

Roberto Solano

List of Publications by Year in Descending Order

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

108
papers

18,174
citations

55
h-index

115
g-index

115
ext. papers

21,893
ext. citations

10.4
avg, IF

6.47
L-index

#	Paper	IF	Citations
108	An evolutionarily ancient fatty acid desaturase is required for the synthesis of hexadecatrienoic acid, which is the main source of the bioactive jasmonate in <i>Marchantia polymorpha</i> . <i>New Phytologist</i> , 2021 , 233, 1401	9.8	1
107	SARS-CoV-2 Fears Green: The Chlorophyll Catabolite Pheophorbide A Is a Potent Antiviral. <i>Pharmaceuticals</i> , 2021 , 14,	5.2	1
106	Redox Feedback regulation of ANAC089 signaling alters seed germination and stress response. <i>Cell Reports</i> , 2021 , 35, 109263	10.6	1
105	Potato CYCLING DOF FACTOR1 and its lncRNA counterpart StFLORE link tuber development and drought response. <i>Plant Journal</i> , 2021 , 105, 855-869	6.9	13
104	DNA features beyond the transcription factor binding site specify target recognition by plant MYC2-related bHLH proteins. <i>Plant Communications</i> , 2021 , 2, 100232	9	0
103	A small molecule antagonizes jasmonic acid perception and auxin responses in vascular and nonvascular plants. <i>Plant Physiology</i> , 2021 , 187, 1399-1413	6.6	0
102	<i>Marchantia polymorpha</i> model reveals conserved infection mechanisms in the vascular wilt fungal pathogen <i>Fusarium oxysporum</i> . <i>New Phytologist</i> , 2021 ,	9.8	3
101	Interactions of JAZ Repressors with Anthocyanin Biosynthesis-Related Transcription Factors of <i>Fragaria</i> and <i>Ananassa</i> . <i>Agronomy</i> , 2020 , 10, 1586	3.6	2
100	An Ancient COI1-Independent Function for Reactive Electrophilic Oxylipins in Thermotolerance. <i>Current Biology</i> , 2020 , 30, 962-971.e3	6.3	29
99	Integrated multi-omics framework of the plant response to jasmonic acid. <i>Nature Plants</i> , 2020 , 6, 290-302	11.5	59
98	CUL3 E3 ubiquitin ligases regulate MYC2, MYC3, and MYC4 stability and JA responses. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 6205-6215	11.5	30
97	Isolation of Natural Fungal Pathogens from <i>Marchantia polymorpha</i> Reveals Antagonism between Salicylic Acid and Jasmonate during Liverwort-Fungus Interactions. <i>Plant and Cell Physiology</i> , 2020 , 61, 265-275	4.9	12
96	The JA-pathway MYC transcription factors regulate photomorphogenic responses by targeting HY5 gene expression. <i>Plant Journal</i> , 2020 , 102, 138-152	6.9	24
95	A new functional JAZ degron sequence in strawberry JAZ1 revealed by structural and interaction studies on the COI1-JA-Ile/COR-JAZs complexes. <i>Scientific Reports</i> , 2020 , 10, 11310	4.9	9
94	Characterizing the involvement of FaMADS9 in the regulation of strawberry fruit receptacle development. <i>Plant Biotechnology Journal</i> , 2020 , 18, 929-943	11.6	10
93	Omega hydroxylated JA-Ile is an endogenous bioactive jasmonate that signals through the canonical jasmonate signaling pathway. <i>Biochimica Et Biophysica Acta - Molecular and Cell Biology of Lipids</i> , 2019 , 1864, 158520	5	12
92	PIF transcription factors link a neighbor threat cue to accelerated reproduction in <i>Arabidopsis</i> . <i>Nature Communications</i> , 2019 , 10, 4005	17.4	39

