

# Maria J Harrison

## List of Publications by Year in descending order

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80  
papers

12,989  
citations

26630

56  
h-index

66911

78  
g-index

85  
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85  
docs citations

85  
times ranked

7610  
citing authors

#	ARTICLE	IF	CITATIONS
1	Fifteen compelling open questions in plant cell biology. <i>Plant Cell</i> , 2022, 34, 72-102.	6.6	27
2	<scp><i>KIN3</i></scp> impacts arbuscular mycorrhizal symbiosis and promotes fungal colonisation in <i>Medicago truncatula</i>. <i>Plant Journal</i> , 2022, 110, 513-528.	5.7	9
3	A genetically encoded biosensor reveals spatiotemporal variation in cellular phosphate content in <i>Brachypodium distachyon</i> mycorrhizal roots. <i>New Phytologist</i> , 2022, 234, 1817-1831.	7.3	4
4	Conserved and reproducible bacterial communities associate with extraradical hyphae of arbuscular mycorrhizal fungi. <i>ISME Journal</i> , 2021, 15, 2276-2288.	9.8	91
5	Constitutive Overexpression of RAM1 Leads to an Increase in Arbuscule Density in <i>Brachypodium distachyon</i> . <i>Plant Physiology</i> , 2020, 184, 1263-1272.	4.8	11
6	A CLEâ€“SUNN module regulates strigolactone content and fungal colonization in arbuscular mycorrhiza. <i>Nature Plants</i> , 2019, 5, 933-939.	9.3	65
7	Phytohormones, miRNAs, and peptide signals integrate plant phosphorus status with arbuscular mycorrhizal symbiosis. <i>Current Opinion in Plant Biology</i> , 2019, 50, 132-139.	7.1	70
8	A Phosphate-Dependent Requirement for Transcription Factors IPD3 and IPD3L During Arbuscular Mycorrhizal Symbiosis in <i>Medicago truncatula</i>. <i>Molecular Plant-Microbe Interactions</i> , 2019, 32, 1277-1290.	2.6	11
9	Extensive membrane systems at the hostâ€“arbuscular mycorrhizal fungus interface. <i>Nature Plants</i> , 2019, 5, 194-203.	9.3	85
10	Transcriptomic analysis of field-droughted sorghum from seedling to maturity reveals biotic and metabolic responses. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 27124-27132.	7.1	129
11	Diverse <scp><i>Sorghum bicolor</i></scp> accessions show marked variation in growth and transcriptional responses to arbuscular mycorrhizal fungi. <i>Plant, Cell and Environment</i> , 2019, 42, 1758-1774.	5.7	60
12	Accumulation of phosphoinositides in distinct regions of the periarbuscular membrane. <i>New Phytologist</i> , 2019, 221, 2213-2227.	7.3	24
13	Genome and evolution of the arbuscular mycorrhizal fungus <i>Diversispora epigaea</i> (formerly) Tj ETQq1 1 0.784314 rgBT /Overl	7.3	88
14	A short LysM protein with high molecular diversity from an arbuscular mycorrhizal fungus, <i>Rhizophagus irregularis</i> . <i>Mycoscience</i> , 2019, 60, 63-70.	0.8	15
15	RiArsB and RiMT-11: Two novel genes induced by arsenate in arbuscular mycorrhiza. <i>Fungal Biology</i> , 2018, 122, 121-130.	2.5	13
16	Blumenols as shoot markers of root symbiosis with arbuscular mycorrhizal fungi. <i>ELife</i> , 2018, 7, .	6.0	69
17	Exocytosis for endosymbiosis: membrane trafficking pathways for development of symbiotic membrane compartments. <i>Current Opinion in Plant Biology</i> , 2017, 38, 101-108.	7.1	54
18	Arbuscular mycorrhizaâ€“specific enzymes FatM and <scp>RAM</scp>2 fineâ€“tune lipid biosynthesis to promote development of arbuscular mycorrhiza. <i>New Phytologist</i> , 2017, 214, 1631-1645.	7.3	260

