## Ana Carolina Maisonnave Arisi

List of Publications by Year in descending order

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73 papers

1,942 citations

304368 22 h-index 276539 41 g-index

74 all docs

74 docs citations

times ranked

74

2322 citing authors

#	Article	IF	CITATIONS
1	Manipulation of Glutathione and Amino Acid Biosynthesis in the Chloroplast. Plant Physiology, 1998, 118, 471-482.	2.3	190
2	Robust biological nitrogen fixation in a model grass–bacterial association. Plant Journal, 2015, 81, 907-919.	2.8	171
3	Modification of thiol contents in poplars ( Populus tremula $\tilde{A}-P$ . alba ) overexpressing enzymes involved in glutathione synthesis. Planta, 1997, 203, 362-372.	1.6	117
4	Responses to cadmium in leaves of transformed poplars overexpressing Γ -glutamylcysteine synthetase. Physiologia Plantarum, 2000, 109, 143-149.	2.6	90
5	Overexpression of Iron Superoxide Dismutase in Transformed Poplar Modifies the Regulation of Photosynthesis at Low CO2Partial Pressures or Following Exposure to the Prooxidant Herbicide Methyl Viologen1. Plant Physiology, 1998, 117, 565-574.	2.3	84
6	Light-dependent modulation of foliar glutathione synthesis and associated amino acid metabolism in poplar overexpressing $\hat{I}^3$ -glutamylcysteine synthetase. Planta, 1997, 202, 357-369.	1.6	76
7	Detecting authorized and unauthorized genetically modified organisms containing vip3A by real-time PCR and next-generation sequencing. Analytical and Bioanalytical Chemistry, 2014, 406, 2603-2611.	1.9	64
8	Quantification of Lactobacillus paracasei viable cells in probiotic yoghurt by propidium monoazide combined with quantitative PCR. International Journal of Food Microbiology, 2018, 264, 1-7.	2.1	51
9	Differential effects of short-term lindane administration on parameters related to oxidative stress in rat liver and erythrocytes. Journal of Biochemical Toxicology, 1993, 8, 187-194.	0.5	48
10	Differential growth responses of Brachypodium distachyon genotypes to inoculation with plant growth promoting rhizobacteria. Plant Molecular Biology, 2016, 90, 689-697.	2.0	48
11	Development and validation of real-time PCR screening methods for detection of cry1A.105 and cry2Ab2 genes in genetically modified organisms. Analytical and Bioanalytical Chemistry, 2011, 400, 1433-1442.	1.9	42
12	Cloning, Expression, Purification, and Characterization of a Novel Esterase from Lactobacillus plantarum. Molecular Biotechnology, 2010, 44, 242-249.	1.3	41
13	Ocurrence of Staphylococcus aureus and multiplex pcr detection of classic enterotoxin genes in cheese and meat products. Brazilian Journal of Microbiology, 2009, 40, 145-148.	0.8	38
14	Proteomic Analysis of Four Brazilian MON810 Maize Varieties and Their Four Non-Genetically-Modified Isogenic Varieties. Journal of Agricultural and Food Chemistry, 2011, 59, 11553-11559.	2.4	36
15	Genetic variability in four fish species (Pimelodus maculatus, Prochilodus lineatus, Salminus) Tj ETQq1 1 0.78431 and Technology, 2006, 49, 589-598.	4 rgBT /Ov 0.5	verlock 10 Tf 35
16	Photoinhibition of photosystem II in tobacco plants overexpressing glutathione reductase and poplars overexpressing superoxide dismutase. Physiologia Plantarum, 1999, 105, 409-416.	2.6	34
17	Gene expression analysis of maize seedlings (DKB240 variety) inoculated with plant growth promoting bacterium Herbaspirillum seropedicae. Symbiosis, 2014, 62, 41-50.	1,2	34
18	Genetic mapping of semi-polar metabolites in pepper fruits (Capsicum sp.): towards unravelling the molecular regulation of flavonoid quantitative trait loci. Molecular Breeding, 2014, 33, 503-518.	1.0	33

