Martin Parniske

List of Publications by Year in Descending Order

Source: https://exaly.com/author-pdf/1706585/martin-parniske-publications-by-year.pdf

Version: 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

120	12,782	59	112
papers	citations	h-index	g-index
183	14,819	10.3	6.61
ext. papers	ext. citations	avg, IF	L-index

#	Paper	IF	Citations
120	A CCaMK/Cyclops response element in the promoter of Lotus japonicus Calcium-Binding Protein 1 (CBP1) mediates transcriptional activation in root symbioses <i>New Phytologist</i> , 2022 ,	9.8	1
119	A strawberry accession with elevated methyl anthranilate fruit concentration is naturally resistant to the pest fly Drosophila suzukii. <i>PLoS ONE</i> , 2020 , 15, e0234040	3.7	4
118	Maintenance and Quantitative Phenotyping of the Oomycete-plant Model Pathosystem. <i>Bio-protocol</i> , 2020 , 10, e3661	0.9	
117	as a Model for Studying the Root Symbioses of the Rosaceae. Frontiers in Plant Science, 2019, 10, 661	6.2	7
116	A set of Arabidopsis genes involved in the accommodation of the downy mildew pathogen Hyaloperonospora arabidopsidis. <i>PLoS Pathogens</i> , 2019 , 15, e1007747	7.6	7
115	A unified multi-kingdom Golden Gate cloning platform. Scientific Reports, 2019, 9, 10131	4.9	21
114	LCO Receptors Involved in Arbuscular Mycorrhiza Are Functional for Rhizobia Perception in Legumes. <i>Current Biology</i> , 2019 , 29, 4249-4259.e5	6.3	18
113	Advances and current challenges in calcium signaling. New Phytologist, 2018, 218, 414-431	9.8	263
112	Uptake of bacteria into living plant cells, the unifying and distinct feature of the nitrogen-fixing root nodule symbiosis. <i>Current Opinion in Plant Biology</i> , 2018 , 44, 164-174	9.9	25
111	The Nuclear Pore Complex in Symbiosis and Pathogen Defence 2018 , 229-254		O
110	Phylogenomics reveals multiple losses of nitrogen-fixing root nodule symbiosis. <i>Science</i> , 2018 , 361,	33.3	167
109	The ERN1 transcription factor gene is a target of the CCaMK/CYCLOPS complex and controls rhizobial infection in Lotus japonicus. <i>New Phytologist</i> , 2017 , 215, 323-337	9.8	52
108	Cell Biology: Control of Partner Lifetime in la Plant-Fungus Relationship. Current Biology, 2017 , 27, R420)- <u>R4</u> 23	15
107	Lipid transfer from plants to arbuscular mycorrhiza fungi. ELife, 2017, 6,	8.9	206
106	A quantitative hypermorphic allele confers ectopic calcium flux and impairs cellular development. <i>ELife</i> , 2017 , 6,	8.9	23
105	Author response: Lipid transfer from plants to arbuscular mycorrhiza fungi 2017,		3
104	A CCaMK-CYCLOPS-DELLA Complex Activates Transcription of RAM1 to Regulate Arbuscule Branching. <i>Current Biology</i> , 2016 , 26, 987-98	6.3	114

