Kirankumar S Mysore

List of Publications by Year
in descending order

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252 papers	15,332 citations	13068 68 h-index	24915 109 g-index
271 all docs	271 docs citations	271 times ranked	13492 citing authors

#	Article	IF	Citations
1	Regulation of primary plant metabolism during plant-pathogen interactions and its contribution to plant defense. Frontiers in Plant Science, 2014, 5, 17.	1.7	554
2	Largeâ€scale insertional mutagenesis using the Tnt1 retrotransposon in the model legume Medicago truncatula. Plant Journal, 2008, 54, 335-347.	2.8	442
3	Celebrating 20 Years of Genetic Discoveries in Legume Nodulation and Symbiotic Nitrogen Fixation. Plant Cell, 2020, 32, 15-41.	3.1	416
4	Nonhost resistance: how much do we know?. Trends in Plant Science, 2004, 9, 97-104.	4.3	372
5	The Root Hair â€œInfectomeâ€•of Medicago truncatula Uncovers Changes in Cell Cycle Genes and Reveals a Requirement for Auxin Signaling in Rhizobial Infection. Plant Cell, 2014, 26, 4680-4701.	3.1	313
6	Tobacco rattle virusâ€“based virus-induced gene silencing in Nicotiana benthamiana. Nature Protocols, 2014, 9, 1549-1562.	5.5	283
7	New dimensions for VIGS in plant functional genomics. Trends in Plant Science, 2011, 16, 656-665.	4.3	279
8	A GRAS-Type Transcription Factor with a Specific Function in Mycorrhizal Signaling. Current Biology, 2012, 22, 2236-2241.	1.8	262
9	Plant growth-promoting rhizobacteria systemically protectArabidopsis thalianaagainstCucumber mosaic virusby a salicylic acid and NPR1-independent and jasmonic acid-dependent signaling pathway. Plant Journal, 2004, 39, 381-392.	2.8	242
10	The Phytotoxin Coronatine Contributes to Pathogen Fitness and Is Required for Suppression of Salicylic Acid Accumulation in Tomato Inoculated with Pseudomonas syringae pv. tomato DC3000. Molecular Plant-Microbe Interactions, 2007, 20, 955-965.	1.4	222
11	Glycolate Oxidase Modulates Reactive Oxygen Speciesâ€“Mediated Signal Transduction during Nonhost Resistance in Nicotiana benthamiana and Arabidopsis Â. Plant Cell, 2012, 24, 336-352.	3.1	215
12	Agrodrench: a novel and effective agroinoculation method for virus-induced gene silencing in roots and diverse Solanaceous species. Plant Journal, 2004, 40, 322-331.	2.8	214
13	Vapyrin, a gene essential for intracellular progression of arbuscular mycorrhizal symbiosis, is also essential for infection by rhizobia in the nodule symbiosis of Medicago truncatula. Plant Journal, 2011, 65, 244-252.	2.8	211
14	Computational Estimation and Experimental Verification of Off-Target Silencing during Posttranscriptional Gene Silencing in Plants. Plant Physiology, 2006, 142, 429-440.	2.3	196
15	Identification of T-DNA tagged Arabidopsis mutants that are resistant to transformation by Agrobacterium. Molecular Genetics and Genomics, 1999, 261, 429-438.	2.4	177
16	NODULE INCEPTION Recruits the Lateral Root Developmental Program for Symbiotic Nodule Organogenesis in Medicago truncatula. Current Biology, 2019, 29, 3657-3668.e5.	1.8	177
17	Salicylic Acid and Systemic Acquired Resistance Play a Role in Attenuating Crown Gall Disease Caused byAgrobacterium tumefaciensÂ. Plant Physiology, 2008, 146, 323-324.	2.3	163
18	An Arabidopsis histone H2A mutant is deficient in Agrobacterium T-DNA integration. Proceedings of the National Academy of Sciences of the United States of America, 2000, 97, 948-953.	3.3	162

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19	Identification of Arabidopsis rat Mutants. Plant Physiology, 2003, 132, 494-505.	2.3	159
20	Nonhost Resistance Against Bacterial Pathogens: Retrospectives and Prospects. Annual Review of Phytopathology, 2013, 51, 407-427.	3.5	149
21	Medicago truncatula IPD3 Is a Member of the Common Symbiotic Signaling Pathway Required for Rhizobial and Mycorrhizal Symbioses. Molecular Plant-Microbe Interactions, 2011, 24, 1345-1358.	1.4	147