91	A Single JAZ Repressor Controls the Jasmonate Pathway in <i>Marchantia polymorpha</i> . <i>Molecular Plant</i> , 2019 , 12, 185-198	14.4	55
90	Jasmonate-Related MYC Transcription Factors Are Functionally Conserved in. <i>Plant Cell</i> , 2019 , 31, 2491-2509	15.0	31
89	An Evolutionarily Ancient Immune System Governs the Interactions between <i>Pseudomonas syringae</i> and an Early-Diverging Land Plant Lineage. <i>Current Biology</i> , 2019 , 29, 2270-2281.e4	6.3	20
88	Design of a bacterial speck resistant tomato by CRISPR/Cas9-mediated editing of SLJAZ2. <i>Plant Biotechnology Journal</i> , 2019 , 17, 665-673	11.6	121
87	Ligand-receptor co-evolution shaped the jasmonate pathway in land plants. <i>Nature Chemical Biology</i> , 2018 , 14, 480-488	11.7	105
86	An OPR3-independent pathway uses 4,5-didehydrojasmonate for jasmonate synthesis. <i>Nature Chemical Biology</i> , 2018 , 14, 171-178	11.7	106
85	Arabidopsis SWC4 Binds DNA and Recruits the SWR1 Complex to Modulate Histone H2A.Z Deposition at Key Regulatory Genes. <i>Molecular Plant</i> , 2018 , 11, 815-832	14.4	32
84	The fungal phytotoxin lasiojasmonate A activates the plant jasmonic acid pathway. <i>Journal of Experimental Botany</i> , 2018 , 69, 3095-3102	7	21
83	Coordination of Chloroplast Development through the Action of the GNC and GLK Transcription Factor Families. <i>Plant Physiology</i> , 2018 , 178, 130-147	6.6	30
82	Characterization of wheat (<i>Triticum aestivum</i>) TIFY family and role of <i>Triticum Durum</i> TdTIFY11a in salt stress tolerance. <i>PLoS ONE</i> , 2018 , 13, e0200566	3.7	32
81	A rationally designed JAZ subtype-selective agonist of jasmonate perception. <i>Nature Communications</i> , 2018 , 9, 3654	17.4	25
80	Identification of plant transcription factor target sequences. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2017 , 1860, 21-30	6	24
79	The transcription factor bZIP14 regulates the TCA cycle in the diatom. <i>EMBO Journal</i> , 2017 , 36, 1559-1576	15.9	39
78	JAZ2 controls stomata dynamics during bacterial invasion. <i>New Phytologist</i> , 2017 , 213, 1378-1392	9.8	80
77	Insights into Land Plant Evolution Garnered from the <i>Marchantia polymorpha</i> Genome. <i>Cell</i> , 2017 , 171, 287-304.e15	56.2	538
76	The WRKY3 Transcription Factor SLWRKY3 Is Involved in Salt Stress Tolerance in Tomato. <i>Frontiers in Plant Science</i> , 2017 , 8, 1343	6.2	57
75	Assessing the Role of ETHYLENE RESPONSE FACTOR Transcriptional Repressors in Salicylic Acid-Mediated Suppression of Jasmonic Acid-Responsive Genes. <i>Plant and Cell Physiology</i> , 2017 , 58, 266-278	4.9	41
74	A non-DNA-binding activity for the ATHB4 transcription factor in the control of vegetation proximity. <i>New Phytologist</i> , 2017 , 216, 798-813	9.8	8

