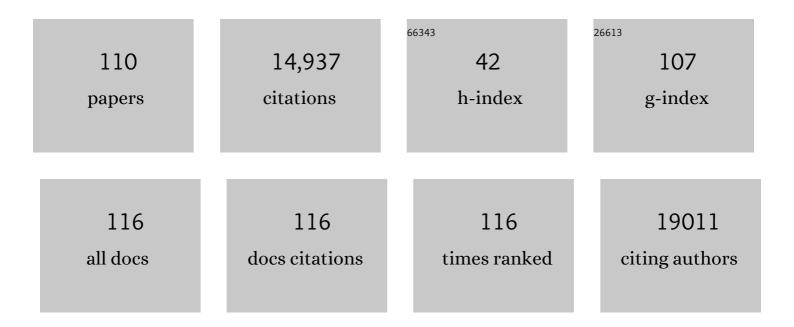
List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Enterotypes of the human gut microbiome. Nature, 2011, 473, 174-180.	27.8	5,800
2	Dietary intervention impact on gut microbial gene richness. Nature, 2013, 500, 585-588.	27.8	1,485
3	Microbiome definition re-visited: old concepts and new challenges. Microbiome, 2020, 8, 103.	11.1	903
4	Stress responses in lactic acid bacteria. Antonie Van Leeuwenhoek, 2002, 82, 187-216.	1.7	598
5	Entry of Listeria monocytogenes into hepatocytes requires expression of InIB, a surface protein of the internalin multigene family. Molecular Microbiology, 1995, 16, 251-261.	2.5	464
6	A reference gene catalogue of the pig gut microbiome. Nature Microbiology, 2016, 1, 16161.	13.3	416
7	The complete genome sequence of <i>Lactobacillus bulgaricus</i> reveals extensive and ongoing reductive evolution. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 9274-9279.	7.1	382
8	Plasmid Addiction Genes of Bacteriophage P1: doc, which Causes Cell Death on Curing of Prophage, and phd, which Prevents Host Death when Prophage is Retained. Journal of Molecular Biology, 1993, 233, 414-428.	4.2	263
9	An M protein with a single C repeat prevents phagocytosis of Streptococcus pyogenes: use of a temperature-sensitive shuttle vector to deliver homologous sequences to the chromosome of S. pyogenes. Molecular Microbiology, 1993, 8, 809-819.	2.5	228
10	Stress responses in lactic acid bacteria. Antonie Van Leeuwenhoek, 2002, 82, 187-216.	1.7	180
11	Indole, a Signaling Molecule Produced by the Gut Microbiota, Negatively Impacts Emotional Behaviors in Rats. Frontiers in Neuroscience, 2018, 12, 216.	2.8	179
12	Acid- and multistress-resistant mutants of Lactococcus lactis : identification of intracellular stress signals. Molecular Microbiology, 2002, 35, 517-528.	2.5	178
13	Lactococcus lactis and stress. Antonie Van Leeuwenhoek, 1996, 70, 243-251.	1.7	158
14	Microbial impact on cholesterol and bile acid metabolism: current status and future prospects. Journal of Lipid Research, 2019, 60, 323-332.	4.2	149
15	Postgenomic Analysis of <i>Streptococcus thermophilus</i> Cocultivated in Milk with <i>Lactobacillus delbrueckii</i> subsp. <i>bulgaricus</i> : Involvement of Nitrogen, Purine, and Iron Metabolism. Applied and Environmental Microbiology, 2009, 75, 2062-2073.	3.1	130
16	Beneficial metabolic effects of selected probiotics on dietâ€induced obesity and insulin resistance in mice are associated with improvement of dysbiotic gut microbiota. Environmental Microbiology, 2016, 18, 1484-1497.	3.8	127
17	The intestinal microbiota regulates host cholesterol homeostasis. BMC Biology, 2019, 17, 94.	3.8	125
18	Prediction of surface exposed proteins in <i>Streptococcus pyogenes</i> , with a potential application to other Gramâ€positive bacteria. Proteomics, 2009, 9, 61-73.	2.2	123

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19	Reduced obesity, diabetes, and steatosis upon cinnamon and grape pomace are associated with changes in gut microbiota and markers of gut barrier. American Journal of Physiology - Endocrinology and Metabolism, 2018, 314, E334-E352.	3.5	119
20	Proteomic characterization of the acid tolerance response inLactococcus lactis MG1363. Proteomics, 2005, 5, 4794-4807.	2.2	98
21	Mucosal-associated invariant T cells promote inflammation and intestinal dysbiosis leading to metabolic dysfunction during obesity. Nature Communications, 2020, 11, 3755.	12.8	97
22	Rerouting of pyruvate metabolism during acid adaptation in <i>Lactobacillus bulgaricus</i> . Proteomics, 2008, 8, 3154-3163.	2.2	93
