

Masayoshi Kawaguchi

List of Publications by Year in descending order

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papers

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all docs

140
docs citations

140
times ranked

5818
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome of an arbuscular mycorrhizal fungus provides insight into the oldest plant symbiosis. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 20117-20122.	7.1	717
2	HAR1 mediates systemic regulation of symbiotic organ development. Nature, 2002, 420, 426-429.	27.8	487
3	Deregulation of a Ca ²⁺ /calmodulin-dependent kinase leads to spontaneous nodule development. Nature, 2006, 441, 1153-1156.	27.8	400
4	CYCLOPS, a mediator of symbiotic intracellular accommodation. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 20540-20545.	7.1	398
5	Plastid proteins crucial for symbiotic fungal and bacterial entry into plant roots. Nature, 2005, 433, 527-531.	27.8	391
6	Nod Factor/Nitrate-Induced CLE Genes that Drive HAR1-Mediated Systemic Regulation of Nodulation. Plant and Cell Physiology, 2009, 50, 67-77.	3.1	342
7	NUCLEOPORIN85 Is Required for Calcium Spiking, Fungal and Bacterial Symbioses, and Seed Production in <i>Lotus japonicus</i> . Plant Cell, 2007, 19, 610-624.	6.6	309
8	Root-derived CLE glycopeptides control nodulation by direct binding to HAR1 receptor kinase. Nature Communications, 2013, 4, 2191.	12.8	292
9	The Sulfate Transporter SST1 Is Crucial for Symbiotic Nitrogen Fixation in <i>Lotus japonicus</i> Root Nodules. Plant Cell, 2005, 17, 1625-1636.	6.6	227
10	How Many Peas in a Pod? Legume Genes Responsible for Mutualistic Symbioses Underground. Plant and Cell Physiology, 2010, 51, 1381-1397.	3.1	227
11	<i>NENA</i> , a <i>Lotus japonicus</i> Homolog of <i>Sec13</i> , Is Required for Rhizodermal Infection by Arbuscular Mycorrhiza Fungi and Rhizobia but Dispensable for Cortical Endosymbiotic Development. Plant Cell, 2010, 22, 2509-2526.	6.6	215
12	Shoot-derived cytokinins systemically regulate root nodulation. Nature Communications, 2014, 5, 4983.	12.8	199
13	Positive and negative regulation of cortical cell division during root nodule development in <i>Lotus japonicus</i> is accompanied by auxin response. Development (Cambridge), 2012, 139, 3997-4006.	2.5	186
14	NODULE INCEPTION creates a long-distance negative feedback loop involved in homeostatic regulation of nodule organ production. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 14607-14612.	7.1	175
15	Long-distance signaling to control root nodule number. Current Opinion in Plant Biology, 2006, 9, 496-502.	7.1	169
16	Root, Root Hair, and Symbiotic Mutants of the Model Legume <i>Lotus japonicus</i> . Molecular Plant-Microbe Interactions, 2002, 15, 17-26.	2.6	150
17	A NIN-LIKE PROTEIN mediates nitrate-induced control of root nodule symbiosis in <i>Lotus japonicus</i> . Nature Communications, 2018, 9, 499.	12.8	144
18	RNA-seq Transcriptional Profiling of an Arbuscular Mycorrhiza Provides Insights into Regulated and Coordinated Gene Expression in <i>Lotus japonicus</i> and <i>Rhizophagus irregularis</i> . Plant and Cell Physiology, 2015, 56, 1490-1511.	3.1	140

