

Yutaka Sato

List of Publications by Year in descending order

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Version: 2024-02-01

24
papers

1,189
citations

687363

13
h-index

642732

23
g-index

24
all docs

24
docs citations

24
times ranked

1519
citing authors

#	ARTICLE	IF	CITATIONS
1	RiceXPro Version 3.0: expanding the informatics resource for rice transcriptome. <i>Nucleic Acids Research</i> , 2013, 41, D1206-D1213.	14.5	312
2	Isolation and characterization of a rice WUSCHEL-type homeobox gene that is specifically expressed in the central cells of a quiescent center in the root apical meristem. <i>Plant Journal</i> , 2003, 35, 429-441.	5.7	231
3	A method for obtaining high quality RNA from paraffin sections of plant tissues by laser microdissection. <i>Journal of Plant Research</i> , 2010, 123, 807-813.	2.4	106
4	OslAA13-mediated auxin signaling is involved in lateral root initiation in rice. <i>Plant Science</i> , 2012, 190, 116-122.	3.6	103
5	Antagonistic regulation of the gibberellic acid response during stem growth in rice. <i>Nature</i> , 2020, 584, 109-114.	27.8	98
6	Position dependent expression of GL2-type homeobox gene, Roc1: significance for protoderm differentiation and radial pattern formation in early rice embryogenesis. <i>Plant Journal</i> , 2002, 29, 497-507.	5.7	78
7	Roles of Rice GL2-type Homeobox Genes in Epidermis Differentiation. <i>Breeding Science</i> , 2003, 53, 245-253.	1.9	27
8	Adaptive reduction of male gamete number in the selfing plant <i>Arabidopsis thaliana</i> . <i>Nature Communications</i> , 2020, 11, 2885.	12.8	27
9	WUSCHEL-related homeobox family genes in rice control lateral root primordium size. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	7.1	26
10	LEAF LATERAL SYMMETRY1, a Member of the WUSCHEL-RELATED HOMEBOX3 Gene Family, Regulates Lateral Organ Development Differentially from Other Paralogs, NARROW LEAF2 and NARROW LEAF3 in Rice. <i>Plant and Cell Physiology</i> , 2018, 59, 376-391.	3.1	25
11	Agrobacterium-Mediated Genetic Transformation of Wild <i>Oryza</i> Species Using Immature Embryos. <i>Rice</i> , 2020, 13, 33.	4.0	25
12	Rice SNF2 family helicase ENL1 is essential for syncytial endosperm development. <i>Plant Journal</i> , 2015, 81, 1-12.	5.7	24
13	Mutation of the imprinted gene OsEMF2a induces autonomous endosperm development and delayed cellularization in rice. <i>Plant Cell</i> , 2021, 33, 85-103.	6.6	23
14	Evolution and diversity of the wild rice <i>Oryza officinalis</i> complex, across continents genome types, and ploidy levels. <i>Genome Biology and Evolution</i> , 2020, 12, 413-428.	2.5	17
15	OryzaGenome2.1: Database of Diverse Genotypes in Wild <i>Oryza</i> Species. <i>Rice</i> , 2021, 14, 24.	4.0	17
16	Specification of the basal region identity after asymmetric zygotic division requires mitogen-activated protein kinase 6 in rice. <i>Development (Cambridge)</i> , 2019, 146, .	2.5	12
17	RNAi of the sesquiterpene cyclase gene for phytoalexin production impairs pre- and post-invasive resistance to potato blight pathogens. <i>Molecular Plant Pathology</i> , 2019, 20, 907-922.	4.2	10
18	Affinity-based high-resolution analysis of DNA binding by VASCULAR-RELATED NAC-DOMAIN7 via fluorescence correlation spectroscopy. <i>Plant Journal</i> , 2019, 100, 298-313.	5.7	8

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19	Collection, preservation and distribution of <i>Oryza</i> genetic resources by the National Bioresource Project RICE (NBRP-RICE). <i>Breeding Science</i> , 2021, 71, 291-298.	1.9	5
20	Temporal changes in transcripts of miniature invertedâ€repeat transposable elements during rice endosperm development. <i>Plant Journal</i> , 2022, 109, 1035-1047.	5.7	5
21	NARROW AND DWARF LEAF 1, the Ortholog of <i>Arabidopsis</i> ENHANCER OF SHOOT REGENERATION1/DORNRA-SCHEN, Mediates Leaf Development and Maintenance of the Shoot Apical Meristem in <i>Oryza sativa</i> L. <i>Plant and Cell Physiology</i> , 2022, 63, 265-278.	3.1	4
22	Measurements of the number of specified and unspecified cells in the shoot apical meristem during a plastochron in rice (<i>Oryza sativa</i>) reveal the robustness of cellular specification process in plant development. <i>PLoS ONE</i> , 2022, 17, e0269374.	2.5	4
23	High-resolution spatiotemporal transcriptome analyses during cellularization of rice endosperm unveil the earliest gene regulation critical for aleurone and starchy endosperm cell fate specification. <i>Journal of Plant Research</i> , 2021, 134, 1061-1081.	2.4	2
24	Perspectives on the use of bioresources in breeding sciences: Lessons from successful studies. <i>Ikushugaku Kenkyu</i> , 2019, 21, 81-85.	0.3	0