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19	A Transcriptional Program for Arbuscule Degeneration during AM Symbiosis Is Regulated by MYB1. <i>Current Biology</i> , 2017, 27, 1206-1212.	3.9	110
20	Plant Signaling and Metabolic Pathways Enabling Arbuscular Mycorrhizal Symbiosis. <i>Plant Cell</i> , 2017, 29, 2319-2335.	6.6	241
21	A comprehensive draft genome sequence for lupin ( <i>Lupinus angustifolius</i> ), an emerging health food: insights into plant-microbe interactions and legume evolution. <i>Plant Biotechnology Journal</i> , 2017, 15, 318-330.	8.3	153
22	Genes conserved for arbuscular mycorrhizal symbiosis identified through phylogenomics. <i>Nature Plants</i> , 2016, 2, 15208.	9.3	206
23	DELLA proteins regulate expression of a subset of AM symbiosis-induced genes in <i>Medicago truncatula</i> . <i>Plant Signaling and Behavior</i> , 2016, 11, e1162369.	2.4	23
24	A CCaMK-CYCLOPS-DELLA Complex Activates Transcription of RAM1 to Regulate Arbuscule Branching. <i>Current Biology</i> , 2016, 26, 987-998.	3.9	182
25	EXO70I Is Required for Development of a Sub-domain of the Periarbuscular Membrane during Arbuscular Mycorrhizal Symbiosis. <i>Current Biology</i> , 2015, 25, 2189-2195.	3.9	120
26	Suppression of Arbuscule Degeneration in <i>Medicago truncatula</i> phosphate transporter4 Mutants Is Dependent on the Ammonium Transporter 2 Family Protein AMT2;3. <i>Plant Cell</i> , 2015, 27, 1352-1366.	6.6	180
27	Hyphal branching during arbuscule development requires RAM1. <i>Plant Physiology</i> , 2015, 169, pp.01155.2015.	4.8	94
28	Signaling events during initiation of arbuscular mycorrhizal symbiosis. <i>Journal of Integrative Plant Biology</i> , 2014, 56, 250-261.	8.5	102
29	A set of fluorescent protein-based markers expressed from constitutive and arbuscular mycorrhiza-inducible promoters to label organelles, membranes and cytoskeletal elements in <i>Medicago truncatula</i> . <i>Plant Journal</i> , 2014, 80, 1151-1163.	5.7	121
30	Gene Silencing in <i>Medicago truncatula</i> Roots Using RNAi. <i>Methods in Molecular Biology</i> , 2013, 1069, 163-177.	0.9	10
31	DELLA proteins regulate arbuscule formation in arbuscular mycorrhizal symbiosis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, E5025-34.	7.1	266
32	Polar localization of a symbiosis-specific phosphate transporter is mediated by a transient reorientation of secretion. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, E665-72.	7.1	164
33	Cellular programs for arbuscular mycorrhizal symbiosis. <i>Current Opinion in Plant Biology</i> , 2012, 15, 691-698.	7.1	151
34	Diversity of morphology and function in arbuscular mycorrhizal symbioses in <i>Brachypodium distachyon</i> . <i>Planta</i> , 2012, 236, 851-865.	3.2	85
35	The half-size ABC transporters STR1 and STR2 are indispensable for mycorrhizal arbuscule formation in rice. <i>Plant Journal</i> , 2012, 69, 906-920.	5.7	131
36	Arsenate induces the expression of fungal genes involved in As transport in arbuscular mycorrhiza. <i>Fungal Biology</i> , 2011, 115, 1197-1209.	2.5	58