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19	Strigolactone-Induced Putative Secreted Protein 1 Is Required for the Establishment of Symbiosis by the Arbuscular Mycorrhizal Fungus <i>Rhizophagus irregularis</i> . <i>Molecular Plant-Microbe Interactions</i> , 2016, 29, 277-286.	2.6	136
20	A shared gene drives lateral root development and root nodule symbiosis pathways in <i>Lotus</i> . <i>Science</i> , 2019, 366, 1021-1023.	12.6	135
21	Leguminous Plants: Inventors of Root Nodules to Accommodate Symbiotic Bacteria. <i>International Review of Cell and Molecular Biology</i> , 2015, 316, 111-158.	3.2	133
22	Positional Cloning Identifies <i>Lotus japonicus</i> NSP2, A Putative Transcription Factor of the GRAS Family, Required for NIN and ENOD40 Gene Expression in Nodule Initiation. <i>DNA Research</i> , 2006, 13, 255-265.	3.4	129
23	Shoot-applied MeJA Suppresses Root Nodulation in <i>Lotus japonicus</i> . <i>Plant and Cell Physiology</i> , 2006, 47, 176-180.	3.1	129
24	Aquaporin-mediated long-distance polyphosphate translocation directed towards the host in arbuscular mycorrhizal symbiosis: application of virus-induced gene silencing. <i>New Phytologist</i> , 2016, 211, 1202-1208.	7.3	122
25	Gibberellins Interfere with Symbiosis Signaling and Gene Expression and Alter Colonization by Arbuscular Mycorrhizal Fungi in <i>Lotus japonicus</i> . <i>Plant Physiology</i> , 2015, 167, 545-557.	4.8	120
26	klavier (klv), A novel hypernodulation mutant of <i>Lotus japonicus</i> affected in vascular tissue organization and floral induction. <i>Plant Journal</i> , 2005, 44, 505-515.	5.7	114
27	<i>TOO MUCH LOVE</i> , a Root Regulator Associated with the Long-Distance Control of Nodulation in <i>Lotus japonicus</i> . <i>Molecular Plant-Microbe Interactions</i> , 2009, 22, 259-268.	2.6	114
28	A <i>Lotus</i> basic leucine zipper protein with a RING-finger motif negatively regulates the developmental program of nodulation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 15206-15210.	7.1	113
29	Conservation of <i>Lotus</i> and <i>Arabidopsis</i> Basic Helix-Loop-Helix Proteins Reveals New Players in Root Hair Development. <i>Plant Physiology</i> , 2009, 151, 1175-1185.	4.8	113
30	The <i>Clavata2</i> genes of pea and <i>Lotus japonicus</i> affect autoregulation of nodulation. <i>Plant Journal</i> , 2011, 65, 861-871.	5.7	110
31	<i>TOO MUCH LOVE</i> , a Novel Kelch Repeat-Containing F-box Protein, Functions in the Long-Distance Regulation of the Legume-Rhizobium Symbiosis. <i>Plant and Cell Physiology</i> , 2013, 54, 433-447.	3.1	110
32	The receptor-like kinase KLAVIER mediates systemic regulation of nodulation and non-symbiotic shoot development in <i>Lotus japonicus</i> . <i>Development (Cambridge)</i> , 2010, 137, 4317-4325.	2.5	109
33	Host plant genome overcomes the lack of a bacterial gene for symbiotic nitrogen fixation. <i>Nature</i> , 2009, 462, 514-517.	27.8	103
34	The Integral Membrane Protein SEN1 is Required for Symbiotic Nitrogen Fixation in <i>Lotus japonicus</i> Nodules. <i>Plant and Cell Physiology</i> , 2012, 53, 225-236.	3.1	95
35	Genetics of Symbiosis in <i>Lotus japonicus</i> : Recombinant Inbred Lines, Comparative Genetic Maps, and Map Position of 35 Symbiotic Loci. <i>Molecular Plant-Microbe Interactions</i> , 2006, 19, 80-91.	2.6	94
36	The genome of <i>Rhizophagus clarus</i> HR1 reveals a common genetic basis for auxotrophy among arbuscular mycorrhizal fungi. <i>BMC Genomics</i> , 2018, 19, 465.	2.8	91