103	Strawberry Accessions with Reduced Emergence From Fruits. Frontiers in Plant Science, 2016, 7, 1880	6.2	13
102	The relationship between thiamine and two symbioses: Root nodule symbiosis and arbuscular mycorrhiza. <i>Plant Signaling and Behavior</i> , 2016 , 11, e1265723	2.5	7
101	The Thiamine Biosynthesis Gene THI1 Promotes Nodule Growth and Seed Maturation. <i>Plant Physiology</i> , 2016 , 172, 2033-2043	6.6	11
100	SCARN a Novel Class of SCAR Protein That Is Required for Root-Hair Infection during Legume Nodulation. <i>PLoS Genetics</i> , 2015 , 11, e1005623	6	47
99	Colonization of root cells and plant growth promotion by Piriformospora indica occurs independently of plant common symbiosis genes. <i>Frontiers in Plant Science</i> , 2015 , 6, 667	6.2	27
98	CYCLOPS, a DNA-binding transcriptional activator, orchestrates symbiotic root nodule development. <i>Cell Host and Microbe</i> , 2014 , 15, 139-52	23.4	222
97	Knowing your friends and foesplant receptor-like kinases as initiators of symbiosis or defence. <i>New Phytologist</i> , 2014 , 204, 791-802	9.8	75
96	Cleavage of the SYMBIOSIS RECEPTOR-LIKE KINASE ectodomain promotes complex formation with Nod factor receptor 5. <i>Current Biology</i> , 2014 , 24, 422-7	6.3	83
95	A modular plasmid assembly kit for multigene expression, gene silencing and silencing rescue in plants. <i>PLoS ONE</i> , 2014 , 9, e88218	3.7	71
94	Spontaneous symbiotic reprogramming of plant roots triggered by receptor-like kinases. <i>ELife</i> , 2014 , 3,	8.9	51
93	Cell and developmental biology of arbuscular mycorrhiza symbiosis. <i>Annual Review of Cell and Developmental Biology</i> , 2013 , 29, 593-617	12.6	355
92	CERBERUS and NSP1 of Lotus japonicus are common symbiosis genes that modulate arbuscular mycorrhiza development. <i>Plant and Cell Physiology</i> , 2013 , 54, 1711-23	4.9	53
91	Host-related metabolic cues affect colonization strategies of a root endophyte. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 13965-70	11.5	118
90	Analysis of the Lotus japonicus nuclear pore NUP107-160 subcomplex reveals pronounced structural plasticity and functional redundancy. <i>Frontiers in Plant Science</i> , 2013 , 4, 552	6.2	11
89	The Nuclear Pore Complex in Symbiosis and Pathogen Defence 2013 , 229-254		5
88	Molecular Mechanisms Governing Arbuscular Mycorrhiza Development and Function 2013, 457-465		1
87	The K+-dependent asparaginase, NSE1, is crucial for plant growth and seed production in Lotus japonicus. <i>Plant and Cell Physiology</i> , 2013 , 54, 107-18	4.9	30
86	Two Lotus japonicus symbiosis mutants impaired at distinct steps of arbuscule development. <i>Plant Journal</i> , 2013 , 75, 117-129	6.9	13

85	Negative regulation of CCaMK is essential for symbiotic infection. <i>Plant Journal</i> , 2012 , 72, 572-84	6.9	39
84	Polymorphic infection and organogenesis patterns induced by a Rhizobium leguminosarum isolate from Lotus root nodules are determined by the host genotype. <i>New Phytologist</i> , 2012 , 196, 561-573	9.8	27
83	Activation of calcium- and calmodulin-dependent protein kinase (CCaMK), the central regulator of plant root endosymbiosis. <i>Current Opinion in Plant Biology</i> , 2012 , 15, 444-53	9.9	116
82	Receptor kinase signaling pathways in plant-microbe interactions. <i>Annual Review of Phytopathology</i> , 2012 , 50, 451-73	10.8	157
81	The recent evolution of a symbiotic ion channel in the legume family altered ion conductance and improved functionality in calcium signaling. <i>Plant Cell</i> , 2012 , 24, 2528-45	11.6	51
80	Two microRNAs linked to nodule infection and nitrogen-fixing ability in the legume Lotus japonicus. <i>Plant Physiology</i> , 2012 , 160, 2137-54	6.6	90
79	Lotus japonicus E3 ligase SEVEN IN ABSENTIA4 destabilizes the symbiosis receptor-like kinase SYMRK and negatively regulates rhizobial infection. <i>Plant Cell</i> , 2012 , 24, 1691-707	11.6	60
78	FRET-based genetically encoded sensors allow high-resolution live cell imaging of Call+ dynamics. <i>Plant Journal</i> , 2012 , 69, 181-92	6.9	175
77	RNA-seq pinpoints a Xanthomonas TAL-effector activated resistance gene in a large-crop genome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, 19480-5	11.5	80
76	A set of Lotus japonicus Gifu x Lotus burttii recombinant inbred lines facilitates map-based cloning and QTL mapping. <i>DNA Research</i> , 2012 , 19, 317-23	4.5	22
75	Functional domain analysis of the Remorin protein LjSYMREM1 in Lotus japonicus. <i>PLoS ONE</i> , 2012 , 7, e30817	3.7	69
74	Phosphoproteome analysis of Lotus japonicus roots reveals shared and distinct components of symbiosis and defense. <i>Molecular Plant-Microbe Interactions</i> , 2011 , 24, 932-7	3.6	27
73	Activation of a Lotus japonicus subtilase gene during arbuscular mycorrhiza is dependent on the common symbiosis genes and two cis-active promoter regions. <i>Molecular Plant-Microbe Interactions</i> , 2011 , 24, 662-70	3.6	19
72	Autophosphorylation is essential for the in vivo function of the Lotus japonicus Nod factor receptor 1 and receptor-mediated signalling in cooperation with Nod factor receptor 5. <i>Plant Journal</i> , 2011 , 65, 404-17	6.9	135
71	The Clavata2 genes of pea and Lotus japonicus affect autoregulation of nodulation. <i>Plant Journal</i> , 2011 , 65, 861-71	6.9	89
70	Lotus japonicus symRK-14 uncouples the cortical and epidermal symbiotic program. <i>Plant Journal</i> , 2011 , 67, 929-40	6.9	57
69	NENA, a Lotus japonicus homolog of Sec13, is required for rhizodermal infection by arbuscular mycorrhiza fungi and rhizobia but dispensable for cortical endosymbiotic development. <i>Plant Cell</i> , 2010 , 22, 2509-26	11.6	174
68	A suite of Lotus japonicus starch mutants reveals both conserved and novel features of starch metabolism. <i>Plant Physiology</i> , 2010 , 154, 643-55	6.6	49