73	The fruit-specific transcription factor FaDOF2 regulates the production of eugenol in ripe fruit receptacles. <i>Journal of Experimental Botany</i> , 2017 , 68, 4529-4543	7	17
72	Cytokinin induces genome-wide binding of the type-B response regulator ARR10 to regulate growth and development in. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017 , 114, E5995-E6004	11.5	99
71	Redundancy and specificity in jasmonate signalling. <i>Current Opinion in Plant Biology</i> , 2016 , 33, 147-156	9.9	193
70	The Venus Flytrap <i>Dionaea muscipula</i> Counts Prey-Induced Action Potentials to Induce Sodium Uptake. <i>Current Biology</i> , 2016 , 26, 286-95	6.3	92
69	The Naming of Names: Guidelines for Gene Nomenclature in Marchantia. <i>Plant and Cell Physiology</i> , 2016 , 57, 257-61	4.9	38
68	How Microbes Twist Jasmonate Signaling around Their Little Fingers. <i>Plants</i> , 2016 , 5,	4.5	41
67	Characterization of the cytokinin-responsive transcriptome in rice. <i>BMC Plant Biology</i> , 2016 , 16, 260	5.3	27
66	The cytokinin response factors modulate root and shoot growth and promote leaf senescence in Arabidopsis. <i>Plant Journal</i> , 2016 , 85, 134-47	6.9	72
65	Deciphering the Molecular Mechanisms Underpinning the Transcriptional Control of Gene Expression by Master Transcriptional Regulators in Arabidopsis Seed. <i>Plant Physiology</i> , 2016 , 171, 1099-112	6.6	43
64	An R2R3-MYB Transcription Factor Regulates Eugenol Production in Ripe Strawberry Fruit Receptacles. <i>Plant Physiology</i> , 2015 , 168, 598-614	6.6	62
63	The RING E3 Ligase KEEP ON GOING Modulates JASMONATE ZIM-DOMAIN12 Stability. <i>Plant Physiology</i> , 2015 , 169, 1405-17	6.6	50
62	FILAMENTOUS FLOWER Is a Direct Target of JAZ3 and Modulates Responses to Jasmonate. <i>Plant Cell</i> , 2015 , 27, 3160-74	11.6	55
61	A MYB/ZML Complex Regulates Wound-Induced Lignin Genes in Maize. <i>Plant Cell</i> , 2015 , 27, 3245-59	11.6	55
60	Novel players fine-tune plant trade-offs. <i>Essays in Biochemistry</i> , 2015 , 58, 83-100	7.6	30
59	DNA-binding specificities of plant transcription factors and their potential to define target genes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 2367-72	11.5	423
58	Structural basis for DNA binding specificity by the auxin-dependent ARF transcription factors. <i>Cell</i> , 2014 , 156, 577-89	56.2	243
57	Rational design of a ligand-based antagonist of jasmonate perception. <i>Nature Chemical Biology</i> , 2014 , 10, 671-6	11.7	63
56	The <i>Solanum lycopersicum</i> Zinc Finger2 cysteine-2/histidine-2 repressor-like transcription factor regulates development and tolerance to salinity in tomato and Arabidopsis. <i>Plant Physiology</i> , 2014 , 164, 1967-90	6.6	41

55	Repression of Jasmonate-Dependent Defenses by Shade Involves Differential Regulation of Protein Stability of MYC Transcription Factors and Their JAZ Repressors in Arabidopsis. <i>Plant Cell</i> , 2014 , 26, 1967-1980	11.6	125
54	High-throughput analysis of protein-DNA binding affinity. <i>Methods in Molecular Biology</i> , 2014 , 1062, 697-709	7.0	6
53	The bacterial effector HopX1 targets JAZ transcriptional repressors to activate jasmonate signaling and promote infection in Arabidopsis. <i>PLoS Biology</i> , 2014 , 12, e1001792	9.7	163
52	The TRANSPLANTA collection of Arabidopsis lines: a resource for functional analysis of transcription factors based on their conditional overexpression. <i>Plant Journal</i> , 2014 , 77, 944-53	6.9	61
51	bHLH003, bHLH013 and bHLH017 are new targets of JAZ repressors negatively regulating JA responses. <i>PLoS ONE</i> , 2014 , 9, e86182	3.7	78
50	Distinct and conserved transcriptomic changes during nematode-induced giant cell development in tomato compared with Arabidopsis: a functional role for gene repression. <i>New Phytologist</i> , 2013 , 197, 1276-1290	9.8	76
49	Pull-down analysis of interactions among jasmonic acid core signaling proteins. <i>Methods in Molecular Biology</i> , 2013 , 1011, 159-71	1.4	12
48	Protein intrinsic disorder in plants. <i>Frontiers in Plant Science</i> , 2013 , 4, 363	6.2	43
47	Nuclear jasmonate and salicylate signaling and crosstalk in defense against pathogens. <i>Frontiers in Plant Science</i> , 2013 , 4, 72	6.2	106
46	Arabidopsis basic helix-loop-helix transcription factors MYC2, MYC3, and MYC4 regulate glucosinolate biosynthesis, insect performance, and feeding behavior. <i>Plant Cell</i> , 2013 , 25, 3117-32	11.6	313
45	Genome-wide analysis of protein disorder in Arabidopsis thaliana: implications for plant environmental adaptation. <i>PLoS ONE</i> , 2013 , 8, e55524	3.7	44
44	Phytochrome interacting factors 4 and 5 control seedling growth in changing light conditions by directly controlling auxin signaling. <i>Plant Journal</i> , 2012 , 71, 699-711	6.9	383
43	Target validation of plant microRNAs. <i>Methods in Molecular Biology</i> , 2011 , 732, 187-208	1.4	20
42	Improved protein-binding microarrays for the identification of DNA-binding specificities of transcription factors. <i>Plant Journal</i> , 2011 , 66, 700-11	6.9	103
41	Genome-wide mapping of Arabidopsis thaliana origins of DNA replication and their associated epigenetic marks. <i>Nature Structural and Molecular Biology</i> , 2011 , 18, 395-400	17.6	105
40	The Arabidopsis bHLH transcription factors MYC3 and MYC4 are targets of JAZ repressors and act additively with MYC2 in the activation of jasmonate responses. <i>Plant Cell</i> , 2011 , 23, 701-15	11.6	700
39	Early transcriptomic events in microdissected Arabidopsis nematode-induced giant cells. <i>Plant Journal</i> , 2010 , 61, 698-712	6.9	173
38	NINJA connects the co-repressor TOPLESS to jasmonate signalling. <i>Nature</i> , 2010 , 464, 788-91	50.4	679