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37	A Positive Regulator of Nodule Organogenesis, NODULE INCEPTION, Acts as a Negative Regulator of Rhizobial Infection in <i>Lotus japonicus</i> . <i>Plant Physiology</i> , 2014, 165, 747-758.	4.8	84
38	<i>Lotus japonicus</i> 'Miyakojima' MG-20: An Early-Flowering Accession Suitable for Indoor Handling. <i>Journal of Plant Research</i> , 2000, 113, 507-509.	2.4	81
39	CERBERUS and NSP1 of <i>Lotus japonicus</i> are Common Symbiosis Genes that Modulate Arbuscular Mycorrhiza Development. <i>Plant and Cell Physiology</i> , 2013, 54, 1711-1723.	3.1	78
40	Responses of a Model Legume <i>Lotus japonicus</i> to Lipochitin Oligosaccharide Nodulation Factors Purified from <i>Mesorhizobium loti</i> JRL501. <i>Molecular Plant-Microbe Interactions</i> , 2001, 14, 848-856.	2.6	77
41	crinkle, a Novel Symbiotic Mutant That Affects the Infection Thread Growth and Alters the Root Hair, Trichome, and Seed Development in <i>Lotus japonicus</i> . <i>Plant Physiology</i> , 2003, 131, 1054-1063.	4.8	77
42	Molecular Framework of a Regulatory Circuit Initiating Two-Dimensional Spatial Patterning of Stomatal Lineage. <i>PLoS Genetics</i> , 2015, 11, e1005374.	3.5	74
43	Characterization of Mycorrhizas Formed by <i>Glomus</i> sp. on Roots of Hypernodulating Mutants of <i>Lotus japonicus</i> . <i>Journal of Plant Research</i> , 2000, 113, 443-448.	2.4	72
44	Plant-Microbe Communications for Symbiosis. <i>Plant and Cell Physiology</i> , 2010, 51, 1377-1380.	3.1	67
45	Myristate can be used as a carbon and energy source for the asymbiotic growth of arbuscular mycorrhizal fungi. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 25779-25788.	7.1	67
46	Genetic basis of cytokinin and auxin functions during root nodule development. <i>Frontiers in Plant Science</i> , 2013, 4, 42.	3.6	65
47	Root nodulation: a developmental program involving cell fate conversion triggered by symbiotic bacterial infection. <i>Current Opinion in Plant Biology</i> , 2014, 21, 16-22.	7.1	64
48	Polyphosphate accumulation is driven by transcriptome alterations that lead to near-synchronous and near-equivalent uptake of inorganic cations in an arbuscular mycorrhizal fungus. <i>New Phytologist</i> , 2014, 204, 638-649.	7.3	63
49	Expression of the CLE-RS3 gene suppresses root nodulation in <i>Lotus japonicus</i> . <i>Journal of Plant Research</i> , 2016, 129, 909-919.	2.4	59
50	Stimulation of asymbiotic sporulation in arbuscular mycorrhizal fungi by fatty acids. <i>Nature Microbiology</i> , 2019, 4, 1654-1660.	13.3	58
51	Long-Distance Control of Nodulation: Molecules and Models. <i>Molecules and Cells</i> , 2009, 27, 129-134.	2.6	57
52	Conserved genetic determinant of motor organ identity in <i>Medicago truncatula</i> and related legumes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 11723-11728.	7.1	57
53	The Novel Symbiotic Phenotype of Enhanced-Nodulating Mutant of <i>Lotus japonicus</i> : astray Mutant is an Early Nodulating Mutant with Wider Nodulation Zone. <i>Plant and Cell Physiology</i> , 2002, 43, 853-859.	3.1	56
54	Two Distinct EIN2 Genes Cooperatively Regulate Ethylene Signaling in <i>Lotus japonicus</i> . <i>Plant and Cell Physiology</i> , 2013, 54, 1469-1477.	3.1	55

