Miguel Gueimonde

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

200 12,228 58 106 g-index

214 14,890 5 6.47 ext. papers ext. citations avg, IF L-index

#	Paper	IF	Citations
200	Needle in a Whey-Stack: PhRACS as a Discovery Tool for Unknown Phage-Host Combinations <i>MBio</i> , 2022 , e0333421	7.8	O
199	Gut Microbiome Characteristics in feral and domesticated horses from different geographic locations <i>Communications Biology</i> , 2022 , 5, 172	6.7	2
198	Preterm Delivery: Microbial Dysbiosis, Gut Inflammation and Hyperpermeability <i>Frontiers in Microbiology</i> , 2021 , 12, 806338	5.7	1
197	Daily ingestion of Akkermansia mucciniphila for one month promotes healthy aging and increases lifespan in old female mice. <i>Biogerontology</i> , 2021 , 1	4.5	4
196	Role of Bifidobacteria on Infant Health <i>Microorganisms</i> , 2021 , 9,	4.9	3
195	Early-Life Development of the Bifidobacterial Community in the Infant Gut. <i>International Journal of Molecular Sciences</i> , 2021 , 22,	6.3	11
194	New players in the relationship between diet and microbiota: the role of macromolecular antioxidant polyphenols. <i>European Journal of Nutrition</i> , 2021 , 60, 1403-1413	5.2	3
193	Effect of inulin-type fructans and galactooligosaccharides on cultures of strains isolated in Algeria from camel's milk and human colostrum. <i>Food Science and Technology International</i> , 2021 , 27, 223-233	2.6	0
192	Impact of Extreme Obesity and Diet-Induced Weight Loss on the Fecal Metabolome and Gut Microbiota. <i>Molecular Nutrition and Food Research</i> , 2021 , 65, e2000030	5.9	8
191	and environmental enrichment reverse cognitive impairment associated with high-fat high-cholesterol consumption in rats. <i>Gut Microbes</i> , 2021 , 13, 1-20	8.8	11
190	Intestinal microbiota alterations by dietary exposure to chemicals from food cooking and processing. Application of data science for risk prediction. <i>Computational and Structural Biotechnology Journal</i> , 2021 , 19, 1081-1091	6.8	3
189	Diet and Microbiota in the Elderly 2021 , 55-55		
188	Selection of Probiotics for Microbiota Modulation in Normal-Weight and Severely Obese Individuals: Focus on Gas Production and Interaction With Intestinal Epithelial Cells. <i>Frontiers in Microbiology</i> , 2021 , 12, 630572	5.7	1
187	Levels of Predominant Intestinal Microorganisms in 1 Month-Old Full-Term Babies and Weight Gain during the First Year of Life. <i>Nutrients</i> , 2021 , 13,	6.7	2
186	Gut microbes and health. <i>Gastroenterolog</i> Y Hepatolog A, 2021, 44, 519-535	0.9	1
185	Gut microbes and health. Gastroenterologa Y Hepatologa (English Edition), 2021, 44, 519-535	0.1	0
184	Effect of Intrapartum Antibiotics Prophylaxis on the Bifidobacterial Establishment within the Neonatal Gut. <i>Microorganisms</i> , 2021 , 9,	4.9	2

