

# Jaime F Martinez-Garcia

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7906765/publications.pdf>

Version: 2024-02-01

58  
papers

4,440  
citations

147801

31  
h-index

149698

56  
g-index

66  
all docs

66  
docs citations

66  
times ranked

4964  
citing authors

#	ARTICLE	IF	CITATIONS
1	Effect of light intensity on steviol glycosides production in leaves of <i>Stevia rebaudiana</i> plants. <i>Phytochemistry</i> , 2022, 194, 113027.	2.9	12
2	Development and carotenoid synthesis in dark-grown carrot taproots require <i>PHYTOCHROME RAPIDLY REGULATED1</i> . <i>Plant Physiology</i> , 2022, , .	4.8	5
3	Light signals generated by vegetation shade facilitate acclimation to low light in shade-avoider plants. <i>Plant Physiology</i> , 2021, 186, 2137-2151.	4.8	13
4	Adjustment of the PIF7-CHFR1 transcriptional module activity controls plant shade adaptation. <i>EMBO Journal</i> , 2021, 40, e104273.	7.8	32
5	The International Symposium on Plant Photobiology 2019: a bright and colourful experience. <i>Physiologia Plantarum</i> , 2020, 169, 297-300.	5.2	0
6	Shedding light on the chromatin changes that modulate shade responses. <i>Physiologia Plantarum</i> , 2020, 169, 407-417.	5.2	12
7	Shade Avoidance and Neighbor Detection. <i>Methods in Molecular Biology</i> , 2019, 2026, 157-168.	0.9	11
8	Photoreceptor Activity Contributes to Contrasting Responses to Shade in Cardamine and Arabidopsis Seedlings. <i>Plant Cell</i> , 2019, 31, tpc.00275.2019.	6.6	23
9	Chloroplasts Modulate Elongation Responses to Canopy Shade by Retrograde Pathways Involving HY5 and Abscisic Acid. <i>Plant Cell</i> , 2019, 31, 384-398.	6.6	40
10	Illuminating colors: regulation of carotenoid biosynthesis and accumulation by light. <i>Current Opinion in Plant Biology</i> , 2017, 37, 49-55.	7.1	142
11	A non-DNA-binding activity for the ATHB4 transcription factor in the control of vegetation proximity. <i>New Phytologist</i> , 2017, 216, 798-813.	7.3	14
12	Approaches to Study Light Effects on Brassinosteroid Sensitivity. <i>Methods in Molecular Biology</i> , 2017, 1564, 39-47.	0.9	3
13	Plant Responses to Vegetation Proximity: A Whole Life Avoiding Shade. <i>Frontiers in Plant Science</i> , 2016, 7, 236.	3.6	92
14	DRACULA2, a dynamic nucleoporin with a role in the regulation of the shade avoidance syndrome in Arabidopsis. <i>Development (Cambridge)</i> , 2016, 143, 1623-31.	2.5	25
15	bZIP and bHLH Family Members Integrate Transcriptional Responses to Light. , 2016, , 329-342.		3
16	Regulation of carotenoid biosynthesis by shade relies on specific subsets of antagonistic transcription factors and co-factors. <i>Plant Physiology</i> , 2015, 169, pp.00552.2015.	4.8	66
17	Meta-Analysis of Arabidopsis KANADI1 Direct Target Genes Identifies a Basic Growth-Promoting Module Acting Upstream of Hormonal Signaling Pathways. <i>Plant Physiology</i> , 2015, 169, 1240-1253.	4.8	26
18	The Shade Avoidance Syndrome in Arabidopsis: The Antagonistic Role of Phytochrome A and B Differentiates Vegetation Proximity and Canopy Shade. <i>PLoS ONE</i> , 2014, 9, e109275.	2.5	83

