## Miguel Gueimonde

List of Publications by Year in descending order

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		14614	16605
208	17,137	66	123
papers	citations	h-index	g-index
214	214	214	17662
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Daily ingestion of Akkermansia mucciniphila for oneÂmonth promotes healthy aging and increases lifespan in old female mice. Biogerontology, 2022, 23, 35-52.	2.0	19
2	In Vitro Probiotic Modulation of the Intestinal Microbiota and 2′Fucosyllactose Consumption in Fecal Cultures from Infants at Two Months of Age. Microorganisms, 2022, 10, 318.	1.6	7
3	Needle in a Whey-Stack: PhRACS as a Discovery Tool for Unknown Phage-Host Combinations. MBio, 2022, 13, e0333421.	1.8	5
4	Gut Microbiome Characteristics in feral and domesticated horses from different geographic locations. Communications Biology, 2022, 5, 172.	2.0	20
5	The Therapeutic Role of Exercise and Probiotics in Stressful Brain Conditions. International Journal of Molecular Sciences, 2022, 23, 3610.	1.8	8
6	Early Life Nutrition and the Role of Complementary Feeding on Later Adherence to the Mediterranean Diet in Children up to 3 Years of Age. Nutrients, 2022, 14, 1664.	1.7	2
7	Maternal Diet Is Associated with Human Milk Oligosaccharide Profile. Molecular Nutrition and Food Research, 2022, 66, .	1.5	13
8	New players in the relationship between diet and microbiota: the role of macromolecular antioxidant polyphenols. European Journal of Nutrition, 2021, 60, 1403-1413.	1.8	10
9	Effect of inulin-type fructans and galactooligosaccharides on cultures of <i>Lactobacillus</i> strains isolated in Algeria from camel's milk and human colostrum. Food Science and Technology International, 2021, 27, 223-233.	1.1	4
10	Impact of Extreme Obesity and Dietâ€Induced Weight Loss on the Fecal Metabolome and Gut Microbiota. Molecular Nutrition and Food Research, 2021, 65, e2000030.	1.5	19
11	<i>Akkermansia muciniphila</i> and environmental enrichment reverse cognitive impairment associated with high-fat high-cholesterol consumption in rats. Gut Microbes, 2021, 13, 1-20.	4.3	47
12	Identification of Nutritional Targets in Spanish Children Belonging to the LAyDI Cohort for the Development of Health Promotion Strategies in the First Two Years of Life. International Journal of Environmental Research and Public Health, 2021, 18, 939.	1.2	3
13	Diet and Microbiota During Pregnancy. , 2021, , .		0
14	Intestinal microbiota alterations by dietary exposure to chemicals from food cooking and processing. Application of data science for risk prediction. Computational and Structural Biotechnology Journal, 2021, 19, 1081-1091.	1.9	4
15	Diet and Microbiota in the Elderly. , 2021, , 55-55.		0
16	In vitro Selection of Probiotics for Microbiota Modulation in Normal-Weight and Severely Obese Individuals: Focus on Gas Production and Interaction With Intestinal Epithelial Cells. Frontiers in Microbiology, 2021, 12, 630572.	1.5	8
17	Longitudinal Study Depicting Differences in Complementary Feeding and Anthropometric Parameters in Late Preterm Infants up to 2 Years of Age. Nutrients, 2021, 13, 982.	1.7	1
18	Early-Life Development of the Bifidobacterial Community in the Infant Gut. International Journal of Molecular Sciences, 2021, 22, 3382.	1.8	28

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19	Influence of 2′-Fucosyllactose on the Microbiota Composition and Metabolic Activity of Fecal Cultures from Breastfed and Formula-Fed Infants at Two Months of Age. Microorganisms, 2021, 9, 1478.	1.6	15
20	Levels of Predominant Intestinal Microorganisms in 1 Month-Old Full-Term Babies and Weight Gain during the First Year of Life. Nutrients, 2021, 13, 2412.	1.7	10
21	Microbiota intestinal y salud. GastroenterologÃa Y HepatologÃa, 2021, 44, 519-535.	0.2	21
22	Gut microbes and health. GastroenterologÃa Y HepatologÃa (English Edition), 2021, 44, 519-535.	0.0	8
23	Effect of Intrapartum Antibiotics Prophylaxis on the Bifidobacterial Establishment within the Neonatal Gut. Microorganisms, 2021, 9, 1867.	1.6	8
24	Identification and Characterization of Human Observational Studies in Nutritional Epidemiology on Gut Microbiomics for Joint Data Analysis. Nutrients, 2021, 13, 3292.	1.7	6
25	Resistance of Bifidobacteria Toward Antibiotics. Methods in Molecular Biology, 2021, 2278, 195-208.	0.4	3
