.). <i>International Journal of Molecular Sciences</i> , 2018, 19, 4069.	1.8	16
17	<i>OsHAC4</i> is critical for arsenate tolerance and regulates arsenic accumulation in rice. <i>New Phytologist</i> , 2017, 215, 1090-1101.	3.5	156
18	A phosphoproteomic landscape of rice ( <i>Oryza sativa</i> ) tissues. <i>Physiologia Plantarum</i> , 2017, 160, 458-475.	2.6	28

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19	DNA demethylation activates genes in seed maternal integument development in rice ( <i>Oryza sativa</i> L.). <i>Plant Physiology and Biochemistry</i> , 2017, 120, 169-178.	2.8	6
20	A Quantitative Proteomic Analysis of Brassinosteroid-induced Protein Phosphorylation in Rice ( <i>Oryza</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 Tf	1.7	24
21	Transcriptomic Analysis of Gibberellin- and Paclobutrazol-Treated Rice Seedlings under Submergence. <i>International Journal of Molecular Sciences</i> , 2017, 18, 2225.	1.8	22
22	A Comprehensive Proteomic Survey of ABA-Induced Protein Phosphorylation in Rice ( <i>Oryza sativa</i> L.). <i>International Journal of Molecular Sciences</i> , 2017, 18, 60.	1.8	31
23	A Quantitative Acetylotomic Analysis of Early Seed Development in Rice ( <i>Oryza sativa</i> L.). <i>International Journal of Molecular Sciences</i> , 2017, 18, 1376.	1.8	20
24	Mapping the N-linked glycosites of rice ( <i>Oryza sativa</i> L.) germinating embryos. <i>PLoS ONE</i> , 2017, 12, e0173853.	1.1	15
25	Serine carboxypeptidase 46 Regulates Grain Filling and Seed Germination in Rice ( <i>Oryza sativa</i> L.). <i>PLoS ONE</i> , 2016, 11, e0159737.	1.1	47
26	Heterologous Expression and Functional Analysis of Rice GLUTAMATE RECEPTOR-LIKE Family Indicates its Role in Glutamate Triggered Calcium Flux in Rice Roots. <i>Rice</i> , 2016, 9, 9.	1.7	33
27	Data set from a comprehensive phosphoproteomic analysis of rice variety IRBB5 in response to bacterial blight. <i>Data in Brief</i> , 2016, 6, 282-285.	0.5	4
28	Quantitative phosphoproteomic analysis of early seed development in rice ( <i>Oryza sativa</i> L.). <i>Plant Molecular Biology</i> , 2016, 90, 249-265.	2.0	38
29	Aequorin-based luminescence imaging reveals differential calcium signalling responses to salt and reactive oxygen species in rice roots. <i>Journal of Experimental Botany</i> , 2015, 66, 2535-2545.	2.4	24
30	The Rice CK2 Kinase Regulates Trafficking of Phosphate Transporters in Response to Phosphate Levels. <i>Plant Cell</i> , 2015, 27, 711-723.	3.1	120
31	Phosphate transporters <scp><scp>OsPHT1</scp></scp>;9 and <scp><scp>OsPHT1</scp></scp>;10 are involved in phosphate uptake in rice. <i>Plant, Cell and Environment</i> , 2014, 37, 1159-1170.	2.8	135
32	OsPHF1 Regulates the Plasma Membrane Localization of Low- and High-Affinity Inorganic Phosphate Transporters and Determines Inorganic Phosphate Uptake and Translocation in Rice Å Å. <i>Plant Physiology</i> , 2011, 157, 269-278.	2.3	144
33	Chromosome-level genome assembly defines female-biased genes associated with sex determination and differentiation in the human blood fluke <i>Schistosoma japonicum</i>. <i>Molecular Ecology Resources</i> , 0, , .	2.2	1