183	Resistance of Bifidobacteria Toward Antibiotics. <i>Methods in Molecular Biology</i> , 2021 , 2278, 195-208	1.4	1	
182	subsp. CECT7210 (IM-1) Displays In Vitro Activity against Some Intestinal Pathogens. <i>Nutrients</i> , 2020 , 12,	6.7	7	
181	Long-Term Coffee Consumption is Associated with Fecal Microbial Composition in Humans. <i>Nutrients</i> , 2020 , 12,	6.7	25	
180	An Overview on Fecal Branched Short-Chain Fatty Acids Along Human Life and as Related With Body Mass Index: Associated Dietary and Anthropometric Factors. <i>Frontiers in Microbiology</i> , 2020 , 11, 973	5.7	50	
179	In Vitro Evaluation of Different Prebiotics on the Modulation of Gut Microbiota Composition and Function in Morbid Obese and Normal-Weight Subjects. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	16	
178	Microbiome: Effects of Ageing and Diet. Current Issues in Molecular Biology, 2020, 36, 33-62	2.9	20	
177	Donated Human Milk as a Determinant Factor for the Gut Bifidobacterial Ecology in Premature Babies. <i>Microorganisms</i> , 2020 , 8,	4.9	6	
176	Use of Fecal Slurry Cultures to Study In Vitro Effects of Bacteriocins on the Gut Bacterial Populations of Infants. <i>Probiotics and Antimicrobial Proteins</i> , 2020 , 12, 1218-1225	5.5	1	
175	How strong is the evidence that gut microbiota composition can be influenced by lifestyle interventions in a cardio-protective way?. <i>Atherosclerosis</i> , 2020 , 311, 124-142	3.1	7	
174	The Relationship between Choline Bioavailability from Diet, Intestinal Microbiota Composition, and Its Modulation of Human Diseases. <i>Nutrients</i> , 2020 , 12,	6.7	25	
173	Comparison of Different Dietary Indices as Predictors of Inflammation, Oxidative Stress and Intestinal Microbiota in Middle-Aged and Elderly Subjects. <i>Nutrients</i> , 2020 , 12,	6.7	9	
172	Neurobehavioral dysfunction in non-alcoholic steatohepatitis is associated with hyperammonemia, gut dysbiosis, and metabolic and functional brain regional deficits. <i>PLoS ONE</i> , 2019 , 14, e0223019	3.7	21	
171	Intestinal Immunomodulation and Shifts on the Gut Microbiota of BALB/c Mice Promoted by Two and Strains Isolated from Human Samples. <i>BioMed Research International</i> , 2019 , 2019, 2323540	3	1	
170	Fermented Dairy Foods: Impact on Intestinal Microbiota and Health-Linked Biomarkers. <i>Frontiers in Microbiology</i> , 2019 , 10, 1046	5.7	41	
169	Reply: "Letter to the editor Re: Diaz M., et al. 2018, , 1481". Nutrients, 2019, 11,	6.7		
168	Nutritional composition of processed baby foods targeted at infants from 012 months. <i>Journal of Food Composition and Analysis</i> , 2019 , 79, 55-62	4.1	4	
167	Age-Associated Changes in Gut Microbiota and Dietary Components Related with the Immune System in Adulthood and Old Age: A Cross-Sectional Study. <i>Nutrients</i> , 2019 , 11,	6.7	55	
166	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. <i>Frontiers in Immunology</i> , 2019 , 10, 1819	8.4	9	

165	Baby First Microbes: The Microbiome of Human Milk 2019 , 3-33		1
164	Impact of probiotics on development and behaviour in Drosophila melanogaster - a potential in vivo model to assess probiotics. <i>Beneficial Microbes</i> , 2019 , 10, 179-188	4.9	3
163	In-vitro characterization of potentially probiotic Lactobacillus strains isolated from human microbiota: interaction with pathogenic bacteria and the enteric cell line HT29. <i>Annals of Microbiology</i> , 2019 , 69, 61-72	3.2	20
162	Exploring the interactions between serum free fatty acids and fecal microbiota in obesity through a machine learning algorithm. <i>Food Research International</i> , 2019 , 121, 533-541	7	15
161	Supplementation with grape pomace in healthy women: Changes in biochemical parameters, gut microbiota and related metabolic biomarkers. <i>Journal of Functional Foods</i> , 2018 , 45, 34-46	5.1	21
160	Probiotics for Prevention and Treatment of Clostridium difficile Infection. <i>Advances in Experimental Medicine and Biology</i> , 2018 , 1050, 161-176	3.6	17
159	Transmission and persistence of IncF conjugative plasmids in the gut microbiota of full-term infants. <i>FEMS Microbiology Ecology</i> , 2018 , 94,	4.3	8
158	Bifidobacterium breve IPLA20005 affects in vitro the expression of hly and luxS genes, related to the virulence of Listeria monocytogenes Lm23. <i>Canadian Journal of Microbiology</i> , 2018 , 64, 215-221	3.2	10
157	Bioactive compounds from regular diet and faecal microbial metabolites. <i>European Journal of Nutrition</i> , 2018 , 57, 487-497	5.2	11
156	Early microbiota, antibiotics and health. Cellular and Molecular Life Sciences, 2018, 75, 83-91	10.3	54
155	C-section and the Neonatal Gut Microbiome Acquisition: Consequences for Future Health. <i>Annals of Nutrition and Metabolism</i> , 2018 , 73 Suppl 3, 17-23	4.5	22
154			
<i>J</i> I	Real-time monitoring of HT29 epithelial cells as an in vitro model for assessing functional differences among intestinal microbiotas from different human population groups. <i>Journal of Microbiological Methods</i> , 2018 , 152, 210-216	2.8	5
153	differences among intestinal microbiotas from different human population groups. <i>Journal of</i>	2.8	25
	differences among intestinal microbiotas from different human population groups. <i>Journal of Microbiological Methods</i> , 2018 , 152, 210-216		
153	differences among intestinal microbiotas from different human population groups. <i>Journal of Microbiological Methods</i> , 2018 , 152, 210-216 The role of yogurt in food-based dietary guidelines. <i>Nutrition Reviews</i> , 2018 , 76, 29-39	6.4	25
153 152	differences among intestinal microbiotas from different human population groups. <i>Journal of Microbiological Methods</i> , 2018 , 152, 210-216 The role of yogurt in food-based dietary guidelines. <i>Nutrition Reviews</i> , 2018 , 76, 29-39 Fecal microbiota profile in a group of myasthenia gravis patients. <i>Scientific Reports</i> , 2018 , 8, 14384 Microbiota and Derived Parameters in Fecal Samples of Infants with Non-IgE Cow's Milk Protein	6.4 4.9	25
153 152 151	differences among intestinal microbiotas from different human population groups. <i>Journal of Microbiological Methods</i> , 2018 , 152, 210-216 The role of yogurt in food-based dietary guidelines. <i>Nutrition Reviews</i> , 2018 , 76, 29-39 Fecal microbiota profile in a group of myasthenia gravis patients. <i>Scientific Reports</i> , 2018 , 8, 14384 Microbiota and Derived Parameters in Fecal Samples of Infants with Non-IgE Cow's Milk Protein Allergy under a Restricted Diet. <i>Nutrients</i> , 2018 , 10, Could Fecal Phenylacetic and Phenylpropionic Acids Be Used as Indicators of Health Status?.	6.4 4.9 6.7	25 19 26