23	Electrotransformation of <i>Lactobacillus delbrueckii</i> subsp. <i>bulgaricus</i> and <i>L. delbrueckii</i> subsp. <i>lactis</i> with Various Plasmids. Applied and Environmental Microbiology, 2002, 68, 46-52.	3.1	90
24	Production of a Heterologous Nonheme Catalase by Lactobacillus casei : an Efficient Tool for Removal of H 2 O 2 and Protection of Lactobacillus bulgaricus from Oxidative Stress in Milk. Applied and Environmental Microbiology, 2006, 72, 5143-5149.	3.1	90
25	Physiological Study of Lactobacillus delbrueckii subsp. bulgaricus Strains in a Novel Chemically Defined Medium. Applied and Environmental Microbiology, 2000, 66, 5306-5311.	3.1	86
26	AGMIAL: implementing an annotation strategy for prokaryote genomes as a distributed system. Nucleic Acids Research, 2006, 34, 3533-3545.	14.5	84
27	Identification and Disruption of Two Discrete Loci Encoding Hyaluronic Acid Capsule Biosynthesis GeneshasA, hasB, and hasC inStreptococcus uberis. Infection and Immunity, 2001, 69, 392-399.	2.2	82
28	Isolation and Characterization of a Plasmid fromLactobacillus fermentumConferring Erythromycin Resistance. Plasmid, 1997, 37, 199-203.	1.4	81
29	Identification of stress-inducible proteins inLactobacillus delbrueckii subsp.bulgaricus. Electrophoresis, 2000, 21, 2557-2561.	2.4	78
30	Characterization of urease genes cluster of Streptococcus thermophilus. Journal of Applied Microbiology, 2004, 96, 209-219.	3.1	78
31	Glyphosate and glyphosateâ€based herbicide exposure during the peripartum period affects maternal brain plasticity, maternal behaviour and microbiome. Journal of Neuroendocrinology, 2019, 31, e12731.	2.6	69
32	Anti-inflammatory properties of dairy lactobacilli. Inflammatory Bowel Diseases, 2012, 18, 657-666.	1.9	68
33	Comparison of the thickening properties of fourLactobacillus delbrueckiisubsp.bulgaricusstrains and physicochemical characterization of their exopolysaccharides. FEMS Microbiology Letters, 2003, 221, 285-291.	1.8	66
34	Lactic acid bacteria and proteomics: current knowledge and perspectives. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2002, 771, 329-342.	2.3	63
35	Fecal Serine Protease Profiling in Inflammatory Bowel Diseases. Frontiers in Cellular and Infection Microbiology, 2020, 10, 21.	3.9	62
36	The acid tolerant l-arabinose isomerase from the food grade Lactobacillus sakei 23K is an attractive d-tagatose producer. Bioresource Technology, 2010, 101, 9171-9177.	9.6	60

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37	Characterization of glucansucrase and dextran from Weissella sp. TN610 with potential as safe food additives. International Journal of Biological Macromolecules, 2013, 52, 125-132.	7.5	60
38	Combining selected immunomodulatory <i>Propionibacterium freudenreichii</i> and <i>Lactobacillus delbrueckii</i> strains: Reverse engineering development of an antiâ€inflammatory cheese. Molecular Nutrition and Food Research, 2016, 60, 935-948.	3.3	60
39	Lactobacillus delbrueckii ssp. lactis and ssp. bulgaricus: a chronicle of evolution in action. BMC Genomics, 2014, 15, 407.	2.8	59
40	Microbiome innovations for a sustainable future. Nature Microbiology, 2021, 6, 138-142.	13.3	53
41	Physiology of <i>Streptococcus thermophilus</i> during the late stage of milk fermentation with special regard to sulfur aminoâ€acid metabolism. Proteomics, 2008, 8, 4273-4286.	2.2	50
42	Production of growth-inhibiting factors by Lactobacillus delbrueckii. Journal of Applied Microbiology, 2001, 91, 147-153.	3.1	49
