

Albert Ferrer

List of Publications by Year in descending order

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65
papers

3,626
citations

134610

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150775

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66
docs citations

66
times ranked

3701
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#	ARTICLE	IF	CITATIONS
1	<i>Pseudomonas germanica</i> sp. nov., isolated from <i>Iris germanica</i> rhizomes. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2022, 72, .	0.8	4
2	Structural and functional analysis of tomato sterol C22 desaturase. <i>BMC Plant Biology</i> , 2021, 21, 141.	1.6	3
3	Phytosterol metabolism in plant positive-strand RNA virus replication. <i>Plant Cell Reports</i> , 2021, , 1.	2.8	3
4	Distinct metabolic pathways drive monoterpenoid biosynthesis in a natural population of <i>Pelargonium graveolens</i> . <i>Journal of Experimental Botany</i> , 2020, 71, 258-271.	2.4	18
5	Inactivation of UDP-Glucose Sterol Glucosyltransferases Enhances <i>Arabidopsis</i> Resistance to <i>Botrytis cinerea</i> . <i>Frontiers in Plant Science</i> , 2019, 10, 1162.	1.7	17
6	Nerolidol production in agroinfiltrated tobacco: Impact of protein stability and membrane targeting of strawberry (<i>Fragaria ananassa</i>) <i>NEROLIDOL SYNTHASE1</i> . <i>Plant Science</i> , 2018, 267, 112-123.	1.7	4
7	Identification and Characterization of Sterol Acyltransferases Responsible for Steryl Ester Biosynthesis in Tomato. <i>Frontiers in Plant Science</i> , 2018, 9, 588.	1.7	15
8	Complex interplays between phytosterols and plastid development. <i>Plant Signaling and Behavior</i> , 2017, 12, e1387708.	1.2	4
9	Emerging roles for conjugated sterols in plants. <i>Progress in Lipid Research</i> , 2017, 67, 27-37.	5.3	161
10	Tomato UDP-Glucose Sterol Glycosyltransferases: A Family of Developmental and Stress Regulated Genes that Encode Cytosolic and Membrane-Associated Forms of the Enzyme. <i>Frontiers in Plant Science</i> , 2017, 8, 984.	1.7	37
11	Suppressing Farnesyl Diphosphate Synthase Alters Chloroplast Development and Triggers Sterol-Dependent Induction of Jasmonate- and Fe-Related Responses. <i>Plant Physiology</i> , 2016, 172, 93-117.	2.3	32
12	Strategies and Methodologies for the Co-expression of Multiple Proteins in Plants. <i>Advances in Experimental Medicine and Biology</i> , 2016, 896, 263-285.	0.8	5
13	Elucidation of the biosynthesis of carnosic acid and its reconstitution in yeast. <i>Nature Communications</i> , 2016, 7, 12942.	5.8	122
14	Towards Elucidating Carnosic Acid Biosynthesis in Lamiaceae: Functional Characterization of the Three First Steps of the Pathway in <i>Salvia fruticosa</i> and <i>Rosmarinus officinalis</i> . <i>PLoS ONE</i> , 2015, 10, e0124106.	1.1	67
15	<i>Arabidopsis</i> Squalene Epoxidase 3 (SQE3) Complements SQE1 and Is Important for Embryo Development and Bulk Squalene Epoxidase Activity. <i>Molecular Plant</i> , 2015, 8, 1090-1102.	3.9	59
16	Elucidation and in planta reconstitution of the parthenolide biosynthetic pathway. <i>Metabolic Engineering</i> , 2014, 23, 145-153.	3.6	68
17	Characterization of two genes for the biosynthesis of abietane-type diterpenes in rosemary (<i>Rosmarinus officinalis</i>) glandular trichomes. <i>Phytochemistry</i> , 2014, 101, 52-64.	1.4	106
18	Determination of 3-Hydroxy-3-methylglutaryl CoA Reductase Activity in Plants. <i>Methods in Molecular Biology</i> , 2014, 1153, 21-40.	0.4	7