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37	<i>Medicago truncatula</i> <i>mtpt4</i> mutants reveal a role for nitrogen in the regulation of arbuscule degeneration in arbuscular mycorrhizal symbiosis. <i>Plant Journal</i> , 2011, 68, 954-965.	5.7	103
38	Genetic variation for root architecture, nutrient uptake and mycorrhizal colonisation in <i>Medicago truncatula</i> accessions. <i>Plant and Soil</i> , 2010, 336, 113-128.	3.7	13
39	<i>Medicago truncatula</i> Vapyrin is a novel protein required for arbuscular mycorrhizal symbiosis. <i>Plant Journal</i> , 2010, 61, 482-494.	5.7	198
40	Two <i>Medicago truncatula</i> Half-ABC Transporters Are Essential for Arbuscule Development in Arbuscular Mycorrhizal Symbiosis. <i>Plant Cell</i> , 2010, 22, 1483-1497.	6.6	223
41	Phosphate Transporters in Arbuscular Mycorrhizal Symbiosis. , 2010, , 117-135.		12
42	Reprogramming Plant Cells for Endosymbiosis. <i>Science</i> , 2009, 324, 753-754.	12.6	160
43	Live-Cell Imaging Reveals Periarbuscular Membrane Domains and Organelle Location in <i>Medicago truncatula</i> Roots during Arbuscular Mycorrhizal Symbiosis. <i>Plant Physiology</i> , 2009, 151, 809-819.	4.8	215
44	<i>Medicago truncatula</i> and <i>Glomus intraradices</i> gene expression in cortical cells harboring arbuscules in the arbuscular mycorrhizal symbiosis. <i>BMC Plant Biology</i> , 2009, 9, 10.	3.6	277
45	Laser microdissection and its application to analyze gene expression in arbuscular mycorrhizal symbiosis. <i>Pest Management Science</i> , 2009, 65, 504-511.	3.4	45
46	Novel plant and fungal AGP-like proteins in the <i>Medicago truncatula</i> - <i>Glomus intraradices</i> arbuscular mycorrhizal symbiosis. <i>Mycorrhiza</i> , 2008, 18, 403-412.	2.8	17
47	The <i>Medicago truncatula</i> ortholog of Arabidopsis EIN2, <i>sickle</i> , is a negative regulator of symbiotic and pathogenic microbial associations. <i>Plant Journal</i> , 2008, 55, 580-595.	5.7	272
48	Closely Related Members of the <i>Medicago truncatula</i> PHT1 Phosphate Transporter Gene Family Encode Phosphate Transporters with Distinct Biochemical Activities. <i>Journal of Biological Chemistry</i> , 2008, 283, 24673-24681.	3.4	87
49	A <i>Medicago truncatula</i> phosphate transporter indispensable for the arbuscular mycorrhizal symbiosis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 1720-1725.	7.1	634
50	Phosphate in the arbuscular mycorrhizal symbiosis: transport properties and regulatory roles. <i>Plant, Cell and Environment</i> , 2007, 30, 310-322.	5.7	339
51	Arbuscular mycorrhizal symbiosis is accompanied by local and systemic alterations in gene expression and an increase in disease resistance in the shoots. <i>Plant Journal</i> , 2007, 50, 529-544.	5.7	430
52	Loss of <i>At4</i> function impacts phosphate distribution between the roots and the shoots during phosphate starvation. <i>Plant Journal</i> , 2006, 45, 712-726.	5.7	205
53	Defensin gene family in <i>Medicago truncatula</i> : structure, expression and induction by signal molecules. <i>Plant Molecular Biology</i> , 2005, 58, 385-399.	3.9	73
54	RNA Interference Identifies a Calcium-Dependent Protein Kinase Involved in <i>Medicago truncatula</i> Root Development. <i>Plant Cell</i> , 2005, 17, 2911-2921.	6.6	147