43	Local and Systemic Immune Mechanisms Underlying the Anti-Colitis Effects of the Dairy Bacterium Lactobacillus delbrueckii. PLoS ONE, 2014, 9, e85923.	2.5	45
44	Rational design of Bacillus stearothermophilus US100 l-arabinose isomerase: Potential applications for d-tagatose production. Biochimie, 2009, 91, 650-653.	2.6	44
45	Production of d-tagatose, a low caloric sweetener during milk fermentation using l-arabinose isomerase. Bioresource Technology, 2011, 102, 3309-3315.	9.6	43
46	In Vitro Characterization of Gut Microbiota-Derived Commensal Strains: Selection of Parabacteroides distasonis Strains Alleviating TNBS-Induced Colitis in Mice. Cells, 2020, 9, 2104.	4.1	43
47	Conservation of key elements of natural competence inLactococcus lactisssp FEMS Microbiology Letters, 2006, 257, 32-42.	1.8	37
48	Efficient bioconversion of lactose in milk and whey: immobilization and biochemical characterization of a β-galactosidase from the dairy Streptococcus thermophilus LMD9 strain. Research in Microbiology, 2010, 161, 515-525.	2.1	36
49	Genome Sequence of " <i>Candidatus</i> Arthromitus―sp. Strain SFB-Mouse-NL, a Commensal Bacterium with a Key Role in Postnatal Maturation of Gut Immune Functions. Genome Announcements, 2014, 2, .	0.8	35
50	A catalog of microbial genes from the bovine rumen unveils a specialized and diverse biomass-degrading environment. GigaScience, 2020, 9, .	6.4	35
51	Transcriptional analysis of the cyclopropane fatty acid synthase gene ofLactococcus lactisMG1363 at low pH. FEMS Microbiology Letters, 2005, 250, 189-194.	1.8	33
52	Proteome phenotyping of acid stress-resistant mutants ofLactococcus lactis MG1363. Proteomics, 2007, 7, 2038-2046.	2.2	33
53	Siropins, novel serine protease inhibitors from gut microbiota acting on human proteases involved in inflammatory bowel diseases. Microbial Cell Factories, 2016, 15, 201.	4.0	33
54	Bile Salt Hydrolases: At the Crossroads of Microbiota and Human Health. Microorganisms, 2021, 9, 1122.	3.6	33

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55	Development of Microbiome Biobanks – Challenges and Opportunities. Trends in Microbiology, 2021, 29, 89-92.	7.7	31
56	Extensive horizontal transfer of core genome genes between two Lactobacillus species found in the gastrointestinal tract. BMC Evolutionary Biology, 2007, 7, 141.	3.2	29
57	Induction of Heavy-Metal-Transporting CPX-Type ATPases during Acid Adaptation in Lactobacillus bulgaricus. Applied and Environmental Microbiology, 2006, 72, 7445-7454.	3.1	28
58	The acid-tolerant L-arabinose isomerase from the mesophilic Shewanella sp. ANA-3 is highly active at low temperatures. Microbial Cell Factories, 2011, 10, 96.	4.0	28
59	Microbial Reduction of Cholesterol to Coprostanol: An Old Concept and New Insights. Catalysts, 2019, 9, 167.	3.5	28
60	Stress responses in lactic acid bacteria. , 2002, , 187-216.		28
61	Development of an EfficientIn VivoSystem (Pjunc-TpaseIS1223) for Random Transposon Mutagenesis of Lactobacillus casei. Applied and Environmental Microbiology, 2012, 78, 5417-5423.	3.1	27
62	Heterologous expression and optimization using experimental designs allowed highly efficient production of the PHY US417 phytase in Bacillus subtilis 168. AMB Express, 2012, 2, 10.	3.0	27
63	Crucial role of Pro 257 in the thermostability of Bacillus phytases: Biochemical and structural investigation. International Journal of Biological Macromolecules, 2013, 54, 9-15.	7.5	26
64	Genetic structure and transcriptional analysis of the arginine deiminase (ADI) cluster inLactococcus lactisMG1363. Canadian Journal of Microbiology, 2006, 52, 617-622.	1.7	25
