

Manuel Martinez

List of Publications by Year in descending order

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89
papers

4,546
citations

117453

34
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110170

64
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all docs

89
docs citations

89
times ranked

4509
citing authors

#	ARTICLE	IF	CITATIONS
1	Hydroxynitrile lyase defends Arabidopsis against <i>Tetranychus urticae</i> . <i>Plant Physiology</i> , 2022, 189, 2244-2258.	2.3	9
2	Plant Kinases in the Perception and Signaling Networks Associated With Arthropod Herbivory. <i>Frontiers in Plant Science</i> , 2022, 13, .	1.7	5
3	Comparative transcriptomics reveals hidden issues in the plant response to arthropod herbivores. <i>Journal of Integrative Plant Biology</i> , 2021, 63, 312-326.	4.1	19
4	Repression of barley cathepsins, HvPap-19 and HvPap-1, differentially alters grain composition and delays germination. <i>Journal of Experimental Botany</i> , 2021, 72, 3474-3485.	2.4	4
5	Disentangling transcriptional responses in plant defense against arthropod herbivores. <i>Scientific Reports</i> , 2021, 11, 12996.	1.6	9
6	The co-chaperone HOP3 participates in jasmonic acid signaling by regulating CORONATINE-INSENSITIVE 1 activity. <i>Plant Physiology</i> , 2021, 187, 1679-1689.	2.3	7
7	Spider mite egg extract modifies Arabidopsis response to future infestations. <i>Scientific Reports</i> , 2021, 11, 17692.	1.6	5
8	Saving time maintaining reliability: a new method for quantification of <i>Tetranychus urticae</i> damage in Arabidopsis whole rosettes. <i>BMC Plant Biology</i> , 2020, 20, 397.	1.6	11
9	Factores de riesgo y protecci3n del estr3s traum3tico secundario en los cuidados intensivos: un estudio exploratorio en un hospital terciario de Madrid. <i>Medicina Intensiva</i> , 2020, 44, 420-428.	0.4	2
10	Plant Defenses Against <i>Tetranychus urticae</i> : Mind the Gaps. <i>Plants</i> , 2020, 9, 464.	1.6	43
11	The Price of the Induced Defense Against Pests: A Meta-Analysis. <i>Frontiers in Plant Science</i> , 2020, 11, 615122.	1.7	20
12	Plant Defenses Against Pests Driven by a Bidirectional Promoter. <i>Frontiers in Plant Science</i> , 2019, 10, 930.	1.7	6
13	Editorial for Special Issue "Molecular Advances in Wheat and Barley" <i>International Journal of Molecular Sciences</i> , 2019, 20, 3501.	1.8	10
14	Plant Proteases: From Key Enzymes in Germination to Allies for Fighting Human Gluten-Related Disorders. <i>Frontiers in Plant Science</i> , 2019, 10, 721.	1.7	30
15	Insights on the Proteases Involved in Barley and Wheat Grain Germination. <i>International Journal of Molecular Sciences</i> , 2019, 20, 2087.	1.8	31
16	An Arabidopsis TIR-Lectin Two-Domain Protein Confers Defense Properties against <i>Tetranychus urticae</i> . <i>Plant Physiology</i> , 2019, 179, 1298-1314.	2.3	38
17	Repression of drought-induced cysteine-protease genes alters barley leaf structure and responses to abiotic and biotic stresses. <i>Journal of Experimental Botany</i> , 2019, 70, 2143-2155.	2.4	26
18	Silencing barley cystatins HvCPI2 and HvCPI4 specifically modifies leaf responses to drought stress. <i>Plant, Cell and Environment</i> , 2018, 41, 1776-1790.	2.8	20