(2016-2017)

147	Different Intestinal Microbial Profile in Over-Weight and Obese Subjects Consuming a Diet with Low Content of Fiber and Antioxidants. <i>Nutrients</i> , 2017 , 9,	6.7	28
146	Impact of intrapartum antimicrobial prophylaxis upon the intestinal microbiota and the prevalence of antibiotic resistance genes in vaginally delivered full-term neonates. <i>Microbiome</i> , 2017 , 5, 93	16.6	110
145	Correlation between in vitro and in vivo assays in selection of probiotics from traditional species of bacteria. <i>Trends in Food Science and Technology</i> , 2017 , 68, 83-90	15.3	43
144	The First Microbial Colonizers of the Human Gut: Composition, Activities, and Health Implications of the Infant Gut Microbiota. <i>Microbiology and Molecular Biology Reviews</i> , 2017 , 81,	13.2	626
143	In vitro fermentation of different fructo-oligosaccharides by Bifidobacterium strains for the selection of synbiotic combinations. <i>International Journal of Food Microbiology</i> , 2017 , 242, 19-23	5.8	38
142	Nutrition and the gut microbiome in the elderly. <i>Gut Microbes</i> , 2017 , 8, 82-97	8.8	121
141	Probiotics, gut microbiota, and their influence on host health and disease. <i>Molecular Nutrition and Food Research</i> , 2017 , 61, 1600240	5.9	442
140	Intestinal Dysbiosis Is Associated with Altered Short-Chain Fatty Acids and Serum-Free Fatty Acids in Systemic Lupus Erythematosus. <i>Frontiers in Immunology</i> , 2017 , 8, 23	8.4	53
139	Free Fatty Acids Profiles Are Related to Gut Microbiota Signatures and Short-Chain Fatty Acids. <i>Frontiers in Immunology</i> , 2017 , 8, 823	8.4	45
138	Intestinal Microbiota and Weight-Gain in Preterm Neonates. Frontiers in Microbiology, 2017, 8, 183	5.7	23
137	Shaping the Metabolism of Intestinal Population through Diet to Improve Human Health. <i>Frontiers in Microbiology</i> , 2017 , 8, 376	5.7	93
136	Safety of Novel Microbes for Human Consumption: Practical Examples of Assessment in the European Union. <i>Frontiers in Microbiology</i> , 2017 , 8, 1725	5.7	84
135	Glucolytic fingerprinting reveals metabolic groups within the genus Bifidobacterium: an exploratory study. <i>Beneficial Microbes</i> , 2016 , 7, 265-73	4.9	6
134	Exopolysaccharides Produced by Lactic Acid Bacteria and Bifidobacteria as Fermentable Substrates by the Intestinal Microbiota. <i>Critical Reviews in Food Science and Nutrition</i> , 2016 , 56, 1440-53	11.5	97
133	Supplementation of xylitol-containing chewing gum with probiotics: a double blind, randomised pilot study focusing on saliva flow and saliva properties. <i>Food and Function</i> , 2016 , 7, 1601-9	6.1	8
132	A proteomic approach towards understanding the cross talk between Bacteroides fragilis and Bifidobacterium longum in coculture. <i>Canadian Journal of Microbiology</i> , 2016 , 62, 623-8	3.2	5
131	Allergic Patients with Long-Term Asthma Display Low Levels of Bifidobacterium adolescentis. <i>PLoS ONE</i> , 2016 , 11, e0147809	3.7	62
130	Intestinal Short Chain Fatty Acids and their Link with Diet and Human Health. <i>Frontiers in Microbiology</i> , 2016 , 7, 185	5.7	934