22	Monolignol ferulate conjugates are naturally incorporated into plant lignins. Science Advances, 2016, 2, e1600393.	4.7	147
23	Phytosterols Play a Key Role in Plant Innate Immunity against Bacterial Pathogens by Regulating Nutrient Efflux into the Apoplast Â Â Â. Plant Physiology, 2012, 158, 1789-1802.	2.3	146
24	Arabidopsis seedling flood-inoculation technique: a rapid and reliable assay for studying plant-bacterial interactions. Plant Methods, 2011, 7, 32.	1.9	145
25	Evolutionarily conserved repressive activity of WOX proteins mediates leaf blade outgrowth and floral organ development in plants. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 366-371.	3.3	144
26	Control of Compound Leaf Development byFLORICAULA/LEAFYOrthologSINGLE LEAFLET1inMedicago truncatulaÂ Â Â Â. Plant Physiology, 2008, 146, 1759-1772.	2.3	139
27	A WD40 Repeat Protein fromMedicago truncatulaIs Necessary for Tissue-Specific Anthocyanin and Proanthocyanidin Biosynthesis But Not for Trichome Development Â Â. Plant Physiology, 2009, 151, 1114-1129.	2.3	137
28	Regulation of anthocyanin and proanthocyanidin biosynthesis by <scp>M</scp>edicago truncatula b<scp>HLH</scp> transcription factor <scp>M</scp>t<scp>TT</scp>8. New Phytologist, 2016, 210, 905-921.	3.5	136
29	MtPAR MYB transcription factor acts as an on switch for proanthocyanidin biosynthesis in Medicago truncatula. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 1766-1771.	3.3	135
30	DELLA-mediated gibberellin signalling regulates Nod factor signalling and rhizobial infection. Nature Communications, 2016, 7, 12636.	5.8	135
31	STENOFOLIARegulates Blade Outgrowth and Leaf Vascular Patterning inMedicago truncatulaandNicotiana sylvestrisÂ Â Â. Plant Cell, 2011, 23, 2125-2142.	3.1	133
32	The Medicago FLOWERING LOCUS T Homolog, MtFTa1, Is a Key Regulator of Flowering Time Â Â Â. Plant Physiology, 2011, 156, 2207-2224.	2.3	133
33	A H+-ATPase That Energizes Nutrient Uptake during Mycorrhizal Symbioses in Rice and Medicago truncatula Â Â Â. Plant Cell, 2014, 26, 1818-1830.	3.1	131
34	Comprehensive transcript profiling of Pto- and Prf-mediated host defense responses to infection byPseudomonas syringaepv.tomato. Plant Journal, 2002, 32, 299-315.	2.8	128
35	Medicago truncatula <scp>DNF</scp>2 is a <scp>PI</scp>â€<scp>PLC</scp>â€<scp>XD</scp>â€containing protein required for bacteroid persistence and prevention of nodule early senescence and defenseâ€like reactions. New Phytologist, 2013, 197, 1250-1261.	3.5	128
36	A systematic study to determine the extent of gene silencing in Nicotiana benthamiana and other Solanaceae species when heterologous gene sequences are used for virusâ€induced gene silencing. New Phytologist, 2007, 176, 782-791.	3.5	118

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37	Host Versus Nonhost Resistance: Distinct Wars with Similar Arsenals. Phytopathology, 2015, 105, 580-587.	1.1	118
38	NODULE ROOT and COCHLEATA Maintain Nodule Development and Are Legume Orthologs of Arabidopsis BLADE-ON-PETIOLE Genes. Plant Cell, 2012, 24, 4498-4510.	3.1	116
39	Virus-induced gene silencing is a versatile tool for unraveling the functional relevance of multiple abiotic-stress-responsive genes in crop plants. Frontiers in Plant Science, 2014, 5, 323.	1.7	114
40	An efficient reverse genetics platform in the model legume <scp>M</scp>edicago truncatula. New Phytologist, 2014, 201, 1065-1076.	3.5	113
41	Loss of Abaxial Leaf Epicuticular Wax inMedicago truncatula irg1/palm1Mutants Results in Reduced Spore Differentiation of Anthracnose and Nonhost Rust Pathogens. Plant Cell, 2012, 24, 353-370.	3.1	112
42	Insertional mutagenesis: a Swiss Army knife for functional genomics of Medicago truncatula. Trends in Plant Science, 2005, 10, 229-235.	4.3	111
