Wanqi Liang

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91 4,890 38 69 g-index

97 6,391 8.3 5.33 L-index

#	Paper	IF	Citations
91	Genome-wide analysis of basic/helix-loop-helix transcription factor family in rice and Arabidopsis. <i>Plant Physiology</i> , 2006 , 141, 1167-84	6.6	396
90	Cytochrome P450 family member CYP704B2 catalyzes the {omega}-hydroxylation of fatty acids and is required for anther cutin biosynthesis and pollen exine formation in rice. <i>Plant Cell</i> , 2010 , 22, 173-90	11.6	265
89	The ABORTED MICROSPORES regulatory network is required for postmeiotic male reproductive development in Arabidopsis thaliana. <i>Plant Cell</i> , 2010 , 22, 91-107	11.6	235
88	EAT1 promotes tapetal cell death by regulating aspartic proteases during male reproductive development in rice. <i>Nature Communications</i> , 2013 , 4, 1445	17.4	201
87	OsC6, encoding a lipid transfer protein, is required for postmeiotic anther development in rice. <i>Plant Physiology</i> , 2010 , 154, 149-62	6.6	200
86	Rice MADS3 regulates ROS homeostasis during late anther development. <i>Plant Cell</i> , 2011 , 23, 515-33	11.6	191
85	PERSISTENT TAPETAL CELL1 encodes a PHD-finger protein that is required for tapetal cell death and pollen development in rice. <i>Plant Physiology</i> , 2011 , 156, 615-30	6.6	186
84	Defective pollen wall is required for anther and microspore development in rice and encodes a fatty acyl carrier protein reductase. <i>Plant Cell</i> , 2011 , 23, 2225-46	11.6	180
83	Carbon starved anther encodes a MYB domain protein that regulates sugar partitioning required for rice pollen development. <i>Plant Cell</i> , 2010 , 22, 672-89	11.6	174
82	Jasmonic acid regulates spikelet development in rice. <i>Nature Communications</i> , 2014 , 5, 3476	17.4	150
81	The SEPALLATA-like gene OsMADS34 is required for rice inflorescence and spikelet development. <i>Plant Physiology</i> , 2010 , 153, 728-40	6.6	148
80	ABORTED MICROSPORES Acts as a Master Regulator of Pollen Wall Formation in Arabidopsis. <i>Plant Cell</i> , 2014 , 26, 1544-1556	11.6	136
79	The Rice Basic Helix-Loop-Helix Transcription Factor TDR INTERACTING PROTEIN2 Is a Central Switch in Early Anther Development. <i>Plant Cell</i> , 2014 , 26, 1512-1524	11.6	120
78	The AGL6-like gene OsMADS6 regulates floral organ and meristem identities in rice. <i>Cell Research</i> , 2010 , 20, 299-313	24.7	108
77	Rice MADS6 interacts with the floral homeotic genes SUPERWOMAN1, MADS3, MADS58, MADS13, and DROOPING LEAF in specifying floral organ identities and meristem fate. <i>Plant Cell</i> , 2011 , 23, 2536-2	5 2 1.6	108
76	Rice CYP703A3, a cytochrome P450 hydroxylase, is essential for development of anther cuticle and pollen exine. <i>Journal of Integrative Plant Biology</i> , 2014 , 56, 979-94	8.3	107
75	Mutation in CSA creates a new photoperiod-sensitive genic male sterile line applicable for hybrid rice seed production. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 76-81	11.5	87