129	Screening of Bifidobacteria and Lactobacilli Able to Antagonize the Cytotoxic Effect of Clostridium difficile upon Intestinal Epithelial HT29 Monolayer. <i>Frontiers in Microbiology</i> , 2016 , 7, 577	5.7	31
128	Effect of Bifidobacterium upon Clostridium difficile Growth and Toxicity When Co-cultured in Different Prebiotic Substrates. <i>Frontiers in Microbiology</i> , 2016 , 7, 738	5.7	45
127	Impact of Prematurity and Perinatal Antibiotics on the Developing Intestinal Microbiota: A Functional Inference Study. <i>International Journal of Molecular Sciences</i> , 2016 , 17,	6.3	81
126	Bacteroides fragilis metabolises exopolysaccharides produced by bifidobacteria. <i>BMC Microbiology</i> , 2016 , 16, 150	4.5	29
125	Perinatal Microbiomes' Influence on Preterm Birth and Preterms' Health: Influencing Factors and Modulation Strategies. <i>Journal of Pediatric Gastroenterology and Nutrition</i> , 2016 , 63, e193-e203	2.8	19
124	Intestinal microbiota development in preterm neonates and effect of perinatal antibiotics. <i>Journal of Pediatrics</i> , 2015 , 166, 538-44	3.6	250
123	The relationship between phenolic compounds from diet and microbiota: impact on human health. <i>Food and Function</i> , 2015 , 6, 2424-39	6.1	140
122	Red wine consumption is associated with fecal microbiota and malondialdehyde in a human population. <i>Journal of the American College of Nutrition</i> , 2015 , 34, 135-41	3.5	24
121	Monitoring in real time the cytotoxic effect of Clostridium difficile upon the intestinal epithelial cell line HT29. <i>Journal of Microbiological Methods</i> , 2015 , 119, 66-73	2.8	15
120	Enhanced butyrate formation by cross-feeding between Faecalibacterium prausnitzii and Bifidobacterium adolescentis. <i>FEMS Microbiology Letters</i> , 2015 , 362,	2.9	167
119	Insights from genomes of representatives of the human gut commensal Bifidobacterium bifidum. <i>Environmental Microbiology</i> , 2015 , 17, 2515-31	5.2	61
118	Inulin-type fructans modulate intestinal Bifidobacterium species populations and decrease fecal short-chain fatty acids in obese women. <i>Clinical Nutrition</i> , 2015 , 34, 501-7	5.9	162
117	Isolation, Identification and Characterisation of Potential New Probiotics 2015, 3-25		
116	Production of immune response mediators by HT-29 intestinal cell-lines in the presence of Bifidobacterium-treated infant microbiota. <i>Beneficial Microbes</i> , 2015 , 6, 543-52	4.9	12
115	Different metabolic features of Bacteroides fragilis growing in the presence of glucose and exopolysaccharides of bifidobacteria. <i>Frontiers in Microbiology</i> , 2015 , 6, 825	5.7	32
114	Occurrence and Diversity of CRISPR-Cas Systems in the Genus Bifidobacterium. <i>PLoS ONE</i> , 2015 , 10, e0	13366	1 53
113	The effects of Bifidobacterium breve on immune mediators and proteome of HT29 cells monolayers. <i>BioMed Research International</i> , 2015 , 2015, 479140	3	19
112	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. <i>Food Research International</i> , 2015 , 74, 199-207	7	22

(2013-2014)