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55	Evidence of non-tandemly repeated rDNAs and their intragenomic heterogeneity in <i>Rhizophagus irregularis</i> . <i>Communications Biology</i> , 2018, 1, 87.	4.4	55
56	Evolutionary Dynamics of Nitrogen Fixation in the Legume-Rhizobia Symbiosis. <i>PLoS ONE</i> , 2014, 9, e93670.	2.5	53
57	Endoreduplication-mediated initiation of symbiotic organ development in <i>Lotus japonicus</i> . <i>Development (Cambridge)</i> , 2014, 141, 2441-2445.	2.5	52
58	Different DNA-binding specificities of NLP and NIN transcription factors underlie nitrate-induced control of root nodulation. <i>Plant Cell</i> , 2021, 33, 2340-2359.	6.6	52
59	A comprehensive strategy for identifying long-distance mobile peptides in xylem sap. <i>Plant Journal</i> , 2015, 84, 611-620.	5.7	51
60	Oriented cell division shapes carnivorous pitcher leaves of <i>Sarracenia purpurea</i> . <i>Nature Communications</i> , 2015, 6, 6450.	12.8	50
61	Loss of function of <i>ASPARTIC PEPTIDASE NODULE-INDUCED 1 (APN1)</i> in <i>Lotus japonicus</i> restricts efficient nitrogen-fixing symbiosis with specific <i>Mesorhizobium loti</i> strains. <i>Plant Journal</i> , 2018, 93, 5-16.	5.7	46
62	The Presence of an Enzyme that Converts Indole-3-acetamide into IAA in Wild and Cultivated Rice. <i>Plant and Cell Physiology</i> , 1991, 32, 143-149.	3.1	45
63	Reaction-Diffusion Pattern in Shoot Apical Meristem of Plants. <i>PLoS ONE</i> , 2011, 6, e18243.	2.5	45
64	Pattern Dynamics in Adaxial-Abaxial Specific Gene Expression Are Modulated by a Plastid Retrograde Signal during <i>Arabidopsis thaliana</i> Leaf Development. <i>PLoS Genetics</i> , 2013, 9, e1003655.	3.5	44
65	A Set of <i>Lotus japonicus</i> Gifu x <i>Lotus burttii</i> Recombinant Inbred Lines Facilitates Map-based Cloning and QTL Mapping. <i>DNA Research</i> , 2012, 19, 317-323.	3.4	40
66	Two CLE genes are induced by phosphate in roots of <i>Lotus japonicus</i> . <i>Journal of Plant Research</i> , 2011, 124, 155-163.	2.4	39
67	<i>Lotus burttii</i> Takes a Position of the Third Corner in the Lotus Molecular Genetics Triangle. <i>DNA Research</i> , 2005, 12, 69-77.	3.4	38
68	plenty, a Novel Hypernodulation Mutant in <i>Lotus japonicus</i> . <i>Plant and Cell Physiology</i> , 2010, 51, 1425-1435.	3.1	38
69	The Thiamine Biosynthesis Gene <i>THI1</i> Promotes Nodule Growth and Seed Maturation. <i>Plant Physiology</i> , 2016, 172, 2033-2043.	4.8	38
70	The Excessive Production of Indole-3-Acetic Acid and Its Significance in Studies of the Biosynthesis of This Regulator of Plant Growth and Development. <i>Plant and Cell Physiology</i> , 1996, 37, 1043-1048.	3.1	36
71	The SNARE Protein <i>SYP71</i> Expressed in Vascular Tissues Is Involved in Symbiotic Nitrogen Fixation in <i>Lotus japonicus</i> Nodules. <i>Plant Physiology</i> , 2012, 160, 897-905.	4.8	36
72	Function and evolution of a <i>Lotus japonicus</i> AP2/ERF family transcription factor that is required for development of infection threads. <i>DNA Research</i> , 2016, 24, dsw052.	3.4	36