65	The attractive recombinant phytase from Bacillus licheniformis: biochemical and molecular characterization. Applied Microbiology and Biotechnology, 2014, 98, 5937-5947.	3.6	24
66	Exploring the acidotolerance of β-galactosidase from Lactobacillus delbrueckii subsp. bulgaricus: an attractive enzyme for lactose bioconversion. Research in Microbiology, 2009, 160, 775-784.	2.1	23
67	Discriminatory antibacterial effects of calix[n]arene capped silver nanoparticles with regard to Gram positive and Gram negative bacteria. Chemical Communications, 2013, 49, 7150.	4.1	21
68	Serine protease inhibitors and human wellbeing interplay: new insights for old friends. PeerJ, 2019, 7, e7224.	2.0	20
69	Multiple Selection Criteria for Probiotic Strains with High Potential for Obesity Management. Nutrients, 2021, 13, 713.	4.1	19
70	Bile Acids: Key Players in Inflammatory Bowel Diseases?. Cells, 2022, 11, 901.	4.1	19
71	The acid tolerant and cold-active β-galactosidase from Lactococcus lactis strain is an attractive biocatalyst for lactose hydrolysis. Antonie Van Leeuwenhoek, 2013, 103, 701-712.	1.7	18
72	Serine proteases at the cutting edge of IBD: Focus on gastrointestinal inflammation. FASEB Journal, 2020, 34, 7270-7282.	0.5	18

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73	Exploring the Bacterial Impact on Cholesterol Cycle: A Numerical Study. Frontiers in Microbiology, 2020, 11, 1121.	3.5	17
74	Identification of New Potential Biotherapeutics from Human Gut Microbiota-Derived Bacteria. Microorganisms, 2021, 9, 565.	3.6	16
75	csp-like genes ofLactobacillus delbrueckiissp.bulgaricusand their response to cold shock. FEMS Microbiology Letters, 2003, 226, 323-330.	1.8	15
76	Bacillus phytases: Current status and future prospects. Bioengineered, 2015, 6, 233-236.	3.2	15
77	Proteomic analysis of spontaneous mutants of <i><scp>L</scp>actococcus lactis</i> : Involvement of <scp>GAPDH</scp> and arginine deiminase pathway in H ₂ O ₂ resistance. Proteomics, 2012, 12, 1792-1805.	2.2	14
78	High-Throughput System for the Presentation of Secreted and Surface-Exposed Proteins from Gram-Positive Bacteria in Functional Metagenomics Studies. PLoS ONE, 2013, 8, e65956.	2.5	14
79	Transposition in Lactobacillus delbrueckii subsp. bulgaricus: identification of two thermosensitive replicons and two functional insertion sequences. Microbiology (United Kingdom), 2003, 149, 1503-1511.	1.8	13
80	P1016 The pig's other genome: A reference gene catalog of the gut microbiome as a new resource for deep studies of the interplay between the host and its microbiome. Journal of Animal Science, 2016, 94, 22-22.	0.5	13
81	Bacterial L-Arabinose Isomerases: Industrial Application for D-Tagatose Production. Recent Patents on DNA & Gene Sequences, 2011, 5, 194-201.	0.7	13
82	The secreted l-arabinose isomerase displays anti-hyperglycemic effects in mice. Microbial Cell Factories, 2015, 14, 204.	4.0	12
83	Highly efficient production of the staphylococcal nuclease reporter in <i>Lactobacillus bulgaricus</i> governed by the promoter of the <i>hlbA</i> gene. FEMS Microbiology Letters, 2009, 293, 232-239.	1.8	11
84	Selection of Lactobacillus plantarum TN627 as a new probiotic candidate based on in vitro functional properties. Biotechnology and Bioprocess Engineering, 2011, 16, 1115-1123.	2.6	11
85	Fatâ€Shaped Microbiota Affects Lipid Metabolism, Liver Steatosis, and Intestinal Homeostasis in Mice Fed a Lowâ€Protein Diet. Molecular Nutrition and Food Research, 2020, 64, e1900835.	3.3	11
86	Digestive Inflammation: Role of Proteolytic Dysregulation. International Journal of Molecular Sciences, 2021, 22, 2817.	4.1	10