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19	Host plant use by two distinct lineages of the tomato red spider mite, <i>Tetranychus evansi</i> , differing in their distribution range. <i>Journal of Pest Science</i> , 2018, 91, 169-179.	1.9	14
20	Vacuolar processing enzyme 4 contributes to maternal control of grain size in barley by executing programmed cell death in the pericarp. <i>New Phytologist</i> , 2018, 218, 1127-1142.	3.5	30
21	Differential response of silencing <i>HvCly2</i> barley plants against <i>Magnaporthe oryzae</i> infection and light deprivation. <i>BMC Plant Biology</i> , 2018, 18, 337.	1.6	5
22	<i>Arabidopsis</i> response to the spider mite <i>Tetranychus urticae</i> depends on the regulation of reactive oxygen species homeostasis. <i>Scientific Reports</i> , 2018, 8, 9432.	1.6	33
23	<i>Arabidopsis</i> Kunitz Trypsin Inhibitors in Defense Against Spider Mites. <i>Frontiers in Plant Science</i> , 2018, 9, 986.	1.7	47
24	Dehydration Stress Contributes to the Enhancement of Plant Defense Response and Mite Performance on Barley. <i>Frontiers in Plant Science</i> , 2018, 9, 458.	1.7	23
25	Overexpression of <i>HvCly6</i> in Barley Enhances Resistance against <i>Tetranychus urticae</i> and Entails Partial Transcriptomic Reprogramming. <i>International Journal of Molecular Sciences</i> , 2018, 19, 697.	1.8	21
26	Plant Perception and Short-Term Responses to Phytophagous Insects and Mites. <i>International Journal of Molecular Sciences</i> , 2018, 19, 1356.	1.8	70
27	Senescence-Associated Genes in Response to Abiotic/Biotic Stresses. <i>Progress in Botany Fortschritte Der Botanik</i> , 2017, , 89-109.	0.1	5
28	MATI, a Novel Protein Involved in the Regulation of Herbivore-Associated Signaling Pathways. <i>Frontiers in Plant Science</i> , 2017, 8, 975.	1.7	42
29	<i>HvPap-1 C1A</i> Protease Participates Differentially in the Barley Response to a Pathogen and an Herbivore. <i>Frontiers in Plant Science</i> , 2017, 8, 1585.	1.7	18
30	Insights into the molecular evolution of peptidase inhibitors in arthropods. <i>PLoS ONE</i> , 2017, 12, e0187643.	1.1	1
31	Plant senescence and proteolysis: two processes with one destiny. <i>Genetics and Molecular Biology</i> , 2016, 39, 329-338.	0.6	124
32	Phytocystatins: Defense Proteins against Phytophagous Insects and Acari. <i>International Journal of Molecular Sciences</i> , 2016, 17, 1747.	1.8	65
33	<i>HvPap-1 C1A</i> protease actively participates in barley proteolysis mediated by abiotic stresses. <i>Journal of Experimental Botany</i> , 2016, 67, 4297-4310.	2.4	24
34	A Developmental Switch of Gene Expression in the Barley Seed Mediated by <i>HvVP1</i> (Viviparous-1) and <i>HvGAMYB</i> Interactions. <i>Plant Physiology</i> , 2016, 170, 2146-2158.	2.3	38
35	<i>HvPap-1 C1A</i> Protease and <i>HvCPI-2</i> Cystatin Contribute to Barley Grain Filling and Germination. <i>Plant Physiology</i> , 2016, 170, 2511-2524.	2.3	33
36	Synchronizing atomic force microscopy force mode and fluorescence microscopy in real time for immune cell stimulation and activation studies. <i>Ultramicroscopy</i> , 2016, 160, 168-181.	0.8	29