43	NIN interacts with NLPs to mediate nitrate inhibition of nodulation in Medicago truncatula. Nature Plants, 2018, 4, 942-952.	4.7	111
44	Arabidopsis VIRE2 INTERACTING PROTEIN2 Is Required for Agrobacterium T-DNA Integration in Plants. Plant Cell, 2007, 19, 1695-1708.	3.1	109
45	A Medicago truncatula Tobacco Retrotransposon Insertion Mutant Collection with Defects in Nodule Development and Symbiotic Nitrogen Fixation Â Â. Plant Physiology, 2012, 159, 1686-1699.	2.3	109
46	Virusâ€induced gene silencing can persist for more than 2â€ƒyears and also be transmitted to progeny seedlings in Nicotiana benthamiana and tomato. Plant Biotechnology Journal, 2011, 9, 797-806.	4.1	108
47	Role of the Agrobacterium tumefaciens VirD2 Protein in T-DNA Transfer and Integration. Molecular Plant-Microbe Interactions, 1998, 11, 668-683.	1.4	107
48	Role of proline and pyrroline-5-carboxylate metabolism in plant defense against invading pathogens. Frontiers in Plant Science, 2015, 6, 503.	1.7	102
49	Local and Systemic Regulation of Plant Root System Architecture and Symbiotic Nodulation by a Receptor-Like Kinase. PLoS Genetics, 2014, 10, e1004891.	1.5	101
50	Monitoring in planta bacterial infection at both cellular and wholeâ€plant levels using the green fluorescent protein variant GFPuv. New Phytologist, 2007, 174, 212-223.	3.5	98
51	A non<scp>RD</scp> receptorâ€like kinase prevents nodule early senescence and defenseâ€like reactions during symbiosis. New Phytologist, 2014, 203, 1305-1314.	3.5	97
52	Drought Stress Acclimation Imparts Tolerance to Sclerotinia sclerotiorum and Pseudomonas syringae in Nicotiana benthamiana. International Journal of Molecular Sciences, 2013, 14, 9497-9513.	1.8	95
53	Ornithineâ€deltaâ€aminotransferase and proline dehydrogenase genes play a role in nonâ€host disease resistance by regulating pyrrolineâ€5â€carboxylate metabolismâ€induced hypersensitive response. Plant, Cell and Environment, 2012, 35, 1329-1343.	2.8	93
54	Rhizobial Infection Is Associated with the Development of Peripheral Vasculature in Nodules ofMedicago truncatulaÂ Â Â. Plant Physiology, 2013, 162, 107-115.	2.3	92

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55	NODULES WITH ACTIVATED DEFENSE 1 is required for maintenance of rhizobial endosymbiosis in Medicago truncatula. New Phytologist, 2016, 212, 176-191.	3.5	90
56	DiVenn: An Interactive and Integrated Web-Based Visualization Tool for Comparing Gene Lists. Frontiers in Genetics, 2019, 10, 421.	1.1	85
57	A symbiosisâ€dedicated SYNTAXIN OF PLANTS 13II isoform controls the formation of a stable hostâ€“microbe interface in symbiosis. New Phytologist, 2016, 211, 1338-1351.	3.5	83
58	Diverse functions of multidrug and toxin extrusion (<scp>MATE</scp>) transporters in citric acid efflux and metal homeostasis in Medicago truncatula. Plant Journal, 2017, 90, 79-95.	2.8	83
59	Expression of a Finger Millet Transcription Factor, EcNAC1, in Tobacco Confers Abiotic Stress-Tolerance. PLoS ONE, 2012, 7, e40397.	1.1	83
60	Developmental Analysis of aMedicago truncatula smooth leaf margin1Mutant Reveals Context-Dependent Effects on Compound Leaf Development Â. Plant Cell, 2011, 23, 2106-2124.	3.1	82
61	Reverse Genetics in Medicago truncatula Using Tnt1 Insertion Mutants. Methods in Molecular Biology, 2011, 678, 179-190.	0.4	81
62	Abscisic Acid Promotion of Arbuscular Mycorrhizal Colonization Requires a Component of the PROTEIN PHOSPHATASE 2A Complex Â Â. Plant Physiology, 2014, 166, 2077-2090.	2.3	81
63	Control of dissected leaf morphology by a Cys(2)His(2) zinc finger transcription factor in the model legume Medicago truncatula. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 10754-10759.	3.3	80