#	ARTICLE	IF	CITATIONS
19	<i>Cardamine hirsuta</i> : a versatile genetic system for comparative studies. <i>Plant Journal</i> , 2014, 78, 1-15.	5.7	78
20	Plant proximity perception dynamically modulates hormone levels and sensitivity in <i>Arabidopsis</i> . <i>Journal of Experimental Botany</i> , 2014, 65, 2937-2947.	4.8	79
21	The bHLH proteins BEE and BIM positively modulate the shade avoidance syndrome in <i>Arabidopsis</i> seedlings. <i>Plant Journal</i> , 2013, 75, 989-1002.	5.7	90
22	A Dual Mechanism Controls Nuclear Localization in the Atypical Basic-Helix-Loop-Helix Protein PAR1 of <i>Arabidopsis thaliana</i> . <i>Molecular Plant</i> , 2012, 5, 669-677.	8.3	17
23	ATHB4 and HAT3, two class II HD-ZIP transcription factors, control leaf development in <i>Arabidopsis</i> . <i>Plant Signaling and Behavior</i> , 2012, 7, 1382-1387.	2.4	80
24	A Light-Regulated Genetic Module Was Recruited to Carpel Development in <i>Arabidopsis</i> following a Structural Change to SPATULA. <i>Plant Cell</i> , 2012, 24, 2812-2825.	6.6	66
25	Genome-wide binding site analysis of REVOLUTA reveals a link between leaf patterning and light-mediated growth responses. <i>Plant Journal</i> , 2012, 72, 31-42.	5.7	120
26	Gibberellin A1 Metabolism Contributes to the Control of Photoperiod-Mediated Tuberization in Potato. <i>PLoS ONE</i> , 2011, 6, e24458.	2.5	44
27	The shade avoidance syndrome in <i>Arabidopsis</i> : a fundamental role for atypical basic helix-loop-helix proteins as transcriptional cofactors. <i>Plant Journal</i> , 2011, 66, 258-267.	5.7	92
28	A DELLA in Disguise: SPATULA Restrains the Growth of the Developing <i>Arabidopsis</i> Seedling. <i>Plant Cell</i> , 2011, 23, 1337-1351.	6.6	77
29	A novel high-throughput in vivo molecular screen for shade avoidance mutants identifies a novel phyA mutation. <i>Journal of Experimental Botany</i> , 2011, 62, 2973-2987.	4.8	20
30	Regulatory Components of Shade Avoidance Syndrome. <i>Advances in Botanical Research</i> , 2010, 53, 65-116.	1.1	61
31	Light Signalling in Plant Developmental Regulation. , 2010, , 255-274.		1
32	Genome-Wide Classification and Evolutionary Analysis of the bHLH Family of Transcription Factors in <i>Arabidopsis</i> , Poplar, Rice, Moss, and Algae. <i>Plant Physiology</i> , 2010, 153, 1398-1412.	4.8	493
33	ATHB4, a regulator of shade avoidance, modulates hormone response in <i>Arabidopsis</i> seedlings. <i>Plant Journal</i> , 2009, 59, 266-277.	5.7	111
34	Integration of Light and Auxin Signaling. <i>Cold Spring Harbor Perspectives in Biology</i> , 2009, 1, a001586-a001586.	5.5	149
35	<i>PROCERA</i> encodes a DELLA protein that mediates control of dissected leaf form in tomato. <i>Plant Journal</i> , 2008, 56, 603-612.	5.7	110
36	Light signaling: back to space. <i>Trends in Plant Science</i> , 2008, 13, 108-114.	8.8	41