(2006-2009)

67	A cytosolic invertase is required for normal growth and cell development in the model legume, Lotus japonicus. <i>Journal of Experimental Botany</i> , 2009 , 60, 3353-65	7	76
66	TILLING in Lotus japonicus identified large allelic series for symbiosis genes and revealed a bias in functionally defective ethyl methanesulfonate alleles toward glycine replacements. <i>Plant Physiology</i> , 2009 , 151, 1281-91	6.6	68
65	The temperature-sensitive brush mutant of the legume Lotus japonicus reveals a link between root development and nodule infection by rhizobia. <i>Plant Physiology</i> , 2009 , 149, 1785-96	6.6	19
64	The unbearable naivety of legumes in symbiosis. Current Opinion in Plant Biology, 2009, 12, 491-9	9.9	55
63	Apoplastic plant subtilases support arbuscular mycorrhiza development in Lotus japonicus. <i>Plant Journal</i> , 2009 , 58, 766-77	6.9	95
62	Evolution of root endosymbiosis with bacteria: How novel are nodules?. <i>Trends in Plant Science</i> , 2009 , 14, 77-86	13.1	159
61	Identification of candidate genome regions controlling disease resistance in Arachis. <i>BMC Plant Biology</i> , 2009 , 9, 112	5.3	98
60	Lotus japonicus CASTOR and POLLUX are ion channels essential for perinuclear calcium spiking in legume root endosymbiosis. <i>Plant Cell</i> , 2008 , 20, 3467-79	11.6	167
59	Arbuscular mycorrhiza: the mother of plant root endosymbioses. <i>Nature Reviews Microbiology</i> , 2008 , 6, 763-75	22.2	1316
58	SymRK defines a common genetic basis for plant root endosymbioses with arbuscular mycorrhiza fungi, rhizobia, and Frankiabacteria. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008 , 105, 4928-32	11.5	227
57	CYCLOPS, a mediator of symbiotic intracellular accommodation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008 , 105, 20540-5	11.5	321
56	Functional adaptation of a plant receptor-kinase paved the way for the evolution of intracellular root symbioses with bacteria. <i>PLoS Biology</i> , 2008 , 6, e68	9.7	179
55	Proteases in plant root symbiosis. <i>Phytochemistry</i> , 2007 , 68, 111-21	4	35
54	NUCLEOPORIN85 is required for calcium spiking, fungal and bacterial symbioses, and seed production in Lotus japonicus. <i>Plant Cell</i> , 2007 , 19, 610-24	11.6	274
53	TILLING mutants of Lotus japonicus reveal that nitrogen assimilation and fixation can occur in the absence of nodule-enhanced sucrose synthase. <i>Plant Physiology</i> , 2007 , 144, 806-20	6.6	83
52	Lotus japonicus nodulation requires two GRAS domain regulators, one of which is functionally conserved in a non-legume. <i>Plant Physiology</i> , 2006 , 142, 1739-50	6.6	201
51	Transcriptome analysis of Arabidopsis clubroots indicate a key role for cytokinins in disease development. <i>Molecular Plant-Microbe Interactions</i> , 2006 , 19, 480-94	3.6	245
50	The most widespread symbiosis on Earth. <i>PLoS Biology</i> , 2006 , 4, e239	9.7	59