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74	Brassinosteroids promote development of rice pollen grains and seeds by triggering expression of Carbon Starved Anther, a MYB domain protein. <i>Plant Journal</i> , 2015 , 82, 570-81	6.9	78
73	Defective Tapetum Cell Death 1 (DTC1) Regulates ROS Levels by Binding to Metallothionein during Tapetum Degeneration. <i>Plant Physiology</i> , 2016 , 170, 1611-23	6.6	76
72	RICE MORPHOLOGY DETERMINANT encodes the type II formin FH5 and regulates rice morphogenesis. <i>Plant Cell</i> , 2011 , 23, 681-700	11.6	75
71	Identification of gamyb-4 and analysis of the regulatory role of GAMYB in rice anther development. <i>Journal of Integrative Plant Biology</i> , 2010 , 52, 670-8	8.3	72
70	Development of japonica Photo-Sensitive Genic Male Sterile Rice Lines by Editing Carbon Starved Anther Using CRISPR/Cas9. <i>Journal of Genetics and Genomics</i> , 2016 , 43, 415-9	4	69
69	Genetic interaction of OsMADS3, DROOPING LEAF, and OsMADS13 in specifying rice floral organ identities and meristem determinacy. <i>Plant Physiology</i> , 2011 , 156, 263-74	6.6	69
68	Dynamic changes of small RNAs in rice spikelet development reveal specialized reproductive phasiRNA pathways. <i>Journal of Experimental Botany</i> , 2016 , 67, 6037-6049	7	69
67	MTR1 encodes a secretory fasciclin glycoprotein required for male reproductive development in rice. <i>Developmental Cell</i> , 2012 , 22, 1127-37	10.2	67
66	Two ATP Binding Cassette G Transporters, Rice ATP Binding Cassette G26 and ATP Binding Cassette G15, Collaboratively Regulate Rice Male Reproduction. <i>Plant Physiology</i> , 2015 , 169, 2064-79	6.6	66
65	Defective Pollen Wall 2 (DPW2) Encodes an Acyl Transferase Required for Rice Pollen Development. <i>Plant Physiology</i> , 2017 , 173, 240-255	6.6	61
64	Rice auxin influx carrier OsAUX1 facilitates root hair elongation in response to low external phosphate. <i>Nature Communications</i> , 2018 , 9, 1408	17.4	61
63	Rice actin-binding protein RMD is a key link in the auxin-actin regulatory loop that controls cell growth. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 1037	77 ⁻¹ 82	58
62	Two rice receptor-like kinases maintain male fertility under changing temperatures. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017 , 114, 12327-12332	11.5	56
61	Interactions of OsMADS1 with Floral Homeotic Genes in Rice Flower Development. <i>Molecular Plant</i> , 2015 , 8, 1366-84	14.4	54
60	Post-meiotic deficient anther1 (PDA1) encodes an ABC transporter required for the development of anther cuticle and pollen exine in rice 2013 , 56, 59-68		50
59	Rice fatty acyl-CoA synthetase OsACOS12 is required for tapetum programmed cell death and male fertility. <i>Planta</i> , 2017 , 246, 105-122	4.7	42
58	Proteomic and phosphoproteomic analyses reveal extensive phosphorylation of regulatory proteins in developing rice anthers. <i>Plant Journal</i> , 2015 , 84, 527-44	6.9	41
57	MYB56 encoding a R2R3 MYB transcription factor regulates seed size in Arabidopsis thaliana. Journal of Integrative Plant Biology, 2013 , 55, 1166-78	8.3	41

56	Rice actin binding protein RMD controls crown root angle in response to external phosphate. <i>Nature Communications</i> , 2018 , 9, 2346	17.4	40
55	Dwarf Tiller1, a Wuschel-related homeobox transcription factor, is required for tiller growth in rice. <i>PLoS Genetics</i> , 2014 , 10, e1004154	6	40
54	A Rice Ca2+ Binding Protein Is Required for Tapetum Function and Pollen Formation. <i>Plant Physiology</i> , 2016 , 172, 1772-1786	6.6	40
53	Glycerol-3-Phosphate Acyltransferase 3 (OsGPAT3) is required for anther development and male fertility in rice. <i>Journal of Experimental Botany</i> , 2017 , 68, 513-526	7	38
52	Rice No Pollen 1 (NP1) is required for anther cuticle formation and pollen exine patterning. <i>Plant Journal</i> , 2017 , 91, 263-277	6.9	37
51	OsMADS16 genetically interacts with OsMADS3 and OsMADS58 in specifying floral patterning in rice. <i>Molecular Plant</i> , 2013 , 6, 743-56	14.4	36
50	MEIOTIC F-BOX Is Essential for Male Meiotic DNA Double-Strand Break Repair in Rice. <i>Plant Cell</i> , 2016 , 28, 1879-93	11.6	34
49	Loss of LOFSEP Transcription Factor Function Converts Spikelet to Leaf-Like Structures in Rice. <i>Plant Physiology</i> , 2018 , 176, 1646-1664	6.6	33
48	The polyketide synthase OsPKS2 is essential for pollen exine and Ubisch body patterning in rice. Journal of Integrative Plant Biology, 2017 , 59, 612-628	8.3	30
47	Dynamic Regulation of Auxin Response during Rice Development Revealed by Newly Established Hormone Biosensor Markers. <i>Frontiers in Plant Science</i> , 2017 , 8, 256	6.2	30
46	Regulatory Role of a Receptor-Like Kinase in Specifying Anther Cell Identity. <i>Plant Physiology</i> , 2016 , 171, 2085-100	6.6	30
45	PERSISTENT TAPETAL CELL2 Is Required for Normal Tapetal Programmed Cell Death and Pollen Wall Patterning. <i>Plant Physiology</i> , 2020 , 182, 962-976	6.6	24
44	Ostkpr1 functions in anther cuticle development and pollen wall formation in rice. <i>BMC Plant Biology</i> , 2019 , 19, 104	5.3	23
43	Postmeiotic development of pollen surface layers requires two Arabidopsis ABCG-type transporters. <i>Plant Cell Reports</i> , 2016 , 35, 1863-73	5.1	22
42	Defective Pollen Wall 3 (DPW3), a novel alpha integrin-like protein, is required for pollen wall formation in rice. <i>New Phytologist</i> , 2020 , 225, 807-822	9.8	21
41	The Post-meiotic Deficicent Anther1 (PDA1) gene is required for post-meiotic anther development in rice. <i>Journal of Genetics and Genomics</i> , 2010 , 37, 37-46	4	20
40	The DNA Topoisomerase VI-B Subunit OsMTOPVIB Is Essential for Meiotic Recombination Initiation in Rice. <i>Molecular Plant</i> , 2016 , 9, 1539-1541	14.4	19
39	Rice pollen aperture formation is regulated by the interplay between OsINP1 and OsDAF1. <i>Nature Plants</i> , 2020 , 6, 394-403	11.5	16