26	Role of Bifidobacteria on Infant Health. Microorganisms, 2021, 9, 2415.	1.6	40
27	Preterm Delivery: Microbial Dysbiosis, Gut Inflammation and Hyperpermeability. Frontiers in Microbiology, 2021, 12, 806338.	1.5	3
28	Use of Fecal Slurry Cultures to Study In Vitro Effects of Bacteriocins on the Gut Bacterial Populations of Infants. Probiotics and Antimicrobial Proteins, 2020, 12, 1218-1225.	1.9	4
29	How strong is the evidence that gut microbiota composition can be influenced by lifestyle interventions in a cardio-protective way?. Atherosclerosis, 2020, 311, 124-142.	0.4	18
30	The Relationship between Choline Bioavailability from Diet, Intestinal Microbiota Composition, and Its Modulation of Human Diseases. Nutrients, 2020, 12, 2340.	1.7	90
31	Comparison of Different Dietary Indices as Predictors of Inflammation, Oxidative Stress and Intestinal Microbiota in Middle-Aged and Elderly Subjects. Nutrients, 2020, 12, 3828.	1.7	24
32	Bifidobacterium longum subsp. infantis CECT7210 (B. infantis IM-1®) Displays In Vitro Activity against Some Intestinal Pathogens. Nutrients, 2020, 12, 3259.	1.7	13
33	Long-Term Coffee Consumption is Associated with Fecal Microbial Composition in Humans. Nutrients, 2020, 12, 1287.	1.7	53
34	An Overview on Fecal Branched Short-Chain Fatty Acids Along Human Life and as Related With Body Mass Index: Associated Dietary and Anthropometric Factors. Frontiers in Microbiology, 2020, 11, 973.	1.5	126
35	In Vitro Evaluation of Different Prebiotics on the Modulation of Gut Microbiota Composition and Function in Morbid Obese and Normal-Weight Subjects. International Journal of Molecular Sciences, 2020, 21, 906.	1.8	29
36	Microbiome: Effects of Ageing and Diet. Current Issues in Molecular Biology, 2020, 36, 33-62.	1.0	42

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37	Donated Human Milk as a Determinant Factor for the Gut Bifidobacterial Ecology in Premature Babies. Microorganisms, 2020, 8, 760.	1.6	13
38	Age-Associated Changes in Gut Microbiota and Dietary Components Related with the Immune System in Adulthood and Old Age: A Cross-Sectional Study. Nutrients, 2019, 11, 1765.	1.7	113
39	Fecal Changes Following Introduction of Milk in Infants With Outgrowing Non-IgE Cow's Milk Protein Allergy Are Influenced by Previous Consumption of the Probiotic LGG. Frontiers in Immunology, 2019, 10, 1819.	2.2	19
40	Neurobehavioral dysfunction in non-alcoholic steatohepatitis is associated with hyperammonemia, gut dysbiosis, and metabolic and functional brain regional deficits. PLoS ONE, 2019, 14, e0223019.	1.1	44
41	Intestinal Immunomodulation and Shifts on the Gut Microbiota of BALB/c Mice Promoted by Two <i>Bifidobacterium</i> and <i>Lactobacillus</i> Strains Isolated from Human Samples. BioMed Research International, 2019, 2019, 1-8.	0.9	3
42	Fermented Dairy Foods: Impact on Intestinal Microbiota and Health-Linked Biomarkers. Frontiers in Microbiology, 2019, 10, 1046.	1.5	79
43	Reply: "Letter to the editor Re: Diaz M., et al. Nutrients 2018, 10, 1481― Nutrients, 2019, 11, 476.	1.7	1
44	Nutritional composition of processed baby foods targeted at infants from 0–12 months. Journal of Food Composition and Analysis, 2019, 79, 55-62.	1.9	14
45	Impact of probiotics on development and behaviour in Drosophila melanogaster – a potential in vivo model to assess probiotics. Beneficial Microbes, 2019, 10, 179-188.	1.0	5
46	Microbiota, Food, and Health. International Journal of Molecular Sciences, 2019, 20, 6329.	1.8	4
47	In-vitro characterization of potentially probiotic Lactobacillus strains isolated from human microbiota: interaction with pathogenic bacteria and the enteric cell line HT29. Annals of Microbiology, 2019, 69, 61-72.	1.1	39
48	Exploring the interactions between serum free fatty acids and fecal microbiota in obesity through a machine learning algorithm. Food Research International, 2019, 121, 533-541.	2.9	25
49	Baby's First Microbes: The Microbiome of Human Milk. , 2019, , 3-33.		1
50	Supplementation with grape pomace in healthy women: Changes in biochemical parameters, gut microbiota and related metabolic biomarkers. Journal of Functional Foods, 2018, 45, 34-46.	1.6	29
51	Probiotics for Prevention and Treatment of Clostridium difficile Infection. Advances in Experimental Medicine and Biology, 2018, 1050, 161-176.	0.8	31
