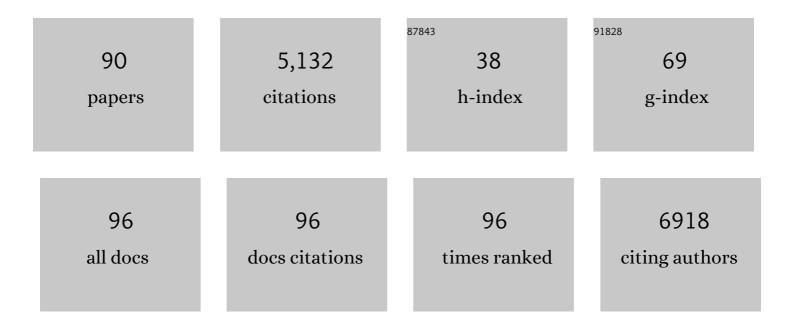
Maddalena Rossi

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Folate Production by Probiotic Bacteria. Nutrients, 2011, 3, 118-134.	1.7	459
2	Fermentation of Fructooligosaccharides and Inulin by Bifidobacteria: a Comparative Study of Pure and Fecal Cultures. Applied and Environmental Microbiology, 2005, 71, 6150-6158.	1.4	434
3	Antioxidant properties of potentially probiotic bacteria: in vitro and in vivo activities. Applied Microbiology and Biotechnology, 2013, 97, 809-817.	1.7	346
4	Folate Production by Bifidobacteria as a Potential Probiotic Property. Applied and Environmental Microbiology, 2007, 73, 179-185.	1.4	263
5	Antibiotic combination therapy in patients with chronic, treatment-resistant pouchitis. Alimentary Pharmacology and Therapeutics, 1999, 13, 713-718.	1.9	214
6	Characterization of Bifidobacterium strains for use in soymilk fermentation. International Journal of Food Microbiology, 1998, 39, 213-219.	2.1	157
7	In vitro transformation of chlorogenic acid by human gut microbiota. Molecular Nutrition and Food Research, 2014, 58, 1122-1131.	1.5	137
8	Wheat bran promotes enrichment within the human colonic microbiota of butyrateâ€producing bacteria that release ferulic acid. Environmental Microbiology, 2016, 18, 2214-2225.	1.8	119
9	Bioconversion of soy isoflavones daidzin and daidzein by Bifidobacterium strains. Applied Microbiology and Biotechnology, 2009, 81, 943-950.	1.7	117
10	Single cell oils of the cold-adapted oleaginous yeast Rhodotorula glacialis DBVPG 4785. Microbial Cell Factories, 2010, 9, 73.	1.9	111
11	Longitudinal Survey of Fungi in the Human Gut: ITS Profiling, Phenotyping, and Colonization. Frontiers in Microbiology, 2019, 10, 1575.	1.5	101
12	In vitro comparison of the prebiotic effects of two inulin-type fructans. Anaerobe, 2008, 14, 280-286.	1.0	99
13	Kinetics and Metabolism of Bifidobacterium adolescentis MB 239 Growing on Glucose, Galactose, Lactose, and Galactooligosaccharides. Applied and Environmental Microbiology, 2007, 73, 3637-3644.	1.4	97
14	Hydrolysis of the Rutinose-Conjugates Flavonoids Rutin and Hesperidin by the Gut Microbiota and Bifidobacteria. Nutrients, 2015, 7, 2788-2800.	1.7	94
15	Administration of Folate-Producing Bifidobacteria Enhances Folate Status in Wistar Rats ,. Journal of Nutrition, 2007, 137, 2742-2746.	1.3	93
16	Growth, lipid accumulation, and fatty acid composition in obligate psychrophilic, facultative psychrophilic, and mesophilic yeasts. FEMS Microbiology Ecology, 2009, 69, 363-372.	1.3	87
17	PCR detection of Bifidobacterium strains and Streptococcus thermophilus in feces of human subjects after oral bacteriotherapy and yogurt consumption. International Journal of Food Microbiology, 2003, 81, 203-209.	2.1	85
18	Cholesterol-lowering probiotics: in vitro selection and in vivo testing of bifidobacteria. Applied Microbiology and Biotechnology, 2013, 97, 8273-8281.	1.7	82