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19	Regression of morphological alterations and oxidative stress-related parameters after acute lindane-induced hepatotoxicity in rats. Toxicology, 1997, 117, 199-205.	2.0	30
20	Real-Time PCR Quantification of the Plant Growth Promoting Bacteria Herbaspirillum seropedicae Strain SmR1 in Maize Roots. Molecular Biotechnology, 2014, 56, 660-70.	1.3	29
21	Monitoring of GMO in Brazilian processed meat and soy-based products from 2007 to 2008. Journal of Food Composition and Analysis, 2010, 23, 226-229.	1.9	27
22	Acute lindane intoxication: A study on lindane tissue concentration and oxidative stress-related parameters in liver and erythrocytes. Journal of Biochemical Toxicology, 1994, 9, 9-15.	0.5	26
23	Comparison of real-time PCR assay and plate count for Lactobacillus paracasei enumeration in yoghurt. Annals of Microbiology, 2016, 66, 597-606.	1.1	26
24	Real time PCR detection targeting nifA gene of plant growth promoting bacteria Azospirillum brasilense strain FP2 in maize roots. Symbiosis, 2013, 61, 125-133.	1.2	23
25	A high-throughput method for GMO multi-detection using a microfluidic dynamic array. Analytical and Bioanalytical Chemistry, 2014, 406, 1397-1410.	1.9	23
26	Extraction of antifreeze proteins from cold acclimated leaves of Drimys angustifolia and their application to star fruit (Averrhoa carambola) freezing. Food Chemistry, 2019, 289, 65-73.	4.2	23
27	Recombinant DNA in meat additives: Specific detection of Roundup Readyâ,,¢ soybean by nested PCR. Journal of the Science of Food and Agriculture, 2007, 87, 1980-1984.	1.7	20
28	Safety assessment of plant varieties using transcriptomics profiling and a one-class classifier. Regulatory Toxicology and Pharmacology, 2014, 70, 297-303.	1.3	20
29	Monitoring of MON810 genetically modified maize in foods in Brazil from 2005 to 2007. Journal of Food Composition and Analysis, 2008, 21, 515-518.	1.9	19
30	Quantification of Roundup Readyâ,,¢ soybean in Brazilian soyâ€derived foods by realâ€time PCR. International Journal of Food Science and Technology, 2008, 43, 1027-1032.	1.3	19
31	Evaluation of polymerase chain reaction and DNA isolation protocols for detection of genetically modified soybean. International Journal of Food Science and Technology, 2007, 42, 1249-1255.	1.3	18
32	Identification of six differentially accumulated proteins of Zea mays seedlings (DKB240 variety) inoculated with Azospirillum brasilense strain FP2. European Journal of Soil Biology, 2013, 58, 45-50.	1.4	18
33	Comparative Proteomic Analysis of Two Varieties of Genetically Modified (GM) Embrapa 5.1 Common Bean ( <i>Phaseolus vulgaris</i> L.) and Their Non-GM Counterparts. Journal of Agricultural and Food Chemistry, 2015, 63, 10569-10577.	2.4	18
34	Proteome comparison for discrimination between honeydew and floral honeys from botanical species <i>Mimosa scabrella</i> Bentham by principal component analysis. Journal of the Science of Food and Agriculture, 2017, 97, 4515-4519.	1.7	18
35	Biochemical and Structural Characterization of Two Site-Directed Mutants of Staphylococcus xylosus Lipase. Molecular Biotechnology, 2010, 46, 168-175.	1.3	16
36	Structural stability of Staphylococcus xylosus lipase is modulated by Zn2+ ions. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2011, 1814, 1120-1126.	1,1	16