87	Gut Serpinome: Emerging Evidence in IBD. International Journal of Molecular Sciences, 2021, 22, 6088.	4.1	10
88	Microbiome Research as an Effective Driver of Success Stories in Agrifood Systems – A Selection of Case Studies. Frontiers in Microbiology, 0, 13, .	3.5	10
89	Supramolecular stabilization of acid tolerant l-arabinose isomerase from Lactobacillus sakei. Chemical Communications, 2011, 47, 12307.	4.1	9
90	The Enterococcus faecalis virulence factor ElrA interacts with the human Four-and-a-Half LIM Domains Protein 2. Scientific Reports, 2017, 7, 4581.	3.3	9

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91	Functional Comparison of Bacteria from the Human Gut and Closely Related Non-Gut Bacteria Reveals the Importance of Conjugation and a Paucity of Motility and Chemotaxis Functions in the Gut Environment. PLoS ONE, 2016, 11, e0159030.	2.5	9
92	Patented Biotechnological Applications of Serpin: an Update. Recent Patents on DNA & Gene Sequences, 2013, 7, 137-143.	0.7	8
93	Tolerogenic Dendritic Cells Shape a Transmissible Gut Microbiota That Protects From Metabolic Diseases. Diabetes, 2021, 70, 2067-2080.	0.6	7
94	Domestic Environment and Gut Microbiota: Lessons from Pet Dogs. Microorganisms, 2022, 10, 949.	3.6	7
95	Unprecedented large inverted repeats at the replication terminus of circular bacterial chromosomes suggest a novel mode of chromosome rescue. Scientific Reports, 2017, 7, 44331.	3.3	6
96	Sildenafil citrate long-term treatment effects on cardiovascular reactivity in a SHR experimental model of metabolic syndrome. PLoS ONE, 2019, 14, e0223914.	2.5	6
97	<i>para</i> -Sulphonato-calix[<i>n</i>]arene capped silver nanoparticles challenge the catalytic efficiency and the stability of a novel human gut serine protease inhibitor. Chemical Communications, 2019, 55, 8935-8938.	4.1	5
98	Identification de protéines de stress chez Lactobacillus delbrueckii bulgaricus par électrophorèse bidimensionnelle. Dairy Science and Technology, 2001, 81, 317-325.	0.9	5
99	Genome Sequence of Lactobacillus delbrueckii subsp. <i>lactis</i> CNRZ327, a Dairy Bacterium with Anti-Inflammatory Properties. Genome Announcements, 2014, 2, .	0.8	4
100	Biotechnological Applications of Serine Proteases: A Patent Review. Recent Patents on Biotechnology, 2018, 12, 280-287.	0.8	4
101	SP-1, a Serine Protease from the Gut Microbiota, Influences Colitis and Drives Intestinal Dysbiosis in Mice. Cells, 2021, 10, 2658.	4.1	4
102	Testing of a whole genome PCR scanning approach to identify genomic variability in four different species of lactic acid bacteria. Research in Microbiology, 2006, 157, 386-394.	2.1	3
103	Effect of a <i>guaA</i> mutation on the acid tolerance of <i>L. lactis</i> . Sciences Des Aliments, 2002, 22, 67-74.	0.2	3
104	Recent Patents on Hypocholesterolemic Therapeutic Strategies: An Update. Recent Advances in DNA & Gene Sequences, 2016, 9, 36-44.	0.7	2
105	Mutants de Lactococcus lactis résistants à l'acidité. Dairy Science and Technology, 1998, 78, 157-163.	0.9	1
106	Exopolysaccharides de Lactobacillus delbrueckii subsp. bulgaricus. Sciences Des Aliments, 2002, 22, 143-149.	0.2	1
107	Functional Metagenomics of Bacterial-Cell Crosstalk. , 2013, , 1-6.		0
108	Relevant Patented Biotechnological Applications of Ecotin: An Update. Recent Patents on Biotechnology, 2018, 12, 233-238.	0.8	0

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109	Targeting the Gut Microbiota in Metabolic Disorders and Juvenile Growth. , 2019, , 441-462.		0
110	Editorial: Probiotic Trigger Molecules in Action. Frontiers in Microbiology, 2021, 12, 789209.	3.5	0