MADDALENA ROSSI

#	Article	IF	CITATIONS
19	Profiling of Protein Degraders in Cultures of Human Gut Microbiota. Frontiers in Microbiology, 2019, 10, 2614.	1.5	74
20	Mining metagenomic whole genome sequences revealed subdominant but constant <i>Lactobacillus</i> population in the human gut microbiota. Environmental Microbiology Reports, 2016, 8, 399-406.	1.0	72
21	Assessment of In-Line Near-Infrared Spectroscopy for Continuous Monitoring of Fermentation Processes. Biotechnology Progress, 2003, 19, 1816-1821.	1.3	68
22	Characterization of the plasmid pMB1 from Bifidobacterium longum and its use for shuttle vector construction. Research in Microbiology, 1996, 147, 133-143.	1.0	67
23	Identification of mucin degraders of the human gut microbiota. Scientific Reports, 2021, 11, 11094.	1.6	67
24	Getting lipids from glycerol: new perspectives on biotechnological exploitation of Candida freyschussii. Microbial Cell Factories, 2014, 13, 83.	1.9	60
25	Fermentation of xylo-oligosaccharides by Bifidobacterium adolescentis DSMZ 18350: kinetics, metabolism, and β-xylosidase activities. Applied Microbiology and Biotechnology, 2013, 97, 3109-3117.	1.7	58
26	Role of bifidobacteria in the hydrolysis of chlorogenic acid. MicrobiologyOpen, 2015, 4, 41-52.	1.2	55
27	Bifidobacteria supplementation: Effects on plasma lipid profiles in dyslipidemic children. Nutrition, 2014, 30, 831-836.	1.1	54
28	Substrate preference of Bifidobacterium adolescentis MB 239: compared growth on single and mixed carbohydrates. Applied Microbiology and Biotechnology, 2006, 73, 654-662.	1.7	53
29	Detection of novel metabolites of flaxseed lignans in vitro and in vivo. Molecular Nutrition and Food Research, 2016, 60, 1590-1601.	1.5	47
30	Specific Detection of Bifidobacterium Strains in a Pharmaceutical Probiotic Product and in Human Feces by Polymerase Chain Reaction. Systematic and Applied Microbiology, 2000, 23, 391-399.	1.2	46
31	Role of bifidobacteria in the activation of the lignan secoisolariciresinol diglucoside. Applied Microbiology and Biotechnology, 2011, 92, 159-168.	1.7	46
32	Characterization of the peptide fraction from digested Parmigiano Reggiano cheese and its effect on growth of lactobacilli and bifidobacteria. International Journal of Food Microbiology, 2017, 255, 32-41.	2.1	46
33	Lactic acid bacteria as protective cultures in fermented pork meat to prevent Clostridium spp. growth. International Journal of Food Microbiology, 2016, 235, 53-59.	2.1	45
34	Nucleotide sequence, expression and transcriptional analysis of the Bifidobacterium longum MB 219 lacZ gene. Archives of Microbiology, 2000, 174, 74-80.	1.0	44
35	Antibiotic Resistance, Virulence Factors, Phenotyping, and Genotyping of E. coli Isolated from the Feces of Healthy Subjects. Microorganisms, 2019, 7, 251.	1.6	43
36	Comparison of formulaâ€fed infants with and without colic revealed significant differences in total bacteria, <i>Enterobacteriaceae</i> and faecal ammonia. Acta Paediatrica, International Journal of Paediatrics, 2017, 106, 573-578.	0.7	42