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73	New Nodulation Mutants Responsible for Infection Thread Development in <i>Lotus japonicus</i> . <i>Molecular Plant-Microbe Interactions</i> , 2006, 19, 801-810.	2.6	32
74	Characterization of the <i>Lotus japonicus</i> Symbiotic Mutant <i>lot1</i> That Shows a Reduced Nodule Number and Distorted Trichomes. <i>Plant Physiology</i> , 2005, 137, 1261-1271.	4.8	31
75	MIR2111-5 locus and shoot-accumulated mature miR2111 systemically enhance nodulation depending on HAR1 in <i>Lotus japonicus</i> . <i>Nature Communications</i> , 2020, 11, 5192.	12.8	31
76	Structure-Specific Regulation of Nutrient Transport and Metabolism in Arbuscular Mycorrhizal Fungi. <i>Plant and Cell Physiology</i> , 2019, 60, 2272-2281.	3.1	30
77	Pollen Development and Tube Growth are Affected in the Symbiotic Mutant of <i>Lotus japonicus</i> , <i>crinkle</i> . <i>Plant and Cell Physiology</i> , 2004, 45, 511-520.	3.1	29
78	Shoot HAR1 mediates nitrate inhibition of nodulation in <i>Lotus japonicus</i> . <i>Plant Signaling and Behavior</i> , 2015, 10, e1000138.	2.4	29
79	The relationship between thiamine and two symbioses: Root nodule symbiosis and arbuscular mycorrhiza. <i>Plant Signaling and Behavior</i> , 2016, 11, e1265723.	2.4	29
80	cDNA Macroarray Analysis of Gene Expression in Ineffective Nodules Induced on the <i>Lotus japonicus</i> <i>sen1</i> Mutant. <i>Molecular Plant-Microbe Interactions</i> , 2004, 17, 1223-1233.	2.6	25
81	Leguminous nodule symbiosis involves recruitment of factors contributing to lateral root development. <i>Current Opinion in Plant Biology</i> , 2021, 59, 102000.	7.1	24
82	LACK OF SYMBIONT ACCOMMODATION controls intracellular symbiont accommodation in root nodule and arbuscular mycorrhizal symbiosis in <i>Lotus japonicus</i> . <i>PLoS Genetics</i> , 2019, 15, e1007865.	3.5	23
83	PLENTY, a hydroxyproline O-arabinosyltransferase, negatively regulates root nodule symbiosis in <i>Lotus japonicus</i> . <i>Journal of Experimental Botany</i> , 2019, 70, 507-517.	4.8	23
84	Asymbiotic mass production of the arbuscular mycorrhizal fungus <i>Rhizophagus clarus</i> . <i>Communications Biology</i> , 2022, 5, 43.	4.4	22
85	<i>TRICOT</i> encodes an AMP1-related carboxypeptidase that regulates root nodule development and shoot apical meristem maintenance in <i>Lotus japonicus</i> . <i>Development (Cambridge)</i> , 2013, 140, 353-361.	2.5	21
86	Nitrate transport via NRT2.1 mediates NIN-LIKE PROTEIN-dependent suppression of root nodulation in <i>Lotus japonicus</i> . <i>Plant Cell</i> , 2022, 34, 1844-1862.	6.6	21
87	Reactions of <i>Lotus japonicus</i> ecotypes and mutants to root parasitic plants. <i>Journal of Plant Physiology</i> , 2009, 166, 353-362.	3.5	20
88	Nodule Organogenesis in <i>Lotus japonicus</i> . <i>Journal of Plant Research</i> , 2000, 113, 489-495.	2.4	19
89	SLEEPLESS, a gene conferring nyctinastic movement in legume. <i>Journal of Plant Research</i> , 2003, 116, 151-154.	2.4	19
90	Spatiotemporal deep imaging of syncytium induced by the soybean cyst nematode <i>Heterodera glycines</i> . <i>Protoplasma</i> , 2017, 254, 2107-2115.	2.1	19

