Zhigang Zhao

List of Publications by Year in descending order

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ZHICANC ZHAO

#	Article	IF	CITATIONS
1	Transcriptional activation and phosphorylation of OsCNGC9 confer enhanced chilling tolerance in rice. Molecular Plant, 2021, 14, 315-329.	8.3	89
2	Heterosis-associated genes confer high yield in super hybrid rice. Theoretical and Applied Genetics, 2020, 133, 3287-3297.	3.6	18
3	OsMFS1/OsHOP2 Complex Participates in Rice Male and Female Development. Frontiers in Plant Science, 2020, 11, 518.	3.6	10
4	A GARP transcription factor anther dehiscence defected 1 (OsADD1) regulates rice anther dehiscence. Plant Molecular Biology, 2019, 101, 403-414.	3.9	11
5	Identification of QTL for seed dormancy from weedy rice and its application to elite rice cultivar â€~Ninggeng 4'. Molecular Breeding, 2019, 39, 1.	2.1	8
6	Rice albino 1, encoding a glycyl-tRNA synthetase, is involved in chloroplast development and establishment of the plastidic ribosome system in rice. Plant Physiology and Biochemistry, 2019, 139, 495-503.	5.8	11
7	<i>Earlier Degraded Tapetum1</i> (<i>EDT1</i>) Encodes an ATP-Citrate Lyase Required for Tapetum Programmed Cell Death. Plant Physiology, 2019, 181, 1223-1238.	4.8	34
8	Genetic analysis and fine mapping of a dominant dwarfness gene from wild rice (<i>Oryza barthii</i>). Plant Breeding, 2018, 137, 50-59.	1.9	4
9	OPEN GLUME1: a key enzyme reducing the precursor of JA, participates in carbohydrate transport of lodicules during anthesis in rice. Plant Cell Reports, 2018, 37, 329-346.	5.6	23
10	A selfish genetic element confers non-Mendelian inheritance in rice. Science, 2018, 360, 1130-1132.	12.6	105
11	SGD1, a key enzyme in tocopherol biosynthesis, is essential for plant development and cold tolerance in rice. Plant Science, 2017, 260, 90-100.	3.6	26
12	Top Bending Panicle1 is involved in brassinosteroid signaling and regulates the plant architecture in rice. Plant Physiology and Biochemistry, 2017, 121, 1-13.	5.8	14
13	Genetic dissection of top three leaf traits in rice using progenies from a <i>japonica </i> × <i> indica</i> cross. Journal of Integrative Plant Biology, 2017, 59, 866-880.	8.5	15
14	The catalytic subunit of magnesium-protoporphyrin IX monomethyl ester cyclase forms a chloroplast complex to regulate chlorophyll biosynthesis in rice. Plant Molecular Biology, 2016, 92, 177-191.	3.9	47
15	Hybrid Sterility in Rice (<i>Oryza sativa</i> L.) Involves the Tetratricopeptide Repeat Domain Containing Protein. Genetics, 2016, 203, 1439-1451.	2.9	52
16	The role of OsMSH4 in male and female gamete development in rice meiosis. Journal of Experimental Botany, 2016, 67, 1447-1459.	4.8	43
17	Genetic dissection of leaf-related traits using 156 chromosomal segment substitution lines. Journal of Plant Biology, 2015, 58, 402-410.	2.1	6
18	Genetic dissection on rice grain shape by the two-dimensional image analysis in one japonicaÂ×Âindica population consisting of recombinant inbred lines. Theoretical and Applied Genetics, 2015, 128, 1969-1986.	3.6	63

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19	Fine mapping of S37, a locus responsible for pollen and embryo sac sterility in hybrids between Oryza sativa L. and O. glaberrima Steud. Plant Cell Reports, 2015, 34, 1885-1897.	5.6	14
20	A gene cluster encoding lectin receptor kinases confers broad-spectrum and durable insect resistance in rice. Nature Biotechnology, 2015, 33, 301-305.	17.5	299
21	Fine mapping of a minor-effect QTL, DTH12, controlling heading date in rice by up-regulation of florigen genes under long-day conditions. Molecular Breeding, 2014, 34, 311-322.	2.1	9
22	Identification of quantitative trait loci for seed storability in rice (<i>Oryza sativa</i> L.). Plant Breeding, 2012, 131, 739-743.	1.9	32
23	A new gene controlling hybrid sterility in rice (Oryza sativa L.). Euphytica, 2012, 184, 15-22.	1.2	5
24	Identification of QTLs for seed dormancy in rice (<i>Oryza sativa</i> L.). Plant Breeding, 2011, 130, 328-332.	1.9	36
25	Marker-assisted breeding of a photoperiod-sensitive male sterile japonica rice with high cross-compatibility with indica rice. Molecular Breeding, 2011, 27, 247-258.	2.1	25
26	Fine mapping of a gene responsible for pollen semi-sterility in hybrids between Oryza sativa L. and O. glaberrima Steud. Molecular Breeding, 2011, 28, 323-334.	2.1	9
27	Dynamic QTL Analysis of Rice Protein Content and Protein Index Using Recombinant Inbred Lines. Journal of Plant Biology, 2011, 54, 321-328.	2.1	35
28	<i>Pollen Semi-Sterility1</i> Encodes a Kinesin-1–Like Protein Important for Male Meiosis, Anther Dehiscence, and Fertility in Rice. Plant Cell, 2011, 23, 111-129.	6.6	113
29	Identification of <i>Japonica</i> Chromosome Segments Associated with Heterosis for Yield in <i>Indica</i> × <i>Japonica</i> Rice Hybrids. Crop Science, 2010, 50, 2328-2337.	1.8	10
30	Identification of a new hybrid sterility gene in rice (bi Oryza sativa L.). Euphytica, 2006, 151, 331-337.	1.2	23