#	ARTICLE	IF	CITATIONS
37	PAR1 and PAR2 integrate shade and hormone transcriptional networks. <i>Plant Signaling and Behavior</i> , 2008, 3, 453-454.	2.4	29
38	Interaction of shade avoidance and auxin responses: a role for two novel atypical bHLH proteins. <i>EMBO Journal</i> , 2007, 26, 4756-4767.	7.8	195
39	Identification of Primary Target Genes of Phytochrome Signaling. Early Transcriptional Control during Shade Avoidance Responses in Arabidopsis. <i>Plant Physiology</i> , 2006, 141, 85-96.	4.8	127
40	Plastid Cues Posttranscriptionally Regulate the Accumulation of Key Enzymes of the Methylerythritol Phosphate Pathway in Arabidopsis. <i>Plant Physiology</i> , 2006, 141, 75-84.	4.8	84
41	Distinct Light-Mediated Pathways Regulate the Biosynthesis and Exchange of Isoprenoid Precursors during Arabidopsis Seedling Development. <i>Plant Cell</i> , 2004, 16, 144-156.	6.6	189
42	PHOR1: A U-Box GA Signaling Component With a Role in Proteasome Degradation?. <i>Journal of Plant Growth Regulation</i> , 2003, 22, 152-162.	5.1	19
43	Potato Tuberization: Evidence for a SD-Dependent and a Gibberellin-Dependent Pathway of Induction. , 2003, , 57-66.		2
44	Control of photoperiod-regulated tuberization in potato by the Arabidopsis flowering-time gene CONSTANS. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 15211-15216.	7.1	183
45	The Interaction of Gibberellins and Photoperiod in the Control of Potato Tuberization. <i>Journal of Plant Growth Regulation</i> , 2001, 20, 377-386.	5.1	68
46	The end-of-day far-red irradiation increases gibberellin A1 content in cowpea ( <i>Vigna sinensis</i> ) epicotyls by reducing its inactivation. <i>Physiologia Plantarum</i> , 2000, 108, 426-434.	5.2	25
47	Direct Targeting of Light Signals to a Promoter Element-Bound Transcription Factor. <i>Science</i> , 2000, 288, 859-863.	12.6	629
48	The end-of-day far-red irradiation increases gibberellin A1 content in cowpea ( <i>Vigna sinensis</i> ) epicotyls by reducing its inactivation. <i>Physiologia Plantarum</i> , 2000, 108, 426-434.	5.2	11
49	The HMG-I/Y protein PF1 stimulates binding of the transcriptional activator GT-2 to the PHYA gene promoter. <i>Plant Journal</i> , 1999, 18, 173-183.	5.7	36
50	A simple, rapid and quantitative method for preparing Arabidopsis protein extracts for immunoblot analysis. <i>Plant Journal</i> , 1999, 20, 251-257.	5.7	172
51	Two bZIP proteins from <i>Antirrhinum</i> flowers preferentially bind a hybrid C <sub>2</sub> H <sub>2</sub> /C <sub>2</sub> H <sub>2</sub> motif and help to define a new subfamily of bZIP transcription factors. <i>Plant Journal</i> , 1998, 13, 489-505.	5.7	67
52	Apparent redundancy in myb gene function provides gearing for the control of flavonoid biosynthesis in <i>antirrhinum</i> flowers.. <i>Plant Cell</i> , 1996, 8, 1519-1532.	6.6	175
53	Apparent Redundancy in myb Gene Function Provides Gearing for the Control of Flavonoid Biosynthesis in <i>Antirrhinum</i> Flowers. <i>Plant Cell</i> , 1996, 8, 1519.	6.6	24
54	An acylcyclohexadione retardant inhibits gibberellin A1 metabolism, thereby nullifying phytochrome-modulation of cowpea epicotyl explants. <i>Physiologia Plantarum</i> , 1995, 94, 708-714.	5.2	9

#	ARTICLE	IF	CITATIONS
55	An acylcyclohexadione retardant inhibits gibberellin A1 metabolism, thereby nullifying phytochrome-modulation of cowpea epicotyl explants. <i>Physiologia Plantarum</i> , 1995, 94, 708-714.	5.2	12
56	Effect of the growth retardant LAB 198 999, an acylcyclohexanedione compound, on epicotyl elongation and metabolism of gibberellins A1 and A20 in cowpea. <i>Planta</i> , 1992, 188, 245-251.	3.2	13
57	Interaction of gibberellins and phytochrome in the control of cowpea epicotyl elongation. <i>Physiologia Plantarum</i> , 1992, 86, 236-244.	5.2	26
58	Chromatin structure of the 5' flanking region of the yeast <i>LEU2</i> gene. <i>Molecular Genetics and Genomics</i> , 1989, 217, 464-470.	2.4	9