64	Isolation and functional analysis of CONSTANS-LIKE genes suggests that a central role for CONSTANS in flowering time control is not evolutionarily conserved in Medicago truncatula. Frontiers in Plant Science, 2014, 5, 486.	1.7	80
65	Symbiotic root infections in Medicago truncatula require remorin-mediated receptor stabilization in membrane nanodomains. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 5289-5294.	3.3	80
66	Pathogenicity of Pseudomonas syringae pv. tomato on Tomato Seedlings: Phenotypic and Gene Expression Analyses of the Virulence Function of Coronatine. Molecular Plant-Microbe Interactions, 2008, 21, 383-395.	1.4	79
67	Medicago truncatula Molybdate Transporter type 1 (MtMOT1.3) is a plasma membrane molybdenum transporter required for nitrogenase activity in root nodules under molybdenum deficiency. New Phytologist, 2017, 216, 1223-1235.	3.5	79
68	Identification and Characterization of Plant Genes Involved in Agrobacterium-Mediated Plant Transformation by Virus-Induced Gene Silencing. Molecular Plant-Microbe Interactions, 2007, 20, 41-52.	1.4	77
69	GBF3 transcription factor imparts drought tolerance in Arabidopsis thaliana. Scientific Reports, 2017, 7, 9148.	1.6	77
70	Novel phosphate deficiency-responsive long non-coding RNAs in the legume model plant Medicago truncatula. Journal of Experimental Botany, 2017, 68, 5937-5948.	2.4	77
71	Caveat of RNAi in Plants: The Off-Target Effect. Methods in Molecular Biology, 2011, 744, 13-25.	0.4	76
72	At<scp>MBP</scp>â€1, an alternative translation product of <scp>LOS</scp>2, affects abscisic acid responses and is modulated by the <scp>E</scp>3 ubiquitin ligase <scp>A</scp>t<scp>SAP</scp>5. Plant Journal, 2013, 76, 481-493.	2.8	76

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73	Plant Ribosomal Proteins, RPL12 and RPL19, Play a Role in Nonhost Disease Resistance against Bacterial Pathogens. Frontiers in Plant Science, 2015, 6, 1192.	1.7	71
74	NO APICAL MERISTEM (MtNAM) regulates floral organ identity and lateral organ separation in Medicago truncatula. New Phytologist, 2012, 195, 71-84.	3.5	68
75	The Nâ€Acylethanolamineâ€Mediated Regulatory Pathway in Plants. Chemistry and Biodiversity, 2007, 4, 1933-1955.	1.0	67
76	The MicroRNA390/TAS3 Pathway Mediates Symbiotic Nodulation and Lateral Root Growth. Plant Physiology, 2017, 174, 2469-2486.	2.3	67
77	Expression of theArabidopsishistone H2A-1gene correlates with susceptibility toAgrobacteriumtransformation. Plant Journal, 2002, 32, 285-298.	2.8	65
78	Mutagenesis and Beyond! Tools for Understanding Legume Biology. Plant Physiology, 2009, 151, 978-984.	2.3	65
79	Global Gene Expression Profiling During Medicago truncatulaâ€“Phymatotrichopsis omnivora Interaction Reveals a Role for Jasmonic Acid, Ethylene, and the Flavonoid Pathway in Disease Development. Molecular Plant-Microbe Interactions, 2009, 22, 7-17.	1.4	65
80	TheTrans-Acting Short Interfering RNA3 Pathway and NO APICAL MERISTEM Antagonistically Regulate Leaf Margin Development and Lateral Organ Separation, as Revealed by Analysis of anargonaute7/lobed leaflet1Mutant inMedicagoÂtruncatulaÂ Â. Plant Cell, 2014, 25, 4845-4862.	3.1	64
81	NIN-like protein transcription factors regulate leghemoglobin genes in legume nodules. Science, 2021, 374, 625-628.	6.0	61
82	The Medicago truncatula LysM receptorâ€like kinase LYK9 plays a dual role in immunity and the arbuscular mycorrhizal symbiosis. New Phytologist, 2019, 223, 1516-1529.	3.5	59
83	Overexpression of a fatty acid amide hydrolase compromises innate immunity in Arabidopsis. Plant Journal, 2008, 56, 336-349.	2.8	58
84	Overexpression of the Disease Resistance Gene Pto in Tomato Induces Gene Expression Changes Similar to Immune Responses in Human and Fruitfly Â. Plant Physiology, 2003, 132, 1901-1912.	2.3	57