49	Genetics of symbiosis in Lotus japonicus: recombinant inbred lines, comparative genetic maps, and map position of 35 symbiotic loci. <i>Molecular Plant-Microbe Interactions</i> , 2006 , 19, 80-91	3.6	87
48	Genetic suppressors of the Lotus japonicus har1-1 hypernodulation phenotype. <i>Molecular Plant-Microbe Interactions</i> , 2006 , 19, 1082-91	3.6	41
47	Identification of symbiotically defective mutants of Lotus japonicus affected in infection thread growth. <i>Molecular Plant-Microbe Interactions</i> , 2006 , 19, 1444-50	3.6	28
46	Deregulation of a Ca2+/calmodulin-dependent kinase leads to spontaneous nodule development. <i>Nature</i> , 2006 , 441, 1153-6	50.4	359
45	Common symbiosis genes of Lotus japonicus are not required for intracellular accommodation of the rust fungus Uromyces loti. <i>New Phytologist</i> , 2006 , 170, 641-4	9.8	13
44	Lotus japonicus: legume research in the fast lane. <i>Trends in Plant Science</i> , 2005 , 10, 222-8	13.1	110
43	Arbuscular mycorrhiza 2005 , 87-95		7
42	Plastid proteins crucial for symbiotic fungal and bacterial entry into plant roots. <i>Nature</i> , 2005 , 433, 527	'- 3 5'b.4	346
41	Regulation of plant symbiosis receptor kinase through serine and threonine phosphorylation. <i>Journal of Biological Chemistry</i> , 2005 , 280, 9203-9	5.4	75
40	Seven Lotus japonicus genes required for transcriptional reprogramming of the root during fungal and bacterial symbiosis. <i>Plant Cell</i> , 2005 , 17, 2217-29	11.6	252
39	Regulatory Mechanisms of Symrk Kinase Activity 2005 , 183-185		1
38	96-Well DNA isolation method 2005 , 129-131		1
37	Mutagenesis 2005 , 177-186		2
36	Tilling 2005 , 197-210		5
35	Genetic variation at the tomato Cf-4/Cf-9 locus induced by EMS mutagenesis and intralocus recombination. <i>Genetics</i> , 2004 , 167, 459-70	4	30
34	Distinct roles of Lotus japonicus SYMRK and SYM15 in root colonization and arbuscule formation. <i>New Phytologist</i> , 2004 , 163, 381-392	9.8	86
33	Molecular genetics of the arbuscular mycorrhizal symbiosis. <i>Current Opinion in Plant Biology</i> , 2004 , 7, 414-21	9.9	149
32	Exploitation of colinear relationships between the genomes of Lotus japonicus, Pisum sativum and Arabidopsis thaliana, for positional cloning of a legume symbiosis gene. <i>Theoretical and Applied Genetics</i> , 2004 , 108, 442-9	6	39