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37	Leaf proteome comparison of two ⟨scp⟩GM⟨ scp⟩ common bean varieties and their nonâ€ <scp⟩gm⟨ scp⟩ 2016,="" 927-932.<="" 96,="" agriculture,="" analysis.="" and="" by="" component="" counterparts="" food="" journal="" of="" principal="" science="" td="" the=""><td>1.7</td><td>16</td></scp⟩gm⟨ scp⟩>	1.7	16
38	Nested PCR detection of genetically modified soybean in soybean flour, infant formula and soymilk. LWT - Food Science and Technology, 2007, 40, 748-751.	2.5	15
39	Phenotypic and molecular characterization of Staphylococcus xylosus: technological potential for use in fermented sausage. Brazilian Archives of Biology and Technology, 2009, 52, 737-746.	0.5	15
40	The regulation of transcription of genes related to oxidative stress and glutathione synthesis in Zea mays leaves by nitric oxide. Biologia Plantarum, 2013, 57, 620-626.	1.9	15
41	Herbaspirillum seropedicae promotes maize growth but fails to control the maize leaf anthracnose. Physiology and Molecular Biology of Plants, 2019, 25, 167-176.	1.4	15
42	Development of Plasmid DNA Reference Material for the Quantification of Genetically Modified Common Bean Embrapa 5.1. Journal of Agricultural and Food Chemistry, 2013, 61, 4921-4926.	2.4	14
43	Expressed Proteins of Herbaspirillum seropedicae in Maize (DKB240) Roots-Bacteria Interaction Revealed Using Proteomics. Applied Biochemistry and Biotechnology, 2014, 174, 2267-2277.	1.4	14
44	Dose-dependent effects of acute lindane treatment on Kupffer cell function assessed in the isolated perfused rat liver. Xenobiotica, 1997, 27, 747-757.	0.5	13
45	Heterologous Expression and Purification of a Heat-Tolerant Staphylococcus xylosus Lipase. Molecular Biotechnology, 2010, 44, 110-119.	1.3	13
46	Primers and Probes Development for Specific PCR Detection of Genetically Modified Common Bean ( <i>Phaseolus vulgaris</i> ) Embrapa 5.1. Journal of Agricultural and Food Chemistry, 2012, 60, 4672-4677.	2.4	13
47	Microscopic and proteomic analysis of Zea mays roots (P30F53 variety) inoculated with Azospirillum brasilense strain FP2. Journal of Crop Science and Biotechnology, 2015, 18, 63-71.	0.7	13
48	Optimization of random amplified polymorphic DNA protocol for molecular identification of Lophius gastrophysus. Food Science and Technology, 2005, 25, 733-735.	0.8	11
49	Monitoring of Bt11 and Bt176 genetically modified maize in food sold commercially in Brazil from 2005 to 2007. Journal of the Science of Food and Agriculture, 2010, 90, 1566-1569.	1.7	11
50	Brain and liver lipid peroxidation levels following acute and short-term lindane administration in the rat. Toxicology Letters, 1994, 74, 61-68.	0.4	10
51	Comparison of Grain Proteome Profiles of Four Brazilian Common Bean ( <i>Phaseolus vulgaris</i> L.) Cultivars. Journal of Agricultural and Food Chemistry, 2017, 65, 7588-7597.	2.4	10
52	Application of propidium monoazide coupled with quantitative PCR to evaluate cell viability of Bifidobacterium animalis subsp. lactis in a non-dairy probiotic beverage. Annals of Microbiology, 2020, 70, .	1.1	10
53	Azospirillum brasilense viable cells enumeration using propidium monoazide-quantitative PCR. Archives of Microbiology, 2020, 202, 1653-1662.	1.0	10
54	Prolonged phenobarbital pretreatment abolishes the early oxidative stress component induced in the liver by acute lindane intoxication. Toxicology Letters, 2000, 115, 45-51.	0.4	9