52	Transmission and persistence of IncF conjugative plasmids in the gut microbiota of full-term infants. FEMS Microbiology Ecology, 2018, 94, .	1.3	15
53	<i>Bifidobacterium breve</i> IPLA20005 affects in vitro the expression of <i>hly</i> and <i>luxS</i> genes, related to the virulence of <i>Listeria monocytogenes</i> Lm23. Canadian Journal of Microbiology, 2018, 64, 215-221.	0.8	12
54	Bioactive compounds from regular diet and faecal microbial metabolites. European Journal of Nutrition, 2018, 57, 487-497.	1.8	18

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55	Early microbiota, antibiotics and health. Cellular and Molecular Life Sciences, 2018, 75, 83-91.	2.4	76
56	The role of yogurt in food-based dietary guidelines. Nutrition Reviews, 2018, 76, 29-39.	2.6	46
57	Fecal microbiota profile in a group of myasthenia gravis patients. Scientific Reports, 2018, 8, 14384.	1.6	45
58	Microbiota and Derived Parameters in Fecal Samples of Infants with Non-IgE Cow's Milk Protein Allergy under a Restricted Diet. Nutrients, 2018, 10, 1481.	1.7	40
59	Could Fecal Phenylacetic and Phenylpropionic Acids Be Used as Indicators of Health Status?. Journal of Agricultural and Food Chemistry, 2018, 66, 10438-10446.	2.4	25
60	C-section and the Neonatal Gut Microbiome Acquisition: Consequences for Future Health. Annals of Nutrition and Metabolism, 2018, 73, 17-23.	1.0	37
61	Real-time monitoring of HT29 epithelial cells as an in vitro model for assessing functional differences among intestinal microbiotas from different human population groups. Journal of Microbiological Methods, 2018, 152, 210-216.	0.7	6
62	Selection of potential probiotic bifidobacteria and prebiotics for elderly by using in vitro faecal batch cultures. European Food Research and Technology, 2017, 243, 157-165.	1.6	17
63	Adherence to a Mediterranean Diet Influences the Fecal Metabolic Profile of Microbial-Derived Phenolics in a Spanish Cohort of Middle-Age and Older People. Journal of Agricultural and Food Chemistry, 2017, 65, 586-595.	2.4	63
64	Correlation between inÂvitro and inÂvivo assays in selection of probiotics from traditional species of bacteria. Trends in Food Science and Technology, 2017, 68, 83-90.	7.8	57
65	The First Microbial Colonizers of the Human Gut: Composition, Activities, and Health Implications of the Infant Gut Microbiota. Microbiology and Molecular Biology Reviews, 2017, 81, .	2.9	1,118
66	In vitro fermentation of different fructo-oligosaccharides by Bifidobacterium strains for the selection of synbiotic combinations. International Journal of Food Microbiology, 2017, 242, 19-23.	2.1	50
67	Nutrition and the gut microbiome in the elderly. Gut Microbes, 2017, 8, 82-97.	4.3	191
68	Probiotics, gut microbiota, and their influence on host health and disease. Molecular Nutrition and Food Research, 2017, 61, 1600240.	1.5	678
69	Intestinal Dysbiosis Is Associated with Altered Short-Chain Fatty Acids and Serum-Free Fatty Acids in Systemic Lupus Erythematosus. Frontiers in Immunology, 2017, 8, 23.	2.2	95
70	Free Fatty Acids Profiles Are Related to Gut Microbiota Signatures and Short-Chain Fatty Acids. Frontiers in Immunology, 2017, 8, 823.	2.2	75
71	Intestinal Microbiota and Weight-Gain in Preterm Neonates. Frontiers in Microbiology, 2017, 8, 183.	1.5	35
72	Shaping the Metabolism of Intestinal Bacteroides Population through Diet to Improve Human Health. Frontiers in Microbiology, 2017, 8, 376.	1.5	140

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73	Safety of Novel Microbes for Human Consumption: Practical Examples of Assessment in the European Union. Frontiers in Microbiology, 2017, 8, 1725.	1.5	125
74	Different Intestinal Microbial Profile in Over-Weight and Obese Subjects Consuming a Diet with Low Content of Fiber and Antioxidants. Nutrients, 2017, 9, 551.	1.7	36
75	Impact of intrapartum antimicrobial prophylaxis upon the intestinal microbiota and the prevalence of antibiotic resistance genes in vaginally delivered full-term neonates. Microbiome, 2017, 5, 93.	4.9	165
76	Intestinal Short Chain Fatty Acids and their Link with Diet and Human Health. Frontiers in Microbiology, 2016, 7, 185.	1.5	1,443
77	Screening of Bifidobacteria and Lactobacilli Able to Antagonize the Cytotoxic Effect of Clostridium difficile upon Intestinal Epithelial HT29 Monolayer. Frontiers in Microbiology, 2016, 7, 577.	1.5	51