37	A central regulatory system largely controls transcriptional activation and repression responses to phosphate starvation in Arabidopsis. <i>PLoS Genetics</i> , 2010 , 6, e1001102	6	408
36	Mitochondrial beta-cyanoalanine synthase is essential for root hair formation in Arabidopsis thaliana. <i>Plant Cell</i> , 2010 , 22, 3268-79	11.6	84
35	The jasmonate pathway: the ligand, the receptor and the core signalling module. <i>Current Opinion in Plant Biology</i> , 2009 , 12, 539-47	9.9	248
34	The ZIM domain mediates homo- and heteromeric interactions between Arabidopsis JAZ proteins. <i>Plant Journal</i> , 2009 , 59, 77-87	6.9	213
33	Isolation of RNA from laser-capture-microdissected giant cells at early differentiation stages suitable for differential transcriptome analysis. <i>Molecular Plant Pathology</i> , 2009 , 10, 523-35	5.7	33
32	Bacterial chemoattraction towards jasmonate plays a role in the entry of <i>Dickeya dadantii</i> through wounded tissues. <i>Molecular Microbiology</i> , 2009 , 74, 662-71	4.1	38
31	Bacterial chemoattraction towards jasmonate plays a role in the entry of <i>Dickeya dadantii</i> through wounded tissues. <i>Molecular Microbiology</i> , 2009 , 74, 1543-1543	4.1	1
30	Genome-wide identification of small RNA targets based on target enrichment and microarray hybridizations. <i>Plant Journal</i> , 2009 , 59, 840-50	6.9	18
29	(+)-7-iso-Jasmonoyl-L-iso-leucine is the endogenous bioactive jasmonate. <i>Nature Chemical Biology</i> , 2009 , 5, 344-50	11.7	669
28	Plant oxylipins: COI1/JAZs/MYC2 as the core jasmonic acid-signalling module. <i>FEBS Journal</i> , 2009 , 276, 4682-92	5.7	133
27	Reactome array: forging a link between metabolome and genome. <i>Science</i> , 2009 , 326, 252-7	33.3	30
26	JAZ repressors set the rhythm in jasmonate signaling. <i>Current Opinion in Plant Biology</i> , 2008 , 11, 486-94	9.9	182
25	The JAZ family of repressors is the missing link in jasmonate signalling. <i>Nature</i> , 2007 , 448, 666-71	50.4	1576
24	A protein phosphatase 2A catalytic subunit is a negative regulator of abscisic acid signalling. <i>Plant Journal</i> , 2007 , 51, 763-78	6.9	85
23	Modulation of Plant Defenses by Ethylene. <i>Journal of Plant Growth Regulation</i> , 2007 , 26, 160-177	4.7	101
22	A mutant of the Arabidopsis phosphate transporter PHT1;1 displays enhanced arsenic accumulation. <i>Plant Cell</i> , 2007 , 19, 1123-33	11.6	246
21	Heart-rate deflection point and the second heart-rate variability threshold during running exercise in trained boys. <i>Pediatric Exercise Science</i> , 2007 , 19, 192-204	2	38
20	ABA is an essential signal for plant resistance to pathogens affecting JA biosynthesis and the activation of defenses in Arabidopsis. <i>Plant Cell</i> , 2007 , 19, 1665-81	11.6	621