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37	Computational Tools for Genomic Studies in Plants. <i>Current Genomics</i> , 2016, 17, 509-514.	0.7	9
38	Tomato Whole Genome Transcriptional Response to <i>Tetranychus urticae</i> Identifies Divergence of Spider Mite-Induced Responses Between Tomato and <i>Arabidopsis</i> . <i>Molecular Plant-Microbe Interactions</i> , 2015, 28, 343-361.	1.4	90
39	Digestive proteases in bodies and faeces of the two-spotted spider mite, <i>Tetranychus urticae</i> . <i>Journal of Insect Physiology</i> , 2015, 78, 69-77.	0.9	71
40	Inhibitory Properties of Cysteine Protease Pro-Peptides from Barley Confer Resistance to Spider Mite Feeding. <i>PLoS ONE</i> , 2015, 10, e0128323.	1.1	32
41	Plant protein peptidase inhibitors: an evolutionary overview based on comparative genomics. <i>BMC Genomics</i> , 2014, 15, 812.	1.2	58
42	C1A cysteine protease-cystatin interactions in leaf senescence. <i>Journal of Experimental Botany</i> , 2014, 65, 3825-3833.	2.4	102
43	Responses to Phytophagous Arthropods. <i>Biotechnology in Agriculture and Forestry</i> , 2014, , 237-248.	0.2	1
44	Understanding plant defence responses against herbivore attacks: an essential first step towards the development of sustainable resistance against pests. <i>Transgenic Research</i> , 2013, 22, 697-708.	1.3	75
45	FROM PLANT GENOMES TO PROTEIN FAMILIES: COMPUTATIONAL TOOLS. <i>Computational and Structural Biotechnology Journal</i> , 2013, 8, e201307001.	1.9	7
46	Plant C1A Cysteine Peptidases in Germination and Senescence. , 2013, , 1852-1858.		10
47	Phylogenetically distant barley legumains have a role in both seed and vegetative tissues. <i>Journal of Experimental Botany</i> , 2013, 64, 2929-2941.	2.4	45
48	A cathepsin F-like peptidase involved in barley grain protein mobilization, HvPap-1, is modulated by its own propeptide and by cystatins. <i>Journal of Experimental Botany</i> , 2012, 63, 4615-4629.	2.4	32
49	Cysteine peptidases and their inhibitors in <i>Tetranychus urticae</i> : a comparative genomic approach. <i>BMC Genomics</i> , 2012, 13, 307.	1.2	38
50	Structural Basis for Specificity of Propeptide-Enzyme Interaction in Barley C1A Cysteine Peptidases. <i>PLoS ONE</i> , 2012, 7, e37234.	1.1	15
51	Co-evolution of Genes for Specification in Arthropod-Plant Interactions: A Bioinformatic Analysis in Plant and Arthropod Genomes. , 2012, , 1-14.		0
52	C1A cysteine-proteases and their inhibitors in plants. <i>Physiologia Plantarum</i> , 2012, 145, 85-94.	2.6	107
53	Gene Pyramiding of Peptidase Inhibitors Enhances Plant Resistance to the Spider Mite <i>Tetranychus urticae</i> . <i>PLoS ONE</i> , 2012, 7, e43011.	1.1	96
54	Plant protein-coding gene families: emerging bioinformatics approaches. <i>Trends in Plant Science</i> , 2011, 16, 558-567.	4.3	31

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55	The genome of <i>Tetranychus urticae</i> reveals herbivorous pest adaptations. <i>Nature</i> , 2011, 479, 487-492.	13.7	897
56	Differential in vitro and in vivo effect of barley cysteine and serine protease inhibitors on phytopathogenic microorganisms. <i>Plant Physiology and Biochemistry</i> , 2011, 49, 1191-1200.	2.8	23
57	A barley cysteine-proteinase inhibitor reduces the performance of two aphid species in artificial diets and transgenic <i>Arabidopsis</i> plants. <i>Transgenic Research</i> , 2011, 20, 305-319.	1.3	91
58	Expression of a barley cystatin gene in maize enhances resistance against phytophagous mites by altering their cysteine-proteases. <i>Plant Cell Reports</i> , 2011, 30, 101-112.	2.8	83
59	Clan CD of cysteine peptidases as an example of evolutionary divergences in related protein families across plant clades. <i>Gene</i> , 2010, 449, 59-69.	1.0	15
60	<i>Leishmania infantum</i> : Antiproliferative effect of recombinant plant cystatins on promastigotes and intracellular amastigotes estimated by direct counting and real-time PCR. <i>Experimental Parasitology</i> , 2009, 123, 341-346.	0.5	9
61	Comparative analysis of immunoglobulin polymerase chain reaction and flow cytometry in fine needle aspiration biopsy differential diagnosis of non-Hodgkin B-cell lymphoid malignancies. <i>Diagnostic Cytopathology</i> , 2009, 37, 647-653.	0.5	14
62	Characterization of the Entire Cystatin Gene Family in Barley and Their Target Cathepsin L-Like Cysteine-Proteases, Partners in the Hordein Mobilization during Seed Germination. <i>Plant Physiology</i> , 2009, 151, 1531-1545.	2.3	133
63	The origin and evolution of plant cystatins and their target cysteine proteinases indicate a complex functional relationship. <i>BMC Evolutionary Biology</i> , 2008, 8, 198.	3.2	129
64	Carboxy terminal extended phytocystatins are bifunctional inhibitors of papain and legumain cysteine proteinases. <i>FEBS Letters</i> , 2007, 581, 2914-2918.	1.3	96
65	Effects of potato plants expressing a barley cystatin on the predatory bug <i>Podisus maculiventris</i> via herbivorous prey feeding on the plant. <i>Transgenic Research</i> , 2007, 16, 1-13.	1.3	74
66	The family of DOF transcription factors: from green unicellular algae to vascular plants. <i>Molecular Genetics and Genomics</i> , 2007, 277, 379-390.	1.0	140
67	Ternary complex formation between HvMYBS3 and other factors involved in transcriptional control in barley seeds. <i>Plant Journal</i> , 2006, 47, 269-281.	2.8	74
68	HvMCB1, a R1MYB transcription factor from barley with antagonistic regulatory functions during seed development and germination. <i>Plant Journal</i> , 2006, 45, 17-30.	2.8	66
69	Structural and functional diversity within the cystatin gene family of <i>Hordeum vulgare</i> . <i>Journal of Experimental Botany</i> , 2006, 57, 4245-4255.	2.4	42
70	The DOF protein, SAD, interacts with GAMYB in plant nuclei and activates transcription of endosperm-specific genes during barley seed development. <i>Plant Journal</i> , 2005, 42, 652-662.	2.8	127
71	Comparative phylogenetic analysis of cystatin gene families from <i>Arabidopsis</i> , rice and barley. <i>Molecular Genetics and Genomics</i> , 2005, 273, 423-432.	1.0	90
72	The barley cystatin gene (<i>lcy</i>) is regulated by DOF transcription factors in aleurone cells upon germination. <i>Journal of Experimental Botany</i> , 2005, 56, 547-556.	2.4	38