78	Effect of Bifidobacterium upon Clostridium difficile Growth and Toxicity When Co-cultured in Different Prebiotic Substrates. Frontiers in Microbiology, 2016, 7, 738.	1.5	66
79	Impact of Prematurity and Perinatal Antibiotics on the Developing Intestinal Microbiota: A Functional Inference Study. International Journal of Molecular Sciences, 2016, 17, 649.	1.8	109
80	Bacteroides fragilis metabolises exopolysaccharides produced by bifidobacteria. BMC Microbiology, 2016, 16, 150.	1.3	48
81	Perinatal Microbiomes <i>'</i> Influence on Preterm Birth and Preterms' Health. Journal of Pediatric Gastroenterology and Nutrition, 2016, 63, e193-e203.	0.9	32
82	Glucolytic fingerprinting reveals metabolic groups within the genus Bifidobacterium: an exploratory study. Beneficial Microbes, 2016, 7, 265-273.	1.0	10
83	Exopolysaccharides Produced by Lactic Acid Bacteria and Bifidobacteria as Fermentable Substrates by the Intestinal Microbiota. Critical Reviews in Food Science and Nutrition, 2016, 56, 1440-1453.	5.4	139
84	Supplementation of xylitol-containing chewing gum with probiotics: a double blind, randomised pilot study focusing on saliva flow and saliva properties. Food and Function, 2016, 7, 1601-1609.	2.1	15
85	A proteomic approach towards understanding the cross talk between <i>Bacteroides fragilis</i> and <i>Bifidobacterium longum</i> in coculture. Canadian Journal of Microbiology, 2016, 62, 623-628.	0.8	8
86	Allergic Patients with Long-Term Asthma Display Low Levels of Bifidobacterium adolescentis. PLoS ONE, 2016, 11, e0147809.	1.1	90
87	Isolation, Identification and Characterisation of Potential New Probiotics. , 2015, , 3-25.		0
88	Production of immune response mediators by HT-29 intestinal cell-lines in the presence of Bifidobacterium-treated infant microbiota. Beneficial Microbes, 2015, 6, 543-552.	1.0	14
89	Intestinal Microbiota as Modulators of the Immune System. Journal of Immunology Research, 2015, 2015, 1-4.	0.9	19
90	Different metabolic features of Bacteroides fragilis growing in the presence of glucose and exopolysaccharides of bifidobacteria. Frontiers in Microbiology, 2015, 6, 825.	1.5	44

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91	Occurrence and Diversity of CRISPR-Cas Systems in the Genus Bifidobacterium. PLoS ONE, 2015, 10, e0133661.	1.1	73
92	The Effects of <i>Bifidobacterium breve</i> on Immune Mediators and Proteome of HT29 Cells Monolayers. BioMed Research International, 2015, 2015, 1-6.	0.9	21
93	Capability of exopolysaccharide-producing Lactobacillus paraplantarum BGCG11 and its non-producing isogenic strain NB1, to counteract the effect of enteropathogens upon the epithelial cell line HT29-MTX. Food Research International, 2015, 74, 199-207.	2.9	31
94	Intestinal Microbiota Development in Preterm Neonates and EffectÂofÂPerinatal Antibiotics. Journal of Pediatrics, 2015, 166, 538-544.	0.9	329
95	The establishment of the infant intestinal microbiome is not affected by rotavirus vaccination. Scientific Reports, 2015, 4, 7417.	1.6	15
96	The relationship between phenolic compounds from diet and microbiota: impact on human health. Food and Function, 2015, 6, 2424-2439.	2.1	180
97	Red Wine Consumption Is Associated with Fecal Microbiota and Malondialdehyde in a Human Population. Journal of the American College of Nutrition, 2015, 34, 135-141.	1.1	26
98	Monitoring in real time the cytotoxic effect of Clostridium difficile upon the intestinal epithelial cell line HT29. Journal of Microbiological Methods, 2015, 119, 66-73.	0.7	20
99	Enhanced butyrate formation by cross-feeding between <i>Faecalibacterium prausnitzii</i> and <i>Bifidobacterium adolescentis</i> . FEMS Microbiology Letters, 2015, 362, fnv176.	0.7	250
100	Insights from genomes of representatives of the human gut commensal <scp><i>B</i></scp> <i>isifidobacterium bifidum</i> . Environmental Microbiology, 2015, 17, 2515-2531.	1.8	80
101	Inulin-type fructans modulate intestinal Bifidobacterium species populations and decrease fecal short-chain fatty acids in obese women. Clinical Nutrition, 2015, 34, 501-507.	2.3	220
102	Effect of an α-Tocopherol-Containing Antioxidant Parenteral Emulsion upon Gut Microbiota in Preterm Infants. International Journal of Child Health and Nutrition, 2015, 4, 90-93.	0.0	1