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38	A Rice Glutamyl-tRNA Synthetase Modulates Early Anther Cell Division and Patterning. <i>Plant Physiology</i> , 2018 , 177, 728-744	6.6	16
37	ATP binding cassette G transporters and plant male reproduction. <i>Plant Signaling and Behavior</i> , 2016 , 11, e1136764	2.5	16
36	Controls Flower Development by Activating Rice. Plant Physiology, 2018, 177, 713-727	6.6	15
35	Chromatin interacting factor OsVIL2 increases biomass and rice grain yield. <i>Plant Biotechnology Journal</i> , 2019 , 17, 178-187	11.6	15
34	Regulatory network and genetic interactions established by OsMADS34 in rice inflorescence and spikelet morphogenesis. <i>Journal of Integrative Plant Biology</i> , 2017 , 59, 693-707	8.3	15
33	Resolvase OsGEN1 Mediates DNA Repair by Homologous Recombination. <i>Plant Physiology</i> , 2017 , 173, 1316-1329	6.6	13
32	Interactions between FLORAL ORGAN NUMBER4 and floral homeotic genes in regulating rice flower development. <i>Journal of Experimental Botany</i> , 2017 , 68, 483-498	7	12
31	Genome-wide analysis of RopGEF gene family to identify genes contributing to pollen tube growth in rice (Oryza sativa). <i>BMC Plant Biology</i> , 2020 , 20, 95	5.3	11
30	Rice Morphology Determinant-Mediated Actin Filament Organization Contributes to Pollen Tube Growth. <i>Plant Physiology</i> , 2018 , 177, 255-270	6.6	11
29	Genome-wide analysis of the barley non-specific lipid transfer protein gene family. <i>Crop Journal</i> , 2019 , 7, 65-76	4.6	11
28	DWT1/DWL2 act together with OsPIP5K1 to regulate plant uniform growth in rice. <i>New Phytologist</i> , 2020 , 225, 1234-1246	9.8	10
27	Transcriptome profiling reveals phase-specific gene expression in the developing barley inflorescence. <i>Crop Journal</i> , 2020 , 8, 71-86	4.6	10
26	A Multiprotein Complex Regulates Interference-Sensitive Crossover Formation in Rice. <i>Plant Physiology</i> , 2019 , 181, 221-235	6.6	9
25	Oral immunization of mice with plant-derived fimbrial adhesin FaeG induces systemic and mucosal K88ad enterotoxigenic Escherichia coli-specific immune responses. <i>FEMS Immunology and Medical Microbiology</i> , 2006 , 46, 393-9		9
24	The Rice Actin-Binding Protein RMD Regulates Light-Dependent Shoot Gravitropism. <i>Plant Physiology</i> , 2019 , 181, 630-644	6.6	9
23	Regulates Spikelet Development by Controlling Regulatory Genes in. <i>Frontiers in Plant Science</i> , 2018 , 9, 102	6.2	8
22	Wheat AGAMOUS LIKE 6 transcription factors function in stamen development by regulating the expression of. <i>Development (Cambridge)</i> , 2019 , 146,	6.6	7
21	Grass-Specific Is Essential for Pollen Exine Patterning in Rice. <i>Plant Cell</i> , 2020 , 32, 3961-3977	11.6	7