Maddalena Rossi

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37	Enhanced production of L-(+)-lactic acid in chemostat byLactobacillus casei DSM 20011 using ion-exchange resins and cross-flow filtration in a fully automated pilot plant controlled via NIR. Biotechnology and Bioengineering, 2000, 67, 147-156.	1.7	41
38	An efficient transformation system for Bifidobacterium spp Letters in Applied Microbiology, 1997, 24, 33-36.	1.0	39
39	Improved cloning vectors for Bifidobacterium spp Letters in Applied Microbiology, 1998, 26, 101-104.	1.0	39
40	Conjugated Linoleic Acid Production by Bifidobacteria: Screening, Kinetic, and Composition. BioMed Research International, 2016, 2016, 1-8.	0.9	39
41	Growth kinetics on oligo- and polysaccharides and promising features of three antioxidative potential probiotic strains. Journal of Applied Microbiology, 2008, 105, 1266-1276.	1.4	35
42	Microbiota of sliced cooked ham packaged in modified atmosphere throughout the shelf life. International Journal of Food Microbiology, 2019, 289, 200-208.	2.1	35
43	Effect of Rearing Temperature on Growth and Microbiota Composition of Hermetia illucens. Microorganisms, 2020, 8, 902.	1.6	33
44	The effect ofn-alkanes in the degradation of dibenzothiophene and of organic sulfur compounds in heavy oil by aPseudomonas sp Biotechnology Letters, 1992, 14, 515-520.	1.1	32
45	Potential Impact of Probiotic Consumption on the Bioactivity of Dietary Phytochemicals. Journal of Agricultural and Food Chemistry, 2013, 61, 130924093716009.	2.4	32
46	Antibiotic Resistance, Virulence Factors, Phenotyping, and Genotyping of Non-Escherichia coli Enterobacterales from the Gut Microbiota of Healthy Subjects. International Journal of Molecular Sciences, 2020, 21, 1847.	1.8	32
47	The Probiotic <i>Bifidobacterium breve</i> B632 Inhibited the Growth of <i>Enterobacteriaceae</i> within Colicky Infant Microbiota Cultures. BioMed Research International, 2014, 2014, 1-7.	0.9	31
48	Thermal adaptability of Kluyveromyces marxianus in recombinant protein production. Microbial Cell Factories, 2013, 12, 34.	1.9	29
49	Comparison of culture-dependent and independent approaches to characterize fecal bifidobacteria and lactobacilli. Anaerobe, 2016, 38, 130-137.	1.0	29
50	Characterization and molecular cloning of Bifidobacterium longum cryptic plasmid pMB1. Letters in Applied Microbiology, 1990, 11, 220-223.	1.0	27
51	Production of l(+) and d(â^') lactic acid isomers by Lactobacillus casei subsp. casei DSM 20011 and Lactobacillus coryniformis subsp. torquens DSM 20004 in continuous fermentation. Journal of Bioscience and Bioengineering, 1996, 81, 548-552.	0.9	27
52	Prebiotic effects of a wheat germ preparation in human healthy subjects. Food Microbiology, 2004, 21, 119-124.	2.1	27
53	Fermentative production of superoxide dismutase with Kluyveromyces marxianus. Journal of Industrial Microbiology and Biotechnology, 2006, 34, 27-34.	1.4	27
54	Evolution of microbial community and chemical properties of a sourdough during the production of Colomba, an Italian sweet leavened baked product. LWT - Food Science and Technology, 2017, 86, 31-39.	2.5	27

MADDALENA ROSSI

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55	Getting Lipids for Biodiesel Production from Oleaginous Fungi. , 0, , .		26
56	Bacterial community of industrial raw sausage packaged in modified atmosphere throughout the shelf life. International Journal of Food Microbiology, 2018, 280, 78-86.	2.1	24
57	SOD1, a New <i>Kluyveromyces lactis</i> Helper Gene for Heterologous Protein Secretion. Applied and Environmental Microbiology, 2008, 74, 7130-7137.	1.4	21
58	Comparison of gluten peptides and potential prebiotic carbohydrates in old and modern Triticum turgidum ssp. genotypes. Food Research International, 2019, 120, 568-576.	2.9	21
59	Zinc Uptake by Lactic Acid Bacteria. ISRN Biotechnology, 2013, 2013, 1-5.	1.9	21
60	Cloning of the gene for cholesterol oxidase in Bacillus spp., Lactobacillus reuteri and its expression in Escherichia coli. Letters in Applied Microbiology, 1993, 17, 61-64.	1.0	20
61	A new approach to the synthesis of rare thiazino[6,5-b]indol-4-one derivatives. First total synthesis of the indole phytoalexin cyclobrassinon. Tetrahedron, 2002, 58, 9029-9039.	1.0	19
62	Secretion of Kluyveromyces lactis Cu/Zn SOD: strategies for enhanced production. Applied Microbiology and Biotechnology, 2010, 86, 871-878.	1.7	19
63	Riboflavin Biosynthesis and Overproduction by a Derivative of the Human Gut Commensal Bifidobacterium longum subsp. infantis ATCC 15697. Frontiers in Microbiology, 2020, 11, 573335.	1.5	18
64	Barrier and carrier effects of n-dodecane on the anaerobic degradation of benzothiophene by Desulfovibrio desulfuricans. Biotechnology Letters, 1993, 15, 527-530.	1.1	17
65	Characterization of the superoxide dismutase SOD1 gene of Kluyveromyces marxianus L3 and improved production of SOD activity. Applied Microbiology and Biotechnology, 2008, 77, 1269-1277.	1.7	17
66	β-Glucuronidase Pattern Predicted From Gut Metagenomes Indicates Potentially Diversified Pharmacomicrobiomics. Frontiers in Microbiology, 2022, 13, 826994.	1.5	17
67	Enoate reductases from non conventional yeasts: Bioconversion, cloning, and functional expression in Saccharomyces cerevisiae. Journal of Biotechnology, 2011, 156, 279-285.	1.9	16
68	Potential prebiotic effect of a long-chain dextran produced by <i>Weissella cibaria</i> : an <i>inÂvitro</i> evaluation. International Journal of Food Sciences and Nutrition, 2020, 71, 563-571.	1.3	16
69	Characterization of Gram-positive broad host-range plasmids carrying a thermophilic replicon. Research in Microbiology, 1991, 142, 389-396.	1.0	15
70	Stability of recombinant plasmids on the continuous culture ofBifidobacterium animalisATCC 27536. Biotechnology and Bioengineering, 2003, 84, 145-150.	1.7	13
71	Investigation on the antimicrobial properties of ceriumâ€doped bioactive glasses. Journal of Biomedical Materials Research - Part A, 2022, 110, 504-508.	2.1	13
72	Study of stability of recombinant plasmids during the continuous culture ofBacillus stearothermophilus NUB3621 in nonselective medium. , 1997, 53, 507-514.		12