103	Intestinal microbiota in health and disease: Role of bifidobacteria in gut homeostasis. World Journal of Gastroenterology, 2014, 20, 15163.	1.4	390
104	Intestinal Dysbiosis Associated with Systemic Lupus Erythematosus. MBio, 2014, 5, e01548-14.	1.8	500
105	Immune Modulating Capability of Two Exopolysaccharide-Producing <i>Bifidobacterium</i> Strains in a Wistar Rat Model. BioMed Research International, 2014, 2014, 1-9.	0.9	32
106	The human intestinal microbiome at extreme ages of life. Dietary intervention as a way to counteract alterations. Frontiers in Genetics, 2014, 5, 406.	1.1	124
107	Assessment of stress tolerance acquisition in the heat-tolerant derivative strains of <i>Bifidobacterium animalis</i> subsp. <i>lactis </i> BB-12 and <i>Lactobacillus rhamnosus</i> GG. Journal of Applied Microbiology, 2014, 117, 239-248.	1.4	18
108	Pilot Study of Diet and Microbiota: Interactive Associations of Fibers and Polyphenols with Human Intestinal Bacteria. Journal of Agricultural and Food Chemistry, 2014, 62, 5330-5336.	2.4	75

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109	Effect of bacteria used in food industry on the proliferation and cytokine production of epithelial intestinal cellular lines. Journal of Functional Foods, 2014, 6, 348-355.	1.6	11
110	<i>In vitro</i> evaluation of the impact of human background microbiota on the response to <i>Bifidobacterium</i> strains and fructo-oligosaccharides. British Journal of Nutrition, 2013, 110, 2030-2036.	1.2	25
111	Interactions between Bifidobacterium and Bacteroides Species in Cofermentations Are Affected by Carbon Sources, Including Exopolysaccharides Produced by Bifidobacteria. Applied and Environmental Microbiology, 2013, 79, 7518-7524.	1.4	82
112	Omics for the study of probiotic microorganisms. Food Research International, 2013, 54, 1061-1071.	2.9	30
113	Catabolism of Glucose and Lactose in Bifidobacterium animalis subsp. lactis, Studied by <sup>13</sup> C Nuclear Magnetic Resonance. Applied and Environmental Microbiology, 2013, 79, 7628-7638.	1.4	44
114	Fiber from a regular diet is directly associated with fecal short-chain fatty acid concentrations in the elderly. Nutrition Research, 2013, 33, 811-816.	1.3	70
115	Adaptation of bifidobacteria to the gastrointestinal tract and functional consequences. Pharmacological Research, 2013, 69, 127-136.	3.1	48
116	Factors involved in the colonization and survival of bifidobacteria in the gastrointestinal tract. FEMS Microbiology Letters, 2013, 340, 1-10.	0.7	68
117	Assessment of intestinal microbiota modulation ability of Bifidobacterium strains in inÂvitro fecal batch cultures from preterm neonates. Anaerobe, 2013, 19, 9-16.	1.0	45
118	Assessment of the effect of stress-tolerance acquisition on some basic characteristics of specific probiotics. International Journal of Food Microbiology, 2013, 165, 51-56.	2.1	30
119	Microbiota of the Intestine: Probiotics. , 2013, , 175-181.		1
120	Antibiotic resistance in probiotic bacteria. Frontiers in Microbiology, 2013, 4, 202.	1.5	417
121	Microbial Targets for the Development of Functional Foods Accordingly with Nutritional and Immune Parameters Altered in the Elderly. Journal of the American College of Nutrition, 2013, 32, 399-406.	1.1	65
122	Insights into the Ropy Phenotype of the Exopolysaccharide-Producing Strain Bifidobacterium animalis subsp. <i>lactis</i> A1dOxR. Applied and Environmental Microbiology, 2013, 79, 3870-3874.	1.4	19
123	Population Dynamics of Some Relevant Intestinal Microbial Groups in Human Fecal Batch Cultures with Added Fermentable Xylooligosaccharides Obtained from Rice Husks. BioResources, 2013, 8, .	0.5	5
124	Assessing the Fecal Microbiota: An Optimized Ion Torrent 16S rRNA Gene-Based Analysis Protocol. PLoS ONE, 2013, 8, e68739.	1.1	257
125	Fatty acids intake and immune parameters in the elderly. Nutricion Hospitalaria, 2013, 28, 474-8.	0.2	8
126	Enhancing probiotic stability in industrial processes. Microbial Ecology in Health and Disease, 2012, 23,	3.8	22

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127	Controlled Gene Expression in Bifidobacteria by Use of a Bile-Responsive Element. Applied and Environmental Microbiology, 2012, 78, 581-585.	1.4	17
128	Genome Sequence of the Antarctic Psychrophile Bacterium Planococcus antarcticus DSM 14505. Journal of Bacteriology, 2012, 194, 4465-4465.	1.0	16