20	MADS1 maintains barley spike morphology at high ambient temperatures. <i>Nature Plants</i> , 2021 , 7, 1093-	·1/11/0 3	7
19	Gibberellins orchestrate panicle architecture mediated by DELLA-KNOX signalling in rice. <i>Plant Biotechnology Journal</i> , 2021 , 19, 2304-2318	11.6	6
18	Transcriptome Analysis Reveals Photoperiod-Associated Genes Expressed in Rice Anthers. <i>Frontiers in Plant Science</i> , 2021 , 12, 621561	6.2	5
17	AUXIN RESPONSE FACTORS 6 and 17 control the flag leaf angle in rice by regulating secondary cell wall biosynthesis of lamina joints. <i>Plant Cell</i> , 2021 , 33, 3120-3133	11.6	5
16	Rice OsBRCA2 Is Required for DNA Double-Strand Break Repair in Meiotic Cells. <i>Frontiers in Plant Science</i> , 2020 , 11, 600820	6.2	4
15	NERD1 is required for primexine formation and plasma membrane undulation during microsporogenesis in Arabidopsis thaliana. <i>ABIOTECH</i> , 2020 , 1, 205-218	3.9	3
14	Two rice MYB transcription factors maintain male fertility in response to photoperiod by modulating sugar partitioning. <i>New Phytologist</i> , 2021 , 231, 1612-1629	9.8	3
13	HSP70-16 and VDAC3 jointly inhibit seed germination under cold stress in Arabidopsis. <i>Plant, Cell and Environment</i> , 2021 , 44, 3616-3627	8.4	3
12	Rice SEPALLATA genes OsMADS5 and OsMADS34 cooperate to limit inflorescence branching by repressing the TERMINAL FLOWER1-like gene RCN4. <i>New Phytologist</i> , 2021 ,	9.8	2
11	Integrating GWAS and transcriptomics to identify genes involved in seed dormancy in rice. <i>Theoretical and Applied Genetics</i> , 2021 , 134, 3553-3562	6	2
10	Function of the pseudo phosphotransfer proteins has diverged between rice and Arabidopsis. <i>Plant Journal</i> , 2021 , 106, 159-173	6.9	2
9	Bright Fluorescent Vacuolar Marker Lines Allow Vacuolar Tracing Across Multiple Tissues and Stress Conditions in Rice. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	1
8	Encodes a Type II Formin Required for Rice Morphogenesis <i>International Journal of Molecular Sciences</i> , 2021 , 22,	6.3	1
7	SMALL REPRODUCTIVE ORGANS, a SUPERMAN-like transcription factor, regulates stamen and pistil growth in rice. <i>New Phytologist</i> , 2021 , 233, 1701	9.8	1
6	Rice Glucose 6-Phosphate/Phosphate Translocator 1 is required for tapetum function and pollen development. <i>Crop Journal</i> , 2021 , 9, 1278-1278	4.6	1
5	Rice transcription factor MADS32 regulates floral patterning through interactions with multiple floral homeotic genes. <i>Journal of Experimental Botany,</i> 2021 , 72, 2434-2449	7	1
4	Carbon Starved Anther modulates sugar and ABA metabolism to protect rice seed germination and seedling fitness. <i>Plant Physiology</i> , 2021 , 187, 2405-2418	6.6	1
3	Rice SIAH E3 Ligases Interact with RMD Formin and Affect Plant Morphology <i>Rice</i> , 2022 , 15, 6	5.8	O

LIST OF PUBLICATIONS

Dissection of the Genetic Basis of Rice Panicle Architecture Using a Genome-wide Association Study. *Rice*, **2021**, 14, 77

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Automated High-Resolution Structure Analysis of Plant Root with a Morphological Image Filtering Algorithm. *Mathematical Problems in Engineering*, **2021**, 2021, 1-14

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