5

Maddalena Rossi

#	Article	IF	CITATIONS
73	Flavone and xanthone derivatives related to fluoroquinolones. Il Farmaco, 1999, 54, 411-415.	0.9	11
74	Comparative Genomics of Leuconostoc carnosum. Frontiers in Microbiology, 2020, 11, 605127.	1.5	11
75	Vaginal and Anal Microbiome during Chlamydia trachomatis Infections. Pathogens, 2021, 10, 1347.	1.2	11
76	A new photocyclization approach to the rare 1,3-thiazino[6,5-b]indol-4-one derivatives. Tetrahedron Letters, 2001, 42, 9281-9283.	0.7	10
77	Recombinant S. cerevisiae expressing Old Yellow Enzymes from non-conventional yeasts: an easy system for selective reduction of activated alkenes. Microbial Cell Factories, 2014, 13, 60.	1.9	10
78	Multivariate Analysis in Microbiome Description: Correlation of Human Gut Protein Degraders, Metabolites, and Predicted Metabolic Functions. Frontiers in Microbiology, 2021, 12, 723479.	1.5	9
79	Allylamine Type Xanthone Antimycotics. Archiv Der Pharmazie, 1998, 331, 225-227.	2.1	8
80	Anti-Listeria Starters: In Vitro Selection and Production Plant Evaluation. Journal of Food Protection, 2014, 77, 837-842.	0.8	8
81	Draft Genome Sequences of 12 Leuconostoc carnosum Strains Isolated from Cooked Ham Packaged in a Modified Atmosphere and from Fresh Sausages. Microbiology Resource Announcements, 2020, 9, .	0.3	6
82	Gut mobilization improves behavioral symptoms and modulates urinary p resol in chronically constipated autistic children: A prospective study. Autism Research, 2021, , .	2.1	6
83	Microbiota Survey of Sliced Cooked Ham During the Secondary Shelf Life. Frontiers in Microbiology, 2022, 13, 842390.	1.5	6
84	Phenotypic Traits and Immunomodulatory Properties of Leuconostoc carnosum Isolated From Meat Products. Frontiers in Microbiology, 2021, 12, 730827.	1.5	5
85	Rapid method for screening enoate reductase activity in yeasts. Journal of Microbiological Methods, 2010, 83, 106-110.	0.7	4
86	Production of Single Cell Oils from Glycerol By Oleaginous Yeasts. Journal of Biotechnology, 2010, 150, 389-389.	1.9	3
87	Complementary microbial approaches for the preparation of optically pure aromatic molecules. Annals of Microbiology, 2013, 63, 1021-1027.	1.1	2
88	Mining metagenomic whole genome sequences revealed subdominant but constant <i>Lactobacillus</i> population in the human gut microbiota. Environmental Microbiology, 2016, , n/a-n/a.	1.8	2
89	In Vitro Assessment of Prebiotic Activity. Methods in Molecular Biology, 2021, 2278, 209-223.	0.4	1
90	Draft Genome Sequence of the Mucin Degrader Clostridium tertium WC0709. Microbiology Resource Announcements, 2021, 10, e0064221.	0.3	1