129	Molecular Clues To Understand the Aerotolerance Phenotype of Bifidobacterium animalis subsp. lactis. Applied and Environmental Microbiology, 2012, 78, 644-650.	1.4	39
130	Genome Sequence of Parascardovia denticolens IPLA 20019, Isolated from Human Breast Milk. Journal of Bacteriology, 2012, 194, 4776-4777.	1.0	9
131	Role of Extracellular Transaldolase from Bifidobacterium bifidum in Mucin Adhesion and Aggregation. Applied and Environmental Microbiology, 2012, 78, 3992-3998.	1.4	109
132	Facultative to strict anaerobes ratio in the preterm infant microbiota. Gut Microbes, 2012, 3, 583-588.	4.3	73
133	Interaction of Bifidobacterium bifidum LMG13195 with HT29 Cells Influences Regulatory-T-Cell-Associated Chemokine Receptor Expression. Applied and Environmental Microbiology, 2012, 78, 2850-2857.	1.4	52
134	Genome Sequence of the Immunomodulatory Strain Bifidobacterium bifidum LMG 13195. Journal of Bacteriology, 2012, 194, 6997-6997.	1.0	3
135	Immune Modulation Capability of Exopolysaccharides Synthesised by Lactic Acid Bacteria and Bifidobacteria. Probiotics and Antimicrobial Proteins, 2012, 4, 227-237.	1.9	156
136	Treg-inducing membrane vesicles from Bifidobacterium bifidum LMG13195 as potential adjuvants in immunotherapy. Vaccine, 2012, 30, 825-829.	1.7	69
137	Metagenomics and probiotics. Clinical Microbiology and Infection, 2012, 18, 32-34.	2.8	46
138	Exopolysaccharide-producing Bifidobacterium strains elicit different in vitro responses upon interaction with human cells. Food Research International, 2012, 46, 99-107.	2.9	102
139	Diversity of Bifidobacteria within the Infant Gut Microbiota. PLoS ONE, 2012, 7, e36957.	1.1	512
140	Toward improving technological and functional properties of probiotics in foods. Trends in Food Science and Technology, 2012, 26, 56-63.	7.8	44
141	Development of probiotic products for nutritional requirements of specific human populations. Engineering in Life Sciences, 2012, 12, 368-376.	2.0	16
142	Establishment and development of intestinal microbiota in preterm neonates. FEMS Microbiology Ecology, 2012, 79, 763-772.	1.3	365
143	Deep 16S rRNA metagenomics and quantitative PCR analyses of the premature infant fecal microbiota. Anaerobe, 2012, 18, 378-380.	1.0	60
144	Manufacturing process influences properties of probiotic bacteria. British Journal of Nutrition, 2011, 105, 887-894.	1.2	101

9

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145	Adhesion of bile-adapted Bifidobacterium strains to the HT29-MTX cell line is modified after sequential gastrointestinal challenge simulated in vitro using human gastric and duodenal juices. Research in Microbiology, 2011, 162, 514-519.	1.0	40
146	Immune Response to Bifidobacterium bifidum Strains Support Treg/Th17 Plasticity. PLoS ONE, 2011, 6, e24776.	1.1	120
147	Current and Future Applications of Probiotics. Current Nutrition and Food Science, 2011, 7, 170-180.	0.3	3
148	Evaluation of the ability of Bifidobacterium longum to metabolize human intestinal mucus. FEMS Microbiology Letters, 2011, 314, 125-130.	0.7	24
149	Assessment of intestinal microbiota of full-term breast-fed infants from two different geographical locations. Early Human Development, 2011, 87, 511-513.	0.8	47
150	Safety and intestinal microbiota modulation by the exopolysaccharide-producing strains Bifidobacterium animalis IPLA R1 and Bifidobacterium longum IPLA E44 orally administered to Wistar rats. International Journal of Food Microbiology, 2011, 144, 342-351.	2.1	66
151	Characterization and in vitro properties of potentially probiotic Bifidobacterium strains isolated from breast-milk. International Journal of Food Microbiology, 2011, 149, 28-36.	2.1	109
152	3rd International Symposium on Propionibacteria and Bifidobacteria: Dairy and Probiotic Applications, Oviedo 1–4 June 2010. International Journal of Food Microbiology, 2011, 149, 1.	2.1	0
153	How do bifidobacteria counteract environmental challenges? Mechanisms involved and physiological consequences. Genes and Nutrition, 2011, 6, 307-318.	1.2	94
154	Establishment and development of lactic acid bacteria and bifidobacteria microbiota in breast-milk and the infant gut. Anaerobe, 2010, 16, 307-310.	1.0	271
155	Distinct Bifidobacterium strains drive different immune responses in vitro. International Journal of Food Microbiology, 2010, 138, 157-165.	2.1	141