19	Molecular players regulating the jasmonate signalling network. <i>Current Opinion in Plant Biology</i> , 2005 , 8, 532-40	9.9	381
18	PHOSPHATE TRANSPORTER TRAFFIC FACILITATOR1 is a plant-specific SEC12-related protein that enables the endoplasmic reticulum exit of a high-affinity phosphate transporter in Arabidopsis. <i>Plant Cell</i> , 2005 , 17, 3500-12	11.6	221
17	JASMONATE-INSENSITIVE1 encodes a MYC transcription factor essential to discriminate between different jasmonate-regulated defense responses in Arabidopsis. <i>Plant Cell</i> , 2004 , 16, 1938-50	11.6	925
16	Five components of the ethylene-response pathway identified in a screen for weak ethylene-insensitive mutants in Arabidopsis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003 , 100, 2992-7	11.5	312
15	Interactions Between Signaling Compounds Involved in Plant Defense. <i>Journal of Plant Growth Regulation</i> , 2003 , 22, 82-98	4.7	177
14	ETHYLENE RESPONSE FACTOR1 integrates signals from ethylene and jasmonate pathways in plant defense. <i>Plant Cell</i> , 2003 , 15, 165-78	11.6	913
13	Constitutive expression of ETHYLENE-RESPONSE-FACTOR1 in Arabidopsis confers resistance to several necrotrophic fungi. <i>Plant Journal</i> , 2002 , 29, 23-32	6.9	580
12	Mutations at CRE1 impair cytokinin-induced repression of phosphate starvation responses in Arabidopsis. <i>Plant Journal</i> , 2002 , 32, 353-60	6.9	149
11	A conserved MYB transcription factor involved in phosphate starvation signaling both in vascular plants and in unicellular algae. <i>Genes and Development</i> , 2001 , 15, 2122-33	12.6	858
10	Influence of cytokinins on the expression of phosphate starvation responsive genes in Arabidopsis. <i>Plant Journal</i> , 2000 , 24, 559-67	6.9	315
9	Ethylene gas: perception, signaling and response. <i>Current Opinion in Plant Biology</i> , 1998 , 1, 393-8	9.9	72
8	Arabidopsis homologs of a c-Jun coactivator are present both in monomeric form and in the COP9 complex, and their abundance is differentially affected by the pleiotropic cop/det/fus mutations. <i>Plant Cell</i> , 1998 , 10, 1779-90	11.6	144
7	Nuclear events in ethylene signaling: a transcriptional cascade mediated by ETHYLENE-INSENSITIVE3 and ETHYLENE-RESPONSE-FACTOR1. <i>Genes and Development</i> , 1998 , 12, 3703-14	12.6	925
6	A single residue substitution causes a switch from the dual DNA binding specificity of plant transcription factor MYB.Ph3 to the animal c-MYB specificity. <i>Journal of Biological Chemistry</i> , 1997 , 272, 2889-95	5.4	39
5	Activation of the ethylene gas response pathway in Arabidopsis by the nuclear protein ETHYLENE-INSENSITIVE3 and related proteins. <i>Cell</i> , 1997 , 89, 1133-44	56.2	776
4	MYB.Ph3 transcription factor from <i>Petunia hybrida</i> induces similar DNA-bending/distortions on its two types of binding site. <i>Plant Journal</i> , 1995 , 8, 673-82	6.9	21
3	Dual DNA binding specificity of a petal epidermis-specific MYB transcription factor (MYB.Ph3) from <i>Petunia hybrida</i> .. <i>EMBO Journal</i> , 1995 , 14, 1773-1784	13	157
2	Organization of repeated sequences in species of the genus <i>Avena</i> . <i>Theoretical and Applied Genetics</i> , 1992 , 83, 602-7	6	36

1 A small molecule antagonizes jasmonic acid perception and auxin responses in vascular and non-vascular plants

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