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91	ERN1 and CYCLOPS coordinately activate NIN signaling to promote infection thread formation in <i>Lotus japonicus</i> . <i>Journal of Plant Research</i> , 2019, 132, 641-653.	2.4	19
92	Gibberellin regulates infection and colonization of host roots by arbuscular mycorrhizal fungi. <i>Plant Signaling and Behavior</i> , 2015, 10, e1028706.	2.4	18
93	Transcriptomic profiles of nodule senescence in <i>Lotus japonicus</i> and <i>Mesorhizobium loti</i> symbiosis. <i>Plant Biotechnology</i> , 2014, 31, 345-349.	1.0	17
94	Endogenous gibberellins affect root nodule symbiosis via transcriptional regulation of NODULE INCEPTION in <i>Lotus japonicus</i> . <i>Plant Journal</i> , 2021, 105, 1507-1520.	5.7	17
95	Down-Regulation of NSP2 Expression in Developmentally Young Regions of <i>Lotus japonicus</i> Roots in Response to Rhizobial Inoculation. <i>Plant and Cell Physiology</i> , 2013, 54, 518-527.	3.1	16
96	Systemic Regulation of Root Nodule Formation. , 0, , .		16
97	Reactive Sulfur Species Interact with Other Signal Molecules in Root Nodule Symbiosis in <i>Lotus japonicus</i> . <i>Antioxidants</i> , 2020, 9, 145.	5.1	16
98	Hairy Root Transformation in <i>Lotus japonicus</i> . <i>Bio-protocol</i> , 2013, 3, .	0.4	15
99	Partial purification of an enzyme hydrolyzing indole-3-acetamide from rice cells. <i>Journal of Plant Research</i> , 2004, 117, 191-8.	2.4	14
100	Isolation and Phenotypic Characterization of <i>Lotus japonicus</i> Mutants Specifically Defective in Arbuscular Mycorrhizal Formation. <i>Plant and Cell Physiology</i> , 2014, 55, 928-941.	3.1	14
101	Requirement for <i>Mesorhizobium loti</i> Ornithine Transcarbamoylase for Successful Symbiosis with <i>Lotus japonicus</i> as Revealed by an Unexpected Long-Range Genome Deletion. <i>Plant and Cell Physiology</i> , 2008, 49, 301-313.	3.1	11
102	Systemic Optimization of Legume Nodulation: A Shoot-Derived Regulator, miR2111. <i>Frontiers in Plant Science</i> , 2021, 12, 682486.	3.6	11
103	Expression and Functional Analysis of a CLV3-Like Gene in the Model Legume <i>Lotus japonicus</i> . <i>Plant and Cell Physiology</i> , 2011, 52, 1211-1221.	3.1	10
104	Autoregulation of nodulation pathway is dispensable for nitrate-induced control of rhizobial infection. <i>Plant Signaling and Behavior</i> , 2020, 15, 1733814.	2.4	10
105	Analysis of two potential long-distance signaling molecules, LjCLE-RS1/2 and jasmonic acid, in a hypernodulating mutant too much love. <i>Plant Signaling and Behavior</i> , 2010, 5, 403-405.	2.4	9
106	Induction of localized auxin response during spontaneous nodule development in <i>Lotus japonicus</i> . <i>Plant Signaling and Behavior</i> , 2013, 8, e23359.	2.4	9
107	Common symbiosis genes CERBERUS and NSP1 provide additional insight into the establishment of arbuscular mycorrhizal and root nodule symbioses in <i>Lotus japonicus</i> . <i>Plant Signaling and Behavior</i> , 2014, 9, e28544.	2.4	9
108	Spatial regularity control of phyllotaxis pattern generated by the mutual interaction between auxin and PIN1. <i>PLoS Computational Biology</i> , 2018, 14, e1006065.	3.2	9