156	Probiotics in Adhesion of Pathogens. , 2010, , 353-370.		28
157	Fecal <i>Bifidobacterium</i> Levels in Elderly Nursing Home Patients. Bioscience and Microflora, 2010, 29, 111-113.	0.5	14
158	Genetic Basis of Tetracycline Resistance in <i>Bifidobacterium animalis</i> subsp. <i>lactis</i> . Applied and Environmental Microbiology, 2010, 76, 3364-3369.	1.4	61
159	Bile Affects the Synthesis of Exopolysaccharides by <i>Bifidobacterium animalis</i> . Applied and Environmental Microbiology, 2009, 75, 1204-1207.	1.4	100
160	Microbial–Host Interactions: Selecting the Right Probiotics and Prebiotics for Infants. Nestle Nutrition Workshop Series Paediatric Programme, 2009, 64, 201-217.	1.5	19
161	Bile-Inducible Efflux Transporter from <i>Bifidobacterium longum</i> NCC2705, Conferring Bile Resistance. Applied and Environmental Microbiology, 2009, 75, 3153-3160.	1.4	66
162	Coculture of Bifidobacterium longum and Bifidobacterium breve alters their protein expression profiles and enzymatic activities. International Journal of Food Microbiology, 2009, 133, 148-153.	2.1	37

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163	Microbiomic analysis of the bifidobacterial population in the human distal gut. ISME Journal, 2009, 3, 745-751.	4.4	128
164	Probiotic fermented milks: Present and future. International Journal of Dairy Technology, 2009, 62, 472-483.	1.3	57
165	Effect of clinical and probiotic Lactobacillus rhamnosus strains on intestinal permeability and bacterial translocation in healthy and colitic rats. Food Research International, 2009, 42, 636-640.	2.9	8
166	Inflammation Markers and Malnutrition as Risk Factors for Infections and Impaired Health-Related Quality of Life Among Older Nursing Home Residents. Journal of the American Medical Directors Association, 2009, 10, 348-353.	1.2	39
167	Safety Assessment of Probiotics. , 2009, , 1193-1235.		14
168	<i>Bifidobacterium</i> microbiota and parameters of immune function in elderly subjects. FEMS Immunology and Medical Microbiology, 2008, 53, 18-25.	2.7	141
169	Degradation of 16S rRNA and attributes of viability of viable but nonculturable probiotic bacteria. Letters in Applied Microbiology, 2008, 46, 693-698.	1.0	83
170	Two Different Tetracycline Resistance Mechanisms, Plasmid-Carried <i>tet</i> (L) and Chromosomally Located Transposon-Associated <i>tet</i> (M), Coexist in <i>Lactobacillus sakei</i> Rits 9. Applied and Environmental Microbiology, 2008, 74, 1394-1401.	1.4	75
171	Mucin Degradation by <i>Bifidobacterium</i> Strains Isolated from the Human Intestinal Microbiota. Applied and Environmental Microbiology, 2008, 74, 1936-1940.	1.4	180
172	Exopolysaccharides Produced by Intestinal <i>Bifidobacterium</i> Strains Act as Fermentable Substrates for Human Intestinal Bacteria. Applied and Environmental Microbiology, 2008, 74, 4737-4745.	1.4	197
173	Breast Milk: A Source of Bifidobacteria for Infant Gut Development and Maturation?. Neonatology, 2007, 92, 64-66.	0.9	235
174	Qualitative and quantitative analyses of the bifidobacterial microbiota in the colonic mucosa of patients with colorectal cancer, diverticulitis and inflammatory bowel disease. World Journal of Gastroenterology, 2007, 13, 3985.	1.4	102
175	Maternal breastâ€milk and intestinal bifidobacteria guide the compositional development of the <i>Bifidobacterium</i> microbiota in infants at risk of allergic disease. Clinical and Experimental Allergy, 2007, 37, 1764-1772.	1.4	252
176	Competitive exclusion of enteropathogens from human intestinal mucus by Bifidobacterium strains with acquired resistance to bile — A preliminary study. International Journal of Food Microbiology, 2007, 113, 228-232.	2.1	71
177	Induction of α-l-arabinofuranosidase activity by monomeric carbohydrates in Bifidobacterium longum and ubiquity of encoding genes. Archives of Microbiology, 2007, 187, 145-153.	1.0	24
178	Short Communication: Effect of Exopolysaccharide Isolated from "Viili―on the Adhesion of Probiotics and Pathogens to Intestinal Mucus. Journal of Dairy Science, 2006, 89, 2355-2358.	1.4	52
179	New methods for selecting and evaluating probiotics. Digestive and Liver Disease, 2006, 38, S242-S247.	0.4	105