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19	Farnesyl Diphosphate Synthase Assay. <i>Methods in Molecular Biology</i> , 2014, 1153, 41-53.	0.4	5
20	Three-step pathway engineering results in more incidence rate and higher emission of nerolidol and improved attraction of <i>Diadegma semiclausum</i> . <i>Metabolic Engineering</i> , 2013, 15, 88-97.	3.6	35
21	The <i>SUD1</i> Gene Encodes a Putative E3 Ubiquitin Ligase and Is a Positive Regulator of 3-Hydroxy-3-Methylglutaryl Coenzyme A Reductase Activity in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2013, 25, 728-743.	3.1	78
22	Biosynthesis of Isoprenoid Precursors in <i>Arabidopsis</i> . , 2012, , 439-456.		5
23	Characterization of <i>Arabidopsis</i> FPS Isozymes and FPS Gene Expression Analysis Provide Insight into the Biosynthesis of Isoprenoid Precursors in Seeds. <i>PLoS ONE</i> , 2012, 7, e49109.	1.1	30
24	Modulation of plant HMG-CoA reductase by protein phosphatase 2A. <i>Plant Signaling and Behavior</i> , 2011, 6, 1127-1131.	1.2	41
25	Multilevel Control of <i>Arabidopsis</i> 3-Hydroxy-3-Methylglutaryl Coenzyme A Reductase by Protein Phosphatase 2A. <i>Plant Cell</i> , 2011, 23, 1494-1511.	3.1	99
26	The <i>Arabidopsis thaliana</i> FPP synthase isozymes have overlapping and specific functions in isoprenoid biosynthesis, and complete loss of FPP synthase activity causes early developmental arrest. <i>Plant Journal</i> , 2010, 63, 512-525.	2.8	80
27	PLEIOTROPIC REGULATORY LOCUS 1 (PRL1) Integrates the Regulation of Sugar Responses with Isoprenoid Metabolism in <i>Arabidopsis</i> . <i>Molecular Plant</i> , 2010, 3, 101-112.	3.9	64
28	Identification of the <i>Arabidopsis dry2/sqe1</i> mutant reveals a central role for sterols in drought tolerance and regulation of reactive oxygen species. <i>Plant Journal</i> , 2009, 59, 63-76.	2.8	114
29	<i>Arabidopsis</i> 3-hydroxy-3-methylglutaryl-CoA reductase is regulated at the post-translational level in response to alterations of the sphingolipid and the sterol biosynthetic pathways. <i>Phytochemistry</i> , 2009, 70, 53-59.	1.4	58
30	<i>Arabidopsis thaliana</i> contains a single gene encoding squalene synthase. <i>Plant Molecular Biology</i> , 2008, 67, 25-36.	2.0	63
31	Mitochondrial targeting of farnesyl diphosphate synthase is a widespread phenomenon in eukaryotes. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2007, 1773, 419-426.	1.9	30
32	<i>Arabidopsis thaliana</i> expresses two functional isoforms of Arvp, a protein involved in the regulation of cellular lipid homeostasis. <i>Biochimica Et Biophysica Acta - Molecular and Cell Biology of Lipids</i> , 2006, 1761, 725-735.	1.2	29
33	Overexpression of Farnesyl Diphosphate Synthase in <i>Arabidopsis</i> Mitochondria Triggers Light-dependent Lesion Formation and Alters Cytokinin Homeostasis. <i>Plant Molecular Biology</i> , 2006, 61, 195-213.	2.0	30
34	Subcellular Localization of <i>Arabidopsis</i> 3-Hydroxy-3-Methylglutaryl-Coenzyme A Reductase. <i>Plant Physiology</i> , 2005, 137, 57-69.	2.3	102
35	Distinct Light-Mediated Pathways Regulate the Biosynthesis and Exchange of Isoprenoid Precursors during <i>Arabidopsis</i> Seedling Development. <i>Plant Cell</i> , 2004, 16, 144-156.	3.1	189
36	The metabolic imbalance underlying lesion formation in <i>Arabidopsis thaliana</i> overexpressing farnesyl diphosphate synthase (isoform $\frac{1}{2}$ IS) leads to oxidative stress and is triggered by the developmental decline of endogenous HMGR activity. <i>Planta</i> , 2004, 219, 982-992.	1.6	65

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37	High Level Expression of Chorismate Pyruvate-Lyase (UbiC) and HMG-CoA Reductase in Hairy Root Cultures of <i>Lithospermum erythrorhizon</i> . <i>Plant and Cell Physiology</i> , 2002, 43, 894-902.	1.5	24
38	Expression and Molecular Analysis of the <i>Arabidopsis</i> DXR Gene Encoding 1-Deoxy-d-Xylulose 5-Phosphate Reductoisomerase, the First Committed Enzyme of the 2-C-Methyl-d-Erythritol 4-Phosphate Pathway. <i>Plant Physiology</i> , 2002, 129, 1581-1591.	2.3	203
39	Overexpression of <i>Arabidopsis thaliana</i> farnesyl diphosphate synthase (FPS1S) in transgenic <i>Arabidopsis</i> induces a cell death/senescence-like response and reduced cytokinin levels. <i>Plant Journal</i> , 2002, 30, 123-132.	2.8	102
40	Contribution of engineered electrostatic interactions to the stability of cytosolic malate dehydrogenase. <i>Protein Engineering, Design and Selection</i> , 2001, 14, 911-917.	1.0	11
41	Molecular cloning and expression analysis of the mevalonate kinase gene from <i>Arabidopsis thaliana</i> . <i>Plant Molecular Biology</i> , 2000, 42, 365-376.	2.0	48
42	The <i>Arabidopsis thaliana</i> PPX/PP4 phosphatases: molecular cloning and structural organization of the genes and immunolocalization of the proteins to plastids. <i>Plant Molecular Biology</i> , 2000, 44, 499-511.	2.0	15
43	Spatial and temporal patterns of GUS expression directed by 5' regions of the <i>Arabidopsis thaliana</i> farnesyl diphosphate synthase genes FPS1 and FPS2. <i>Plant Molecular Biology</i> , 2000, 44, 747-758.	2.0	36
44	Characterization of dehydrodolichyl diphosphate synthase of <i>Arabidopsis thaliana</i> , a key enzyme in dolichol biosynthesis. <i>FEBS Letters</i> , 2000, 477, 170-174.	1.3	82
45	Mevalonate Biosynthesis in Plants. <i>Critical Reviews in Biochemistry and Molecular Biology</i> , 1999, 34, 107-122.	2.3	56
46	Molecular cloning and characterization of two phosphatase 2A catalytic subunit genes from <i>Arabidopsis thaliana</i> . <i>Gene</i> , 1998, 209, 105-112.	1.0	21
47	Protein Phosphatase 2A and Protein Phosphatase X Genes in <i>Arabidopsis thaliana</i> . , 1998, 93, 201-212.		2
48	The <i>Arabidopsis thaliana</i> FPS1 Gene Generates a Novel mRNA That Encodes a Mitochondrial Farnesyl-diphosphate Synthase Isoform. <i>Journal of Biological Chemistry</i> , 1997, 272, 15381-15388.	1.6	152
49	Three spinach leaf nitrate reductase-3-hydroxy-3-methylglutaryl-CoA reductase kinases that are regulated by reversible phosphorylation and/or Ca ²⁺ ions. <i>Biochemical Journal</i> , 1997, 325, 101-109.	1.7	113
50	Cloning and Characterization of the <i>Arabidopsis thaliana</i> SQS1 Gene Encoding Squalene Synthase. Involvement of the C-Terminal Region of the Enzyme in the Channeling of Squalene through the Sterol Pathway. <i>FEBS Journal</i> , 1997, 249, 61-69.	0.2	79
51	<i>Arabidopsis thaliana</i> Contains Two Differentially Expressed Farnesyl-Diphosphate Synthase Genes. <i>Journal of Biological Chemistry</i> , 1996, 271, 7774-7780.	1.6	158
52	Bacterial Expression of the Catalytic Domain of 3-Hydroxy-3-Methylglutaryl-CoA Reductase (Isoform) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5 oleracea 3-Hydroxy-3-Methylglutaryl-CoA Reductase Kinase. <i>FEBS Journal</i> , 1995, 233, 506-513.	0.2	120
53	Molecular characterization of a fourth isoform of the catalytic subunit of protein phosphatase 2A from <i>Arabidopsis thaliana</i> . <i>Plant Molecular Biology</i> , 1994, 26, 523-528.	2.0	39
54	Two Radish Genes for 3-Hydroxy-3-Methylglutaryl-CoA Reductase Isozymes Complement Mevalonate Auxotrophy in a Yeast Mutant and Yield Membrane-Bound Active Enzyme. <i>Journal of Plant Physiology</i> , 1994, 143, 479-487.	1.6	38

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55	Protein phosphatases in higher plants: multiplicity of type 2A phosphatases in <i>Arabidopsis thaliana</i> . <i>Plant Molecular Biology</i> , 1993, 21, 475-485.	2.0	75
56	Identification and molecular cloning of two homologues of protein phosphatase X from <i>Arabidopsis thaliana</i> . <i>Plant Molecular Biology</i> , 1993, 23, 1177-1185.	2.0	24
57	Properties and Molecular Cloning of Plant HMG-CoA Reductase. , 1992, , 29-49.		2
58	Aspects related to mevalonate biosynthesis in plants. <i>Lipids</i> , 1991, 26, 637-648.	0.7	53
59	Expression of catalytically active radish 3-hydroxy-3-methylglutaryl coenzyme A reductase in <i>Escherichia coli</i> . <i>FEBS Letters</i> , 1990, 266, 67-71.	1.3	33
60	Isolation and structural characterization of a cDNA encoding <i>Arabidopsis thaliana</i> 3-hydroxy-3-methylglutaryl coenzyme A reductase. <i>Plant Molecular Biology</i> , 1989, 13, 627-638.	2.0	136
61	Allosteric Activation of Rat Liver Microsomal [Hydroxymethylglutaryl-CoA Reductase (NADPH)] Kinase by Nucleoside Phosphates. <i>Biological Chemistry Hoppe-Seyler</i> , 1987, 368, 249-258.	1.4	6
62	Reaction of 5- ³² P-fluorosulfonylbenzoyl adenosine with the catalytic and AMP allosteric sites of microsomal HMG-CoA reductase kinase. <i>Biochemical and Biophysical Research Communications</i> , 1987, 148, 1009-1016.	1.0	3
63	Activation of rat liver cytosolic 3-hydroxy-3-methylglutaryl Coenzyme A reductase kinase by adenosine 5- ³² P-monophosphate. <i>Biochemical and Biophysical Research Communications</i> , 1985, 132, 497-504.	1.0	116
64	Phosphorylation of 3-hydroxy-3-methylglutaryl coenzyme A reductase by microsomal 3-hydroxy-3-methylglutaryl coenzyme A reductase kinase. <i>Archives of Biochemistry and Biophysics</i> , 1984, 230, 227-237.	1.4	22
65	Inactivation and Reactivation of Rat Liver 3-Hydroxy-3-methylglutaryl-CoA-Reductase Phosphatases: Effect of Phosphate, Pyrophosphate and Divalent Cations. <i>Hoppe-Seyler's Zeitschrift für Physiologische Chemie</i> , 1982, 363, 1217-1224.	1.7	7