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109	Pattern formation by two-layer Turing system with complementary synthesis. <i>Journal of Theoretical Biology</i> , 2013, 322, 33-45.	1.7	8
110	CLE-HAR1 Systemic Signaling and NIN-Mediated Local Signaling Suppress the Increased Rhizobial Infection in the daphne Mutant of <i>Lotus japonicus</i> . <i>Molecular Plant-Microbe Interactions</i> , 2020, 33, 320-327.	2.6	8
111	Auxin methylation by <i>IAMT1</i> , duplicated in the legume lineage, promotes root nodule development in <i>Lotus japonicus</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, e2116549119.	7.1	8
112	Molecular Characterization of LjABCG1, an ATP-Binding Cassette Protein in <i>Lotus japonicus</i> . <i>PLoS ONE</i> , 2015, 10, e0139127.	2.5	7
113	Mutants of <i>Lotus japonicus</i> deficient in flavonoid biosynthesis. <i>Journal of Plant Research</i> , 2021, 134, 341-352.	2.4	6
114	Polymorphisms of E1 and GIGANTEA in wild populations of <i>Lotus japonicus</i> . <i>Journal of Plant Research</i> , 2014, 127, 651-660.	2.4	5
115	Strategy for shoot meristem proliferation in plants. <i>Plant Signaling and Behavior</i> , 2011, 6, 1851-1854.	2.4	3
116	The transcription activation and homodimerization of <i>Lotus japonicus</i> Nod factor Signaling Pathway2 protein. <i>Plant Signaling and Behavior</i> , 2013, 8, e26457.	2.4	3
117	Spatial regulation of resource allocation in response to nutritional availability. <i>Journal of Theoretical Biology</i> , 2020, 486, 110078.	1.7	3
118	Assessment of <i>Polygala paniculata</i> (Polygalaceae) characteristics for evolutionary studies of legume-rhizobia symbiosis. <i>Journal of Plant Research</i> , 2020, 133, 109-122.	2.4	3
119	Mechanisms of Rice Endophytic Bradyrhizobial Cell Differentiation and Its Role in Nitrogen Fixation. <i>Microbes and Environments</i> , 2020, 35, n/a.	1.6	3
120	Taxonomic revision of <i>Termitomyces</i> species found in Ryukyu Archipelago, Japan, based on phylogenetic analyses with three loci. <i>Mycoscience</i> , 2022, 63, 33-38.	0.8	3
121	<i>Lotus japonicus</i> HAR1 regulates root morphology locally and systemically under a moderate nitrate condition in the absence of rhizobia. <i>Planta</i> , 2022, 255, 95.	3.2	3
122	Current Development of <i>Lotus japonicus</i> Research. <i>Journal of Plant Research</i> , 2000, 113, 449-449.	2.4	2
123	Isolation and Characterization of Arbuscules from Roots of an Increased-arbuscule-forming Mutant of <i>Lotus japonicus</i> . <i>Annals of Botany</i> , 2007, 100, 1599-1603.	2.9	2
124	Morphological Effects of Sinefungin, an Inhibitor of S-Adenosylmethionine-Dependent Methyltransferases, on <i>Anabaena</i> sp. PCC 7120. <i>Microbes and Environments</i> , 2008, 23, 346-349.	1.6	2
125	The evolution of symbiotic systems. <i>Cellular and Molecular Life Sciences</i> , 2011, 68, 1283-1284.	5.4	2
126	Activator and Inhibitor Leading to Generation and Stabilization of Symbiotic Organ Development in Legume. , 2005, , 179-182.		1

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127	Grafting analysis indicates that malfunction ofTRICOTin the root causes a nodulation-deficient phenotype in <i>Lotus japonicus</i> . <i>Plant Signaling and Behavior</i> , 2013, 8, e23497.	2.4	0
128	Genes for Autoregulation of Nodulation. <i>Compendium of Plant Genomes</i> , 2014, , 73-78.	0.5	0
129	Fluorescent Labeling of the Cyst Nematode <i>Heterodera glycines</i> in Deep-Tissue Live Imaging. <i>Cytologia</i> , 2017, 82, 251-259.	0.6	0
130	Wild Accessions and Mutant Resources. <i>Compendium of Plant Genomes</i> , 2014, , 211-220.	0.5	0