180	Adhesion and competitive inhibition and displacement of human enteropathogens by selected lactobacilli. Food Research International, 2006, 39, 467-471.	2.9	114

#	Article	IF	CITATIONS
181	Exopolysaccharides Produced by Probiotic Strains Modify the Adhesion of Probiotics and Enteropathogens to Human Intestinal Mucus. Journal of Food Protection, 2006, 69, 2011-2015.	0.8	201
182	Adhesion Properties and Competitive Pathogen Exclusion Ability of Bifidobacteria with Acquired Acid Resistance. Journal of Food Protection, 2006, 69, 1675-1679.	0.8	72
183	Probiotic Intervention in Neonates-Will Permanent Colonization Ensue?. Journal of Pediatric Gastroenterology and Nutrition, 2006, 42, 604-606.	0.9	54
184	Presence of specific antibiotic (tet) resistance genes in infant faecal microbiota. FEMS Immunology and Medical Microbiology, 2006, 48, 21-25.	2.7	53
185	Quantitative assessment of faecal bifidobacterial populations by real-time PCR using lanthanide probes. Journal of Applied Microbiology, 2006, 102, 061120055200067-???.	1.4	29
186	Assuring the continued safety of lactic acid bacteria used as probiotics. Biologia (Poland), 2006, 61, 755-760.	0.8	13
187	Comparison of four methods to enumerate probiotic bifidobacteria in a fermented food product. Food Microbiology, 2006, 23, 571-577.	2.1	93
188	Meat as a First Complementary Food for Breastfed Infants: Feasibility and Impact on Zinc Intake and Status. Journal of Pediatric Gastroenterology and Nutrition, 2006, 42, 207-214.	0.9	265
189	Gut Microbiota in Infants between 6 and 24 Months of Age. , 2005, 56, 43-56.		13
190	Probiotics That Modify Disease Risk. Journal of Nutrition, 2005, 135, 1294-1298.	1.3	155
191	Similar bifidogenic effects of prebiotic-supplemented partially hydrolyzed infant formula and breastfeeding on infant gut microbiota. FEMS Immunology and Medical Microbiology, 2005, 43, 59-65.	2.7	136
192	Ability of Bifidobacterium strains with acquired resistance to bile to adhere to human intestinal mucus. International Journal of Food Microbiology, 2005, 101, 341-346.	2.1	60
193	Adhesion of Selected Bifidobacterium Strains to Human Intestinal Mucus and the Role of Adhesion in Enteropathogen Exclusion. Journal of Food Protection, 2005, 68, 2672-2678.	0.8	173
194	Probiotic Bacteria May Become Dormant during Storage. Applied and Environmental Microbiology, 2005, 71, 1662-1663.	1.4	81
195	Removal of the cyanobacterial toxin microcystin-LR by human probiotics. Toxicon, 2005, 46, 111-114.	0.8	41
196	The genomics of probiotic intestinal microorganisms. Genome Biology, 2005, 6, 225.	13.9	19
197	MICROBIOTA OF THE INTESTINE   Probiotics. , 2005, , 244-251.		1
198	Safety of probiotics. Scandinavian Journal of Nutrition, 2004, 48, 42-48.	0.2	34

#	Article	IF	CITATIONS
199	New Real-Time Quantitative PCR Procedure for Quantification of Bifidobacteria in Human Fecal Samples. Applied and Environmental Microbiology, 2004, 70, 4165-4169.	1.4	154
200	Effect of the adaptation to high bile salts concentrations on glycosidic activity, survival at low PH and cross-resistance to bile salts in Bifidobacterium. International Journal of Food Microbiology, 2004, 94, 79-86.	2.1	125
201	Viability and diversity of probiotic Lactobacillus and Bifidobacterium populations included in commercial fermented milks. Food Research International, 2004, 37, 839-850.	2.9	192
202	Human Studies on Probiotics: What Is Scientifically Proven. Journal of Food Science, 2004, 69, M137.	1.5	29
203	Inhibition of Bacillus cereus growth in carbonated fermented bifidus milk. Food Microbiology, 2003, 20, 519-526.	2.1	13
204	Characterisation of a Bifidobacterium strain with acquired resistance to cholate—A preliminary study. International Journal of Food Microbiology, 2003, 82, 191-198.	2.1	66
205	Quality of plain yoghurt made from refrigerated and CO2-treated milk. Food Research International, 2003, 36, 43-48.	2.9	32
206	Proteolysis in rennet-coagulated Spanish hard cheeses made from milk preserved by refrigeration and addition of carbon dioxide. Journal of Dairy Research, 2003, 70, 115-122.	0.7	4
207	Evolution of carbohydrate fraction in carbonated fermented milks as affected by β-galactosidase activity of starter strains. Journal of Dairy Research, 2002, 69, 125-137.	0.7	10
208	Characteristics of carbonated fermented milk and survival of probiotic bacteria. International Dairy Journal, 2000, 10, 213-220.	1.5	47