31	A TILLING reverse genetics tool and a web-accessible collection of mutants of the legume Lotus japonicus. <i>Plant Physiology</i> , 2003 , 131, 866-71	6.6	291
30	Dual requirement of the LjSym4 gene for mycorrhizal development in epidermal and cortical cells of Lotus japonicus roots. <i>New Phytologist</i> , 2002 , 154, 741-749	9.8	74
29	A plant receptor-like kinase required for both bacterial and fungal symbiosis. <i>Nature</i> , 2002 , 417, 959-62	50.4	737
28	Phase State of a Steel Plate Caused by the Action of Distributed Heat Sources. <i>Materials Science</i> , 2002 , 38, 214-219	0.7	
27	Evolution of signal transduction in intracellular symbiosis. <i>Trends in Plant Science</i> , 2002 , 7, 511-8	13.1	286
26	A genetic linkage map of the model legume Lotus japonicus and strategies for fast mapping of new loci. <i>Genetics</i> , 2002 , 161, 1673-83	4	66
25	The Lotus japonicus LjSym4 gene is required for the successful symbiotic infection of root epidermal cells. <i>Molecular Plant-Microbe Interactions</i> , 2000 , 13, 1109-20	3.6	120
24	Intracellular accommodation of microbes by plants: a common developmental program for symbiosis and disease?. <i>Current Opinion in Plant Biology</i> , 2000 , 3, 320-8	9.9	26 0
23	Pronounced Intraspecific Haplotype Divergence at the RPP5 Complex Disease Resistance Locus of Arabidopsis. <i>Plant Cell</i> , 1999 , 11, 2099-2111	11.6	286
22	Recombination between diverged clusters of the tomato Cf-9 plant disease resistance gene family. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1999 , 96, 5850-5	11.5	102
21	Homologues of the Cf-9 disease resistance gene (Hcr9s) are present at multiple loci on the short arm of tomato chromosome 1. <i>Molecular Plant-Microbe Interactions</i> , 1999 , 12, 93-102	3.6	45
20	Genetic nomenclature guidelines for the model legume Lotus japonicus. <i>Trends in Plant Science</i> , 1999 , 4, 300-301	13.1	11
19	Pronounced Intraspecific Haplotype Divergence at the RPP5 Complex Disease Resistance Locus of Arabidopsis. <i>Plant Cell</i> , 1999 , 11, 2099	11.6	5
18	Exopolysaccharide (EPS) synthesis in Bradyrhizobium japonicum: sequence, operon structure and mutational analysis of an exo gene cluster. <i>Molecular Genetics and Genomics</i> , 1998 , 259, 161-71		27
17	Genetic and molecular analysis of tomato Cf genes for resistance to Cladosporium fulvum. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 1998 , 353, 1413-24	5.8	62
16	Mycorrhiza Mutants of Lotus japonicus Define Genetically Independent Steps During Symbiotic Infection. <i>Molecular Plant-Microbe Interactions</i> , 1998 , 11, 933-936	3.6	106
15	Characterization of the Tomato Cf-4 Gene for Resistance to Cladosporium fulvum Identifies Sequences That Determine Recognitional Specificity in Cf-4 and Cf-9. <i>Plant Cell</i> , 1997 , 9, 2209	11.6	39
14	Novel disease resistance specificities result from sequence exchange between tandemly repeated genes at the Cf-4/9 locus of tomato. <i>Cell</i> , 1997 , 91, 821-32	56.2	507

13	reversed by the plant flavone luteolin. <i>Bulletin of Environmental Contamination and Toxicology</i> , 1995 , 54, 633-9	2.7	5
12	Modes of expression and common structural features of the complete phenylalanine ammonia-lyase gene family in parsley. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1995 , 92, 5905-9	11.5	212
11	Oligopeptide elicitor-mediated defense gene activation in cultured parsley cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1995 , 92, 4150-7	11.5	190
10	The Impairment of the Nodulation Process, Induced by a Bradyrhizobium japonicum Exopolysaccharide Mutant is Determined by the Genotype of the Host Plant. <i>Zeitschrift Fur Naturforschung - Section C Journal of Biosciences</i> , 1994 , 49, 727-736	1.7	7
9	Competitiveness and communication for effective inoculation byRhizobium, Bradyrhizobium and vesicular-arbuscular mycorrhiza fungi. <i>Experientia</i> , 1994 , 50, 884-889		9
8	Plant Defense Responses of Host Plants with Determinate Nodules Induced by EPS-DefectiveexoBMutants ofBradyrhizobium japonicum. <i>Molecular Plant-Microbe Interactions</i> , 1994 , 7, 631	3.6	68
7	Legume root metabolites and VA-mycorrhiza development. <i>Journal of Plant Physiology</i> , 1993 , 141, 54-60	03.6	38
6	ExoB Mutants ofBradyrhizobium japonicumwith Reduced Competitiveness for Nodulation ofGlycine max. <i>Molecular Plant-Microbe Interactions</i> , 1993 , 6, 99	3.6	35
5	Production of the Phytoalexin Glyceollin I by Soybean Roots in Response to Symbiotic and Pathogenic Infection. <i>Botanica Acta</i> , 1992 , 105, 18-25		48
4	Accumulation of the Phytoalexin Glyceollin I in Soybean Nodules Infected by a Bradyrhizobium japonicum nifA Mutant. <i>Zeitschrift Fur Naturforschung - Section C Journal of Biosciences</i> , 1991 , 46, 318-32	26 ^{.7}	26
3	Chemotaxis and nod Gene Activity of Bradyrhizobium japonicum in Response to Hydroxycinnamic Acids and Isoflavonoids. <i>Applied and Environmental Microbiology</i> , 1991 , 57, 316-9	4.8	74
2	Hypersensitive Reaction of Nodule Cells in the Glycine sp./Bradyrhizobium japonicum-Symbiosis Occurs at the Genotype-Specific Level*. <i>Botanica Acta</i> , 1990 , 103, 143-148		39
1	Symbiosis-related genes sustain the development of a downy mildew pathogen on Arabidopsis thaliana		3