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73	The strawberry gene <i>Cyf1</i> encodes a phytocystatin with antifungal properties. <i>Journal of Experimental Botany</i> , 2005, 56, 1821-1829.	2.4	82
74	Synaptic behaviour of hexaploid wheat haploids with different effectiveness of the diploidizing mechanism. <i>Cytogenetic and Genome Research</i> , 2005, 109, 210-214.	0.6	22
75	A single primer pair immunoglobulin polymerase chain reaction assay as a useful tool in fine-needle aspiration biopsy differential diagnosis of lymphoid malignancies. <i>Cancer</i> , 2003, 99, 180-185.	2.0	11
76	SAD: a new DOF protein from barley that activates transcription of a cathepsin B-like thiol protease gene in the aleurone of germinating seeds. <i>Plant Journal</i> , 2003, 33, 329-340.	2.8	80
77	A cathepsin B-like cysteine protease gene from <i>Hordeum vulgare</i> (gene <i>CatB</i>) induced by GA in aleurone cells is under circadian control in leaves. <i>Journal of Experimental Botany</i> , 2003, 54, 951-959.	2.4	50
78	Inhibition of Plant-Pathogenic Fungi by the Barley Cystatin <i>Hv-CPI</i> (Gene <i>Icy</i>) Is Not Associated with Its Cysteine-Proteinase Inhibitory Properties. <i>Molecular Plant-Microbe Interactions</i> , 2003, 16, 876-883.	1.4	68
79	GAMYB and BPBF transcriptional factors in the control of gene expression during development of barley endosperm.. , 2003, , 77-84.		0
80	The GAMYB protein from barley interacts with the DOF transcription factor BPBF and activates endosperm-specific genes during seed development. <i>Plant Journal</i> , 2002, 29, 453-464.	2.8	208
81	The Ph1 and Ph2 loci play different roles in the synaptic behaviour of hexaploid wheat <i>Triticum aestivum</i> . <i>Theoretical and Applied Genetics</i> , 2001, 103, 398-405.	1.8	53
82	The synaptic behaviour of <i>Triticum turgidum</i> with variable doses of the Ph1 locus. <i>Theoretical and Applied Genetics</i> , 2001, 102, 751-758.	1.8	39
83	The synaptic behaviour of the wild forms of <i>Triticum turgidum</i> and <i>T. timopheevii</i> . <i>Genome</i> , 2001, 44, 517-522.	0.9	5
84	The synaptic behaviour of the wild forms of <i>Triticum turgidum</i> and <i>T. timopheevii</i> . <i>Genome</i> , 2001, 44, 517-522.	0.9	3
85	Differences in the synaptic pattern in two autotetraploid cultivars of rye with different quadrivalent frequencies at metaphase I. <i>Genome</i> , 1999, 42, 662-667.	0.9	1
86	Synaptic behaviour of the tetraploid wheat <i>Triticum timopheevii</i> . <i>Theoretical and Applied Genetics</i> , 1996, 93, 1139-1144.	1.8	20
87	Synaptic abnormalities in spread nuclei of <i>Secale</i> . II. <i>Secale vavilovii</i> . <i>Genome</i> , 1995, 38, 772-779.	0.9	1
88	Further insights on chromosomal pairing of autopolyploids: a triploid and tetraploids of rye. <i>Chromosoma</i> , 1995, 104, 298-307.	1.0	22
89	Synaptic abnormalities in spread nuclei of <i>Secale</i> . I. Inbred lines. <i>Genome</i> , 1995, 38, 764-771.	0.9	4