85	Agroinoculation and Agroinfiltration: Simple Tools for Complex Gene Function Analyses. Methods in Molecular Biology, 2011, 678, 65-76.	0.4	57
86	GENERAL CONTROL NONREPRESSIBLE4 Degrades 14-3-3 and the RIN4 Complex to Regulate Stomatal Aperture with Implications on Nonhost Disease Resistance and Drought Tolerance. Plant Cell, 2017, 29, 2233-2248.	3.1	56
87	Jasmonate ZIM-Domain (JAZ) Protein Regulates Host and Nonhost Pathogen-Induced Cell Death in Tomato and Nicotiana benthamiana. PLoS ONE, 2013, 8, e75728.	1.1	56
88	Overexpression ofMedicago SVPgenes causes floral defects and delayed flowering inArabidopsisbut only affects floral development inMedicago. Journal of Experimental Botany, 2014, 65, 429-442.	2.4	55
89	Coronatine inhibits stomatal closure and delays hypersensitive response cell death induced by nonhost bacterial pathogens. PeerJ, 2013, 1, e34.	0.9	55
90	MtMOT1.2 is responsible for molybdate supply to <scp>Medicago truncatula</scp> nodules. Plant, Cell and Environment, 2019, 42, 310-320.	2.8	54

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91	pssRNAit: A Web Server for Designing Effective and Specific Plant siRNAs with Genome-Wide Off-Target Assessment. Plant Physiology, 2020, 184, 65-81.	2.3	54
92	Functional characterization of Nicotiana benthamiana homologs of peanut water deficit-induced genes by virus-induced gene silencing. Planta, 2007, 225, 523-539.	1.6	52
93	Genes involved in nonhost disease resistance as a key to engineer durable resistance in crops. Plant Science, 2019, 279, 108-116.	1.7	52
94	Arabidopsisecotypes and mutants that are recalcitrant toAgrobacteriumroot transformation are susceptible to germ-line transformation. Plant Journal, 2000, 21, 9-16.	2.8	51
95	Arabidopsis Heterotrimeric G-Proteins Play a Critical Role in Host and Nonhost Resistance against Pseudomonas syringae Pathogens. PLoS ONE, 2013, 8, e82445.	1.1	50
96	The Small GTPase ROP10 of Medicago truncatula Is Required for Both Tip Growth of Root Hairs and Nod Factor-Induced Root Hair Deformation. Plant Cell, 2015, 27, 806-822.	3.1	50
97	The SAL-PAP Chloroplast Retrograde Pathway Contributes to Plant Immunity by Regulating Glucosinolate Pathway and Phytohormone Signaling. Molecular Plant-Microbe Interactions, 2017, 30, 829-841.	1.4	50
98	Agrobacterium-Mediated Transformation of Tomato with rolB Gene Results in Enhancement of Fruit Quality and Foliar Resistance against Fungal Pathogens. PLoS ONE, 2014, 9, e96979.	1.1	49
99	Different cytokinin histidine kinase receptors regulate nodule initiation as well as later nodule developmental stages in Medicago truncatula. Plant, Cell and Environment, 2016, 39, 2198-2209.	2.8	49
100	The Symbiosis-Related ERN Transcription Factors Act in Concert to Coordinate Rhizobial Host Root Infection. Plant Physiology, 2016, 171, pp.00230.2016.	2.3	48
101	Comprehensive analysis of small RNA-seq data reveals that combination of miRNA with its isomiRs increase the accuracy of target prediction in Arabidopsis thaliana. RNA Biology, 2014, 11, 1414-1429.	1.5	46
102	Aldoâ€keto reductase enzymes detoxify glyphosate and improve herbicide resistance in plants. Plant Biotechnology Journal, 2017, 15, 794-804.	4.1	46
103	MiR393 and miR390 synergistically regulate lateral root growth in rice under different conditions. BMC Plant Biology, 2018, 18, 261.	1.6	46
104	NTRC and Chloroplast-Generated Reactive Oxygen Species Regulate Pseudomonas syringae pv. tomato Disease Development in Tomato and Arabidopsis. Molecular Plant-Microbe Interactions, 2012, 25, 294-306.	1.4	45
105	Arabidopsis stress associated protein 9 mediates biotic and abiotic stress responsive ABA signaling via the proteasome pathway. Plant, Cell and Environment, 2017, 40, 702-716.	2.8	45
106	Virus-induced gene silencing and its application in characterizing genes involved in water-deficit-stress tolerance. Journal of Plant Physiology, 2008, 165, 1404-1421.	1.6	44