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55	Phenotypic characterization and species-specific PCR of promising starter culture strains of Lactobacillus plantarum isolated from naturally fermented sausages. Brazilian Journal of Microbiology, 2007, 38, 547-552.	0.8	9
56	Development of an Event-Specific Hydrolysis Probe Quantitative Real-Time Polymerase Chain Reaction Assay for Embrapa 5.1 Genetically Modified Common Bean ( <i>Phaseolus vulgaris</i> ). Journal of Agricultural and Food Chemistry, 2014, 62, 11994-12000.	2.4	9
57	The role of glycine in determining the rate of glutathione synthesis in poplar. Possible implications for glutathione production during stress. Physiologia Plantarum, 1997, 100, 255-263.	2.6	8
58	Tuber proteome comparison of five potato varieties by principal component analysis. Journal of the Science of Food and Agriculture, 2016, 96, 3928-3936.	1.7	8
59	Bifidobacterium animalis ssp. lactis BB-12 enumeration by quantitative PCR assay in microcapsules with full-fat goat milk and inulin-type fructans. Food Research International, 2020, 133, 109131.	2.9	8
60	Immobilization of a Recombinant Esterase from Lactobacillus plantarum on Polypropylene Accurel MP1000. Applied Biochemistry and Biotechnology, 2011, 163, 304-312.	1.4	7
61	Sodium nitroprusside modulates gene expression involved in glutathione synthesis in Zea mays leaves. Biologia Plantarum, 2012, 56, 383-388.	1.9	7
62	Chemical characterization of liquid residues from aqueous enzymatic extraction of soybean oil. LWT - Food Science and Technology, 2013, 51, 51-58.	2.5	7
63	Tools to evaluate Herbaspirillum seropedicae abundance and nifH and rpoC expression in inoculated maize seedlings grown in vitro and in soil. Plant Growth Regulation, 2017, 83, 397-408.	1.8	7
64	Genetic similarity of Brazilian hull-less and malting barley varieties evaluated by RAPD markers. Scientia Agricola, 2005, 62, 36-39.	0.6	6
65	Antifreeze proteins in naturally cold acclimated leaves of Drimys angustifolia, Senecio icoglossus, and Eucalyptus ssp Brazilian Journal of Food Technology, 2016, 19, .	0.8	4
66	Protein profile and antioxidant capacity of processed seeds from two common bean ( <i>Phaseolus) Tj ETQq0 0 0</i>	rgBJ /Ove	erlgck 10 Tf 50
67	Azospirillum brasilense FP2 modulates respiratory burst oxidase gene expression in maize seedlings. Indian Journal of Plant Physiology, 2017, 22, 316-323.	0.8	3
68	Proteome Comparison of Grains from Two Maize Genotypes, with Colorless Kernel Pericarp ( <i>P1-ww</i> ) and Red Kernel Pericarp ( <i>P1-rr</i> ). Food Biotechnology, 2016, 30, 110-122.	0.6	2
69	New plasmid calibrators for geminivirus-resistant (EMB-PV051-1 event) common bean (Phaseolus) Tj ETQq1 1 0.7	784314 rg	BT <sub>2</sub> /Overlock
70	Inoculation of Herbaspirillum seropedicae strain SmR1 increases biomass in maize roots DKB 390 variety in the early stages of plant development. Archives of Microbiology, 2022, 204, .	1.0	2
71	High-added value co-products obtained from pecan nut (Carya illinoinensis) using a green extraction technology. Journal of Food Science and Technology, $0$ , $0$ , $1$ .	1.4	1
72	Applicability of quantitative polymerase chain reaction (qPCR) assays for common bean authentication in processed food. International Journal of Food Science and Technology, 2019, 54, 1381-1389.	1.3	0

 #	Article	IF	CITATIONS
73	Distribution of Genes Related to Probiotic Effects Across Lacticaseibacillus rhamnosus Revealed by Population Structure. Probiotics and Antimicrobial Proteins, 2021, , 1.	1.9	O