111	The establishment of the infant intestinal microbiome is not affected by rotavirus vaccination. <i>Scientific Reports</i> , 2014 , 4, 7417	4.9	15
110	Assessment of stress tolerance acquisition in the heat-tolerant derivative strains of Bifidobacterium animalis subsp. lactis BB-12 and Lactobacillus rhamnosus GG. <i>Journal of Applied Microbiology</i> , 2014 , 117, 239-48	4.7	15
109	Pilot study of diet and microbiota: interactive associations of fibers and polyphenols with human intestinal bacteria. <i>Journal of Agricultural and Food Chemistry</i> , 2014 , 62, 5330-6	5.7	62
108	Effect of bacteria used in food industry on the proliferation and cytokine production of epithelial intestinal cellular lines. <i>Journal of Functional Foods</i> , 2014 , 6, 348-355	5.1	10
107	Intestinal microbiota in health and disease: role of bifidobacteria in gut homeostasis. <i>World Journal of Gastroenterology</i> , 2014 , 20, 15163-76	5.6	282
106	Intestinal dysbiosis associated with systemic lupus erythematosus. <i>MBio</i> , 2014 , 5, e01548-14	7.8	309
105	Immune modulating capability of two exopolysaccharide-producing Bifidobacterium strains in a Wistar rat model. <i>BioMed Research International</i> , 2014 , 2014, 106290	3	23
104	The human intestinal microbiome at extreme ages of life. Dietary intervention as a way to counteract alterations. <i>Frontiers in Genetics</i> , 2014 , 5, 406	4.5	96
103	In vitro evaluation of the impact of human background microbiota on the response to Bifidobacterium strains and fructo-oligosaccharides. <i>British Journal of Nutrition</i> , 2013 , 110, 2030-6	3.6	20
102	Interactions between Bifidobacterium and Bacteroides species in cofermentations are affected by carbon sources, including exopolysaccharides produced by bifidobacteria. <i>Applied and Environmental Microbiology</i> , 2013 , 79, 7518-24	4.8	66
101	Omics for the study of probiotic microorganisms. Food Research International, 2013, 54, 1061-1071	7	26
100	Catabolism of glucose and lactose in Bifidobacterium animalis subsp. lactis, studied by 13C Nuclear Magnetic Resonance. <i>Applied and Environmental Microbiology</i> , 2013 , 79, 7628-38	4.8	29
99	Fiber from a regular diet is directly associated with fecal short-chain fatty acid concentrations in the elderly. <i>Nutrition Research</i> , 2013 , 33, 811-6	4	54
98	Adaptation of bifidobacteria to the gastrointestinal tract and functional consequences. <i>Pharmacological Research</i> , 2013 , 69, 127-36	10.2	43
97	Factors involved in the colonization and survival of bifidobacteria in the gastrointestinal tract. <i>FEMS Microbiology Letters</i> , 2013 , 340, 1-10	2.9	46
96	Assessment of intestinal microbiota modulation ability of Bifidobacterium strains in in vitro fecal batch cultures from preterm neonates. <i>Anaerobe</i> , 2013 , 19, 9-16	2.8	35
95	Assessment of the effect of stress-tolerance acquisition on some basic characteristics of specific probiotics. <i>International Journal of Food Microbiology</i> , 2013 , 165, 51-6	5.8	22
94	Microbiota of the Intestine: Probiotics 2013 , 175-181		1

93	Antibiotic resistance in probiotic bacteria. Frontiers in Microbiology, 2013, 4, 202	5.7	273
92	Microbial targets for the development of functional foods accordingly with nutritional and immune parameters altered in the elderly. <i>Journal of the American College of Nutrition</i> , 2013 , 32, 399-406	3.5	52
91	Insights into the ropy phenotype of the exopolysaccharide-producing strain Bifidobacterium animalis subsp. lactis A1dOxR. <i>Applied and Environmental Microbiology</i> , 2013 , 79, 3870-4	4.8	18
90	Population Dynamics of Some Relevant Intestinal Microbial Groups in Human Fecal Batch Cultures with Added Fermentable Xylooligosaccharides Obtained from Rice Husks. <i>BioResources</i> , 2013 , 8,	1.3	5
89	Assessing the fecal microbiota: an optimized ion torrent 16S rRNA gene-based analysis protocol. <i>PLoS ONE</i> , 2013 , 8, e68739	3.7	205
88	Fatty acids intake and immune parameters in the elderly. <i>Nutricion Hospitalaria</i> , 2013 , 28, 474-8	1	6
87	Genome sequence of the immunomodulatory strain Bifidobacterium bifidum LMG 13195. <i>Journal of Bacteriology</i> , 2012 , 194, 6997	3.5	2
86	Immune Modulation Capability of Exopolysaccharides Synthesised by Lactic Acid Bacteria and Bifidobacteria. <i>Probiotics and Antimicrobial Proteins</i> , 2012 , 4, 227-37	5.5	122
85	Treg-inducing membrane vesicles from Bifidobacterium bifidum LMG13195 as potential adjuvants in immunotherapy. <i>Vaccine</i> , 2012 , 30, 825-9	4.1	47
84	Metagenomics and probiotics. Clinical Microbiology and Infection, 2012, 18 Suppl 4, 32-4	9.5	36
83	Exopolysaccharide-producing Bifidobacterium strains elicit different in vitro responