

Catherine Deborde

List of Publications by Year in descending order

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78
papers

3,532
citations

136740

32
h-index

143772

57
g-index

80
all docs

80
docs citations

80
times ranked

5201
citing authors

#	ARTICLE	IF	CITATIONS
1	Gene and Metabolite Regulatory Network Analysis of Early Developing Fruit Tissues Highlights New Candidate Genes for the Control of Tomato Fruit Composition and Development. <i>Plant Physiology</i> , 2009, 149, 1505-1528.	2.3	199
2	Microclimate Influence on Mineral and Metabolic Profiles of Grape Berries. <i>Journal of Agricultural and Food Chemistry</i> , 2006, 54, 6765-6775.	2.4	188
3	¹ H NMR metabolite fingerprints of grape berry: Comparison of vintage and soil effects in Bordeaux grapevine growing areas. <i>Analytica Chimica Acta</i> , 2006, 563, 346-352.	2.6	159
4	¹ H NMR, GC-MS, and Data Set Correlation for Fruit Metabolomics: Application to Spatial Metabolite Analysis in Melon. <i>Analytical Chemistry</i> , 2009, 81, 2884-2894.	3.2	147
5	COordination of Standards in MetabOlogicS (COSMOS): facilitating integrated metabolomics data access. <i>Metabolomics</i> , 2015, 11, 1587-1597.	1.4	140
6	¹ H NMR and Chemometrics To Characterize Mature Grape Berries in Four Wine-Growing Areas in Bordeaux, France. <i>Journal of Agricultural and Food Chemistry</i> , 2005, 53, 6382-6389.	2.4	137
7	NMRProcFlow: a graphical and interactive tool dedicated to 1D spectra processing for NMR-based metabolomics. <i>Metabolomics</i> , 2017, 13, 36.	1.4	128
8	Quantitative metabolic profiles of tomato flesh and seeds during fruit development: complementary analysis with ANN and PCA. <i>Metabolomics</i> , 2007, 3, 273-288.	1.4	119
9	Extensive metabolic cross-talk in melon fruit revealed by spatial and developmental combinatorial metabolomics. <i>New Phytologist</i> , 2011, 190, 683-696.	3.5	111
10	Effects of long-term cadmium exposure on growth and metabolomic profile of tomato plants. <i>Ecotoxicology and Environmental Safety</i> , 2010, 73, 1965-1974.	2.9	96
11	Hyperpolarized NMR of plant and cancer cell extracts at natural abundance. <i>Analyst</i> , 2015, 140, 5860-5863.	1.7	87
12	An inter-laboratory comparison demonstrates that [¹ H]-NMR metabolite fingerprinting is a robust technique for collaborative plant metabolomic data collection. <i>Metabolomics</i> , 2010, 6, 263-273.	1.4	86
13	Plant metabolism as studied by NMR spectroscopy. <i>Progress in Nuclear Magnetic Resonance Spectroscopy</i> , 2017, 102-103, 61-97.	3.9	85
14	Metabolomics and fish nutrition: a review in the context of sustainable feed development. <i>Reviews in Aquaculture</i> , 2020, 12, 261-282.	4.6	84
15	Down-regulation of a single auxin efflux transport protein in tomato induces precocious fruit development. <i>Journal of Experimental Botany</i> , 2012, 63, 4901-4917.	2.4	82
16	Plant Metabolomics and Its Potential for Systems Biology Research. <i>Methods in Enzymology</i> , 2011, 500, 299-336.	0.4	78
17	Sucrose, Glucose, and Fructose Extraction in Aqueous Carrot Root Extracts Prepared at Different Temperatures by Means of Direct NMR Measurements. <i>Journal of Agricultural and Food Chemistry</i> , 2006, 54, 4681-4686.	2.4	75
18	Metabolic acclimation to hypoxia revealed by metabolite gradients in melon fruit. <i>Journal of Plant Physiology</i> , 2010, 167, 242-245.	1.6	75

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19	Metabolomic and elemental profiling of melon fruit quality as affected by genotype and environment. <i>Metabolomics</i> , 2013, 9, 57-77.	1.4	74
20	Metabolomics in melon: A new opportunity for aroma analysis. <i>Phytochemistry</i> , 2014, 99, 61-72.	1.4	66
21	Metabolomic profiling in tomato reveals diel compositional changes in fruit affected by source-sink relationships. <i>Journal of Experimental Botany</i> , 2015, 66, 3391-3404.	2.4	62
22	Metabolic response in roots of <i>Prunus</i> rootstocks submitted to iron chlorosis. <i>Journal of Plant Physiology</i> , 2011, 168, 415-423.	1.6	58
23	Highly Repeatable Dissolution Dynamic Nuclear Polarization for Heteronuclear NMR Metabolomics. <i>Analytical Chemistry</i> , 2016, 88, 6179-6183.	3.2	57
24	MeRy-B: a web knowledgebase for the storage, visualization, analysis and annotation of plant NMR metabolomic profiles. <i>BMC Plant Biology</i> , 2011, 11, 104.	1.6	54
25	Proton NMR quantitative profiling for quality assessment of greenhouse-grown tomato fruit. <i>Metabolomics</i> , 2009, 5, 183-198.	1.4	51
26	Absolute quantification of metabolites in tomato fruit extracts by fast 2D NMR. <i>Metabolomics</i> , 2015, 11, 1231-1242.	1.4	50
27	nmrML: A Community Supported Open Data Standard for the Description, Storage, and Exchange of NMR Data. <i>Analytical Chemistry</i> , 2018, 90, 649-656.	3.2	50
28	Enhanced polyamine accumulation alters carotenoid metabolism at the transcriptional level in tomato fruit over-expressing spermidine synthase. <i>Journal of Plant Physiology</i> , 2011, 168, 242-252.	1.6	48
29	(Homo)glutathione Deficiency Impairs Root-knot Nematode Development in <i>Medicago truncatula</i> . <i>PLoS Pathogens</i> , 2012, 8, e1002471.	2.1	48
30	Hyperpolarized NMR Metabolomics at Natural ¹³ C Abundance. <i>Analytical Chemistry</i> , 2020, 92, 14867-14871.	3.2	44
31	A genomics and multi-platform metabolomics approach to identify new traits of rice quality in traditional and improved varieties. <i>Metabolomics</i> , 2012, 8, 771-783.	1.4	43
32	Optimizing 1D ¹ H-NMR profiling of plant samples for high throughput analysis: extract preparation, standardization, automation and spectra processing. <i>Metabolomics</i> , 2019, 15, 28.	1.4	37
33	Biomass composition explains fruit relative growth rate and discriminates climacteric from non-climacteric species. <i>Journal of Experimental Botany</i> , 2020, 71, 5823-5836.	2.4	35
34	Comparative Metabolomics and Molecular Phylogenetics of Melon (<i>Cucumis melo</i> , Cucurbitaceae) Biodiversity. <i>Metabolites</i> , 2020, 10, 121.	1.3	35
35	Maize metabolome and proteome responses to controlled cold stress partly mimic early sowing effects in the field and differ from those of <i>Arabidopsis</i> . <i>Plant, Cell and Environment</i> , 2021, 44, 1504-1521.	2.8	32
36	Stress and osmoprotection in propionibacteria. <i>Dairy Science and Technology</i> , 1999, 79, 59-69.	0.9	32

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37	An integrative genomics approach for deciphering the complex interactions between ascorbate metabolism and fruit growth and composition in tomato. <i>Comptes Rendus - Biologies</i> , 2009, 332, 1007-1021.	0.1	30
38	¹ H-NMR metabolomic profiling reveals a distinct metabolic recovery response in shoots and roots of temporarily drought-stressed sugar beets. <i>PLoS ONE</i> , 2018, 13, e0196102.	1.1	27
39	Integrative Metabolomics for Assessing the Effect of Insect (<i>Hermetia illucens</i>) Protein Extract on Rainbow Trout Metabolism. <i>Metabolites</i> , 2020, 10, 83.	1.3	27
40	Interactions between Pyruvate and Lactate Metabolism in <i>Propionibacterium freudenreichii</i> subsp. <i>shermanii</i> : In Vivo ¹³ C Nuclear Magnetic Resonance Studies. <i>Applied and Environmental Microbiology</i> , 2000, 66, 2012-2020.	1.4	26
41	Transcriptional and Metabolic Adjustments in ADP-Glucose Pyrophosphorylase-Deficient <i>Zea mays</i> Maize Kernels. <i>Plant Physiology</i> , 2008, 146, 1553-1570.	2.3	25
42	Mycotoxin Biosynthesis and Central Metabolism Are Two Interlinked Pathways in <i>Fusarium graminearum</i> , as Demonstrated by the Extensive Metabolic Changes Induced by Caffeic Acid Exposure. <i>Applied and Environmental Microbiology</i> , 2018, 84, .	1.4	25
43	An efficient spectra processing method for metabolite identification from ¹ H-NMR metabolomics data. <i>Analytical and Bioanalytical Chemistry</i> , 2013, 405, 5049-5061.	1.9	24
44	The peach HECATE3-like gene FLESHY plays a double role during fruit development. <i>Plant Molecular Biology</i> , 2016, 91, 97-114.	2.0	24
45	Honeydew feeding increased the longevity of two egg parasitoids of the pine processionary moth. <i>Journal of Applied Entomology</i> , 2011, 135, 184-194.	0.8	23
46	MeRy-B, a Metabolomic Database and Knowledge Base for Exploring Plant Primary Metabolism. <i>Methods in Molecular Biology</i> , 2014, 1083, 3-16.	0.4	22
47	Deciphering genetic diversity and inheritance of tomato fruit weight and composition through a systems biology approach. <i>Journal of Experimental Botany</i> , 2013, 64, 5737-5752.	2.4	20
48	Omics Data Reveal Putative Regulators of Einkorn Grain Protein Composition under Sulfur Deficiency. <i>Plant Physiology</i> , 2020, 183, 501-516.	2.3	20
49	Characterizing alternative feeds for rainbow trout (<i>O. mykiss</i>) by ¹ H NMR metabolomics. <i>Metabolomics</i> , 2018, 14, 155.	1.4	18
50	Intestinal microbiota in rainbow trout, <i>Oncorhynchus mykiss</i> , fed diets with different levels of fish-based and plant ingredients: A correlative approach with some plasma metabolites. <i>Aquaculture Nutrition</i> , 2018, 24, 1563-1576.	1.1	18
51	NMR-Based Tissue and Developmental Metabolomics of Tomato Fruit. <i>Metabolites</i> , 2019, 9, 93.	1.3	18
52	Metabolomic characterization of sunflower leaf allows discriminating genotype groups or stress levels with a minimal set of metabolic markers. <i>Metabolomics</i> , 2019, 15, 56.	1.4	17
53	The Tomato Guanylate-Binding Protein SIGBP1 Enables Fruit Tissue Differentiation by Maintaining Endopolyloid Cells in a Non-Proliferative State. <i>Plant Cell</i> , 2020, 32, 3188-3205.	3.1	17
54	Precautions for Harvest, Sampling, Storage, and Transport of Crop Plant Metabolomics Samples. <i>Methods in Molecular Biology</i> , 2011, 860, 51-63.	0.4	17

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55	Maturation of nematode-induced galls in <i>Medicago truncatula</i> is related to water status and primary metabolism modifications. <i>Plant Science</i> , 2015, 232, 77-85.	1.7	15
56	Metabotyping of 30 maize hybrids under early-sowing conditions reveals potential marker-metabolites for breeding. <i>Metabolomics</i> , 2018, 14, 132.	1.4	15
57	Putative imbalanced amino acid metabolism in rainbow trout long term fed a plant-based diet as revealed by ¹ H-NMR metabolomics. <i>Journal of Nutritional Science</i> , 2021, 10, e13.	0.7	15
58	High-Resolution ¹ H-NMR Spectroscopy and Beyond to Explore Plant Metabolome. <i>Advances in Botanical Research</i> , 2013, , 1-66.	0.5	14
59	A Systems Biology Study in Tomato Fruit Reveals Correlations between the Ascorbate Pool and Genes Involved in Ribosome Biogenesis, Translation, and the Heat-Shock Response. <i>Frontiers in Plant Science</i> , 2018, 9, 137.	1.7	11
60	Proton-NMR Metabolomics of Rainbow Trout Fed a Plant-Based Diet Supplemented with Graded Levels of a Protein-Rich Yeast Fraction Reveal Several Metabolic Processes Involved in Growth. <i>Journal of Nutrition</i> , 2020, 150, 2268-2277.	1.3	11
61	In Vivo ¹³ C NMR Study of the Bidirectional Reactions of the Wood-Werkman Cycle and around the Pyruvate Node in <i>Propionibacterium freudenreichii</i> subsp. <i>shermanii</i> and <i>Propionibacterium acidipropionici</i> . <i>Metabolic Engineering</i> , 1999, 1, 309-319.	3.6	9
62	Characterization of GMO or glyphosate effects on the composition of maize grain and maize-based diet for rat feeding. <i>Metabolomics</i> , 2018, 14, 36.	1.4	9
63	Central Metabolism Is Tuned to the Availability of Oxygen in Developing Melon Fruit. <i>Frontiers in Plant Science</i> , 2019, 10, 594.	1.7	9
64	Metabolite Fruit Profile Is Altered in Response to Source-Sink Imbalance and Can Be Used as an Early Predictor of Fruit Quality in Nectarine. <i>Frontiers in Plant Science</i> , 2020, 11, 604133.	1.7	9
65	In silico exploration of the fructose-6-phosphate phosphorylation step in glycolysis: genomic evidence of the coexistence of an atypical ATP-dependent along with a PPI-dependent phosphofructokinase in <i>Propionibacterium freudenreichii</i> subsp. <i>shermanii</i> . <i>In Silico Biology</i> , 2004, 4, 517-28.	0.4	8
66	In vivo nuclear magnetic resonance study of citrate metabolism in <i>Propionibacterium freudenreichii</i> subsp. <i>shermanii</i> . <i>Journal of Dairy Research</i> , 1998, 65, 503-514.	0.7	5
67	DETECTION OF QTLs CONTROLLING MAJOR FRUIT QUALITY COMPONENTS IN PEACH WITHIN THE EUROPEAN PROJECT ISAFRUIT. <i>Acta Horticulturae</i> , 2009, , 533-538.	0.1	5
68	PROPIONIBACTERIUM spp., 2002, , 2330-2339.		4
69	¹ H-NMR metabolic profiling of wines from three cultivars, three soil types and two contrasting vintages. <i>Oeno One</i> , 2016, 41, 103.	0.7	4
70	PeakForest: a multi-platform digital infrastructure for interoperable metabolite spectral data and metadata management. <i>Metabolomics</i> , 2022, 18, .	1.4	4
71	MRSI vs CEST MRI to understand tomato metabolism in ripening fruit: is there a better contrast?. <i>Analytical and Bioanalytical Chemistry</i> , 2021, 413, 1251-1257.	1.9	3
72	Critical assessment of metabolism and related growth and quality traits in trout fed spirulina-supplemented plant-based diets. <i>Aquaculture</i> , 2022, 553, 738033.	1.7	3

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73	New Opportunities in Metabolomics and Biochemical Phenotyping for Plant Systems Biology. , 2012, , .		2
74	Aluminium stress disrupts metabolic performance of <i>Plantago almodavensis</i> plantlets transiently. <i>BioMetals</i> , 2015, 28, 997-1007.	1.8	2
75	¹ H NMR METABOLIC FINGERPRINTS OF GRAPE BERRIES PRODUCED IN DIFFERENT PLOTS IN BORDEAUX, FRANCE. <i>Acta Horticulturae</i> , 2005, , 257-264.	0.1	2
76	ISAFRUIT - STUDY OF THE GENETIC BASIS OF PRUNUS FRUIT QUALITY IN TWO PEACH AND TWO APRICOT POPULATIONS. <i>Acta Horticulturae</i> , 2009, , 523-528.	0.1	2
77	STUDY OF THE GENETIC BASIS OF PRUNUS FRUIT QUALITY ON TWO APRICOT AND TWO PEACH POPULATIONS. <i>Acta Horticulturae</i> , 2010, , 183-188.	0.1	0
78	A MULTI-LEVEL OMIC APPROACH OF TOMATO FRUIT QUALITY. <i>Acta Horticulturae</i> , 2015, , 793-800.	0.1	0