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55	SIGNALING IN THE ARBUSCULAR MYCORRHIZAL SYMBIOSIS. Annual Review of Microbiology, 2005, 59, 19-42.	7.3	647
56	Expression of a xyloglucan endotransglucosylase/hydrolase gene, Mt-XTH1, from <i>Medicago truncatula</i> is induced systemically in mycorrhizal roots. Gene, 2005, 345, 191-197.	2.2	53
57	Phosphate transport in <i>Arabidopsis</i> : Pht1;1 and Pht1;4 play a major role in phosphate acquisition from both low- and high-phosphate environments. Plant Journal, 2004, 39, 629-642.	5.7	719
58	cDNA arrays as a tool to identify mycorrhiza-regulated genes: identification of mycorrhiza-induced genes that encode or generate signaling molecules implicated in the control of root growth. Canadian Journal of Botany, 2004, 82, 1177-1185.	1.1	32
59	A phosphate transporter from <i>Medicago truncatula</i> is expressed in the photosynthetic tissues of the plant and located in the chloroplast envelope. New Phytologist, 2003, 157, 291-302.	7.3	46
60	Transcript Profiling Coupled with Spatial Expression Analyses Reveals Genes Involved in Distinct Developmental Stages of an Arbuscular Mycorrhizal Symbiosis [W]. Plant Cell, 2003, 15, 2106-2123.	6.6	309
61	A Phosphate Transporter from <i>Medicago truncatula</i> Involved in the Acquisition of Phosphate Released by Arbuscular Mycorrhizal Fungi. Plant Cell, 2002, 14, 2413-2429.	6.6	733
62	A Chloroplast Phosphate Transporter, PHT2;1, Influences Allocation of Phosphate within the Plant and Phosphate-Starvation Responses. Plant Cell, 2002, 14, 1751-1766.	6.6	310
63	A Phosphate Transporter Gene from the Extra-Radical Mycelium of an Arbuscular Mycorrhizal Fungus <i>Glomus intraradices</i> Is Regulated in Response to Phosphate in the Environment. Molecular Plant-Microbe Interactions, 2001, 14, 1140-1148.	2.6	261
64	Microtubule organization in root cells of <i>Medicago truncatula</i> during development of an arbuscular mycorrhizal symbiosis with <i>Glomus versiforme</i> . Protoplasma, 2001, 217, 154-165.	2.1	76
65	The spatial expression patterns of a phosphate transporter (MtPT1) from <i>Medicago truncatula</i> indicate a role in phosphate transport at the root/soil interface. Plant Journal, 2001, 25, 281-293.	5.7	176
66	Biotrophic interfaces and nutrient transport in plant/fungal symbioses. Journal of Experimental Botany, 1999, 50, 1013-1022.	4.8	84
67	Construction and characterization of genomic libraries of two endomycorrhizal fungi: <i>Glomus versiforme</i> and <i>Gigaspora margarita</i> . Mycological Research, 1999, 103, 955-960.	2.5	29
68	MOLECULAR AND CELLULAR ASPECTS OF THE ARBUSCULAR MYCORRHIZAL SYMBIOSIS. Annual Review of Plant Biology, 1999, 50, 361-389.	14.3	397
69	The Down-Regulation of Mt4-Like Genes by Phosphate Fertilization Occurs Systemically and Involves Phosphate Translocation to the Shoots <sup>1</sup> . Plant Physiology, 1999, 119, 241-248.	4.8	229
70	Novel Genes Induced During an Arbuscular Mycorrhizal (AM) Symbiosis Formed Between <i>Medicago truncatula</i> and <i>Glomus versiforme</i> . Molecular Plant-Microbe Interactions, 1999, 12, 171-181.	2.6	78
71	Development of the arbuscular mycorrhizal symbiosis. Current Opinion in Plant Biology, 1998, 1, 360-365.	7.1	59
72	Characterization of the Mt4 gene from <i>Medicago truncatula</i> . Gene, 1998, 216, 47-53.	2.2	45

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73	Cloning and Characterization of Two Phosphate Transporters from <i>Medicago truncatula</i> Roots: Regulation in Response to Phosphate and to Colonization by Arbuscular Mycorrhizal (AM) Fungi. <i>Molecular Plant-Microbe Interactions</i> , 1998, 11, 14-22.	2.6	264
74	The arbuscular mycorrhizal symbiosis: an underground association. <i>Trends in Plant Science</i> , 1997, 2, 54-60.	8.8	155
75	A novel gene whose expression in <i>Medicago truncatula</i> roots is suppressed in response to colonization by vesicular-arbuscular mycorrhizal (VAM) fungi and to phosphate nutrition. , 1997, 34, 199-208.		113
76	The Arbuscular Mycorrhizal Symbiosis. , 1997, , 1-34.		14
77	A sugar transporter from <i>Medicago truncatula</i> : altered expression pattern in roots during vesicular-arbuscular (VA) mycorrhizal associations. <i>Plant Journal</i> , 1996, 9, 491-503.	5.7	192
78	A phosphate transporter from the mycorrhizal fungus <i>Glomus versiforme</i> . <i>Nature</i> , 1995, 378, 626-629.	27.8	575
79	Spatial patterns of expression of flavonoid/isoflavonoid pathway genes during interactions between roots of <i>Medicago truncatula</i> and the mycorrhizal fungus <i>Glomus versiforme</i> . <i>Plant Journal</i> , 1994, 6, 9-20.	5.7	207
80	Isoflavonoid Accumulation and Expression of Defense Gene Transcripts During the Establishment of Vesicular-Arbuscular Mycorrhizal Associations in Roots of <i>Medicago truncatula</i> . <i>Molecular Plant-Microbe Interactions</i> , 1993, 6, 643.	2.6	244