107	SGT1 positively regulates the process of plant cell death during both compatible and incompatible plantâ€“pathogen interactions. Molecular Plant Pathology, 2010, 11, 597-611.	2.0	44
108	Functional specialization of duplicated AP3â€like genes in Medicago truncatula. Plant Journal, 2013, 73, 663-675.	2.8	43

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109	Agrobacterium Tâ€<scp>DNA</scp> integration into the plant genome can occur without the activity of key nonâ€homologous endâ€joining proteins. Plant Journal, 2015, 81, 934-946.	2.8	43
110	Transcriptomic and metabolomic analyses identify a role for chlorophyll catabolism and phytoalexin during Medicago nonhost resistance against Asian soybean rust. Scientific Reports, 2015, 5, 13061.	1.6	41
111	Characterization of the Rust Fungus, Puccinia emaculata, and Evaluation of Genetic Variability for Rust Resistance in Switchgrass Populations. Bioenergy Research, 2013, 6, 458-468.	2.2	40
112	AtCYP710A1gene-mediated stigmasterol production plays a role in imparting temperature stress tolerance inArabidopsis thaliana. Plant Signaling and Behavior, 2013, 8, e23142.	1.2	40
113	Retroelement insertions at the <scp>M</scp>edicago FTa1 locus in spring mutants eliminate vernalisation but not longâ€day requirements for early flowering. Plant Journal, 2013, 76, 580-591.	2.8	40
114	Strigolactones contribute to shoot elongation and to the formation of leaf margin serrations in Medicago truncatula R108. Journal of Experimental Botany, 2015, 66, 1237-1244.	2.4	40
115	Role of the Nod Factor Hydrolase MtNFH1 in Regulating Nod Factor Levels during Rhizobial Infection and in Mature Nodules of Medicago truncatula. Plant Cell, 2018, 30, 397-414.	3.1	40
116	MtNODULE ROOT1 and MtNODULE ROOT2 Are Essential for Indeterminate Nodule Identity. Plant Physiology, 2018, 178, 295-316.	2.3	40
117	A molecular framework underlying the compound leaf pattern of Medicago truncatula. Nature Plants, 2020, 6, 511-521.	4.7	40
118	Functional characterization of three water deficit stress-induced genes in tobacco and Arabidopsis: An approach based on gene down regulation. Plant Physiology and Biochemistry, 2010, 48, 35-44.	2.8	39
119	Glycolate oxidase is an alternative source for H₂O₂production during plant defense responses and functions independently from NADPH oxidase. Plant Signaling and Behavior, 2012, 7, 752-755.	1.2	38
120	Evolution by gene duplication ofMedicago truncatula PISTILLATA-like transcription factors. Journal of Experimental Botany, 2016, 67, 1805-1817.	2.4	38
121	Sinorhizobium meliloti succinylated highâ€molecularâ€weight succinoglycan and the Medicago truncatula LysM receptorâ€like kinase MtLYK10 participate independently in symbiotic infection. Plant Journal, 2020, 102, 311-326.	2.8	37
122	Two euAGAMOUS Genes Control C-Function in Medicago truncatula. PLoS ONE, 2014, 9, e103770.	1.1	36
123	The CLE53â€“SUNN genetic pathway negatively regulates arbuscular mycorrhiza root colonization in Medicago truncatula. Journal of Experimental Botany, 2020, 71, 4972-4984.	2.4	36
124	Agrobacterium tumefaciens Transformation of the Radiation Hypersensitive Arabidopsis thaliana Mutants uvh1 and rad5. Molecular Plant-Microbe Interactions, 1998, 11, 1136-1141.	1.4	34
125	A Virus-Induced Gene Silencing Screen Identifies a Role for Thylakoid Formation1 in Pseudomonas syringae pv tomato Symptom Development in Tomato and Arabidopsis. Plant Physiology, 2009, 152, 281-292.	2.3	34
126	Forward Genetics Screening of Medicago truncatula Tnt1 Insertion Lines. Methods in Molecular Biology, 2013, 1069, 93-100.	0.4	34

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127	<scp>LOOSE FLOWER</scp>, a <scp>WUSCHEL</scp>â€like Homeobox gene, is required for lateral fusion of floral organs in Medicago truncatula. Plant Journal, 2015, 81, 480-492.	2.8	34
128	Functional characterisation of brassinosteroid receptor MtBRI1 in Medicago truncatula. Scientific Reports, 2017, 7, 9327.	1.6	34
129	IPD3 and IPD3L Function Redundantly in Rhizobial and Mycorrhizal Symbioses. Frontiers in Plant Science, 2018, 9, 267.	1.7	34
130	Apoplastic Extracts from a Transgenic Wheat Line Exhibiting Lesion-Mimic Phenotype Have Multiple Pathogenesis-Related Proteins That Are Antifungal. Molecular Plant-Microbe Interactions, 2004, 17, 1306-1317.	1.4	33
131	Nuclear DNA content in species ofEleusine (Gramineae): A critical re-evaluation using laser flow cytometry. Plant Systematics and Evolution, 1997, 207, 1-11.	0.3	32
132	Pseudomonas Type III Effector AvrPto Suppresses the Programmed Cell Death Induced by Two Nonhost Pathogens in Nicotiana benthamiana and Tomato. Molecular Plant-Microbe Interactions, 2004, 17, 1328-1336.	1.4	32
133	SGT1 contributes to coronatine signaling and Pseudomonas syringae pv. tomato disease symptom development in tomato and Arabidopsis. New Phytologist, 2011, 189, 83-93.	3.5	32
134	Several components of SKP1/Cullin/Fâ€box E3 ubiquitin ligase complex and associated factors play a role in Agrobacteriumâ€mediated plant transformation. New Phytologist, 2012, 195, 203-216.	3.5	32
135	Opposing control by transcription factors MYB61 and MYB3 Increases Freezing Tolerance by relieving C-repeat Binding Factor suppression. Plant Physiology, 2016, 172, pp.00051.2016.	2.3	32
136	A SOC1-like gene MtSOC1a promotes flowering and primary stem elongation in Medicago. Journal of Experimental Botany, 2018, 69, 4867-4880.	2.4	32
137	SLENDER RICE1 and Oryza sativa INDETERMINATE DOMAIN2 Regulating OsmiR396 Are Involved in Stem Elongation. Plant Physiology, 2020, 182, 2213-2227.	2.3	32
138	Agrobacterium expressing aÂtype III secretion system delivers Pseudomonas effectors into plant cells to enhance transformation. Nature Communications, 2022, 13, 2581.	5.8	32
139	A high-throughput virus-induced gene silencing protocol identifies genes involved in multi-stress tolerance. BMC Plant Biology, 2013, 13, 193.	1.6	31
140	DASH transcription factor impacts Medicago truncatula seed size by its action on embryo morphogenesis and auxin homeostasis. Plant Journal, 2015, 81, 453-466.	2.8	31
141	Mt<scp>VRN</scp>2 is a Polycomb <scp>VRN</scp>2â€like gene which represses the transition to flowering in the model legume Medicago truncatula. Plant Journal, 2016, 86, 145-160.	2.8	31
142	Transcriptomeâ€based analyses of phosphiteâ€mediated suppression of rust pathogens Puccinia emaculata and Phakopsora pachyrhizi and functional characterization of selected fungal target genes. Plant Journal, 2018, 93, 894-904.	2.8	31
143	HEADLESS, aWUSCHELhomolog, uncovers novel aspects of shoot meristem regulation and leaf blade development inMedicago truncatula. Journal of Experimental Botany, 2019, 70, 149-163.	2.4	31
144	Phymatotrichum (cotton) root rot caused by Phymatotrichopsis omnivora: retrospects and prospects. Molecular Plant Pathology, 2010, 11, 325-334.	2.0	30

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145	Agrobacterium May Delay Plant Nonhomologous End-Joining DNA Repair via XRCC4 to Favor T-DNA Integration. Plant Cell, 2012, 24, 4110-4123.	3.1	30
146	Medicago truncatula SOC1 Genes Are Up-regulated by Environmental Cues That Promote Flowering. Frontiers in Plant Science, 2018, 9, 496.	1.7	30
147	The future of legume genetic data resources: Challenges, opportunities, and priorities. , 2019, 1, e16.		30
148	Suppression of plant defense responses by extracellular metabolites from Pseudomonas syringae pv. tabaci in Nicotiana benthamiana. BMC Plant Biology, 2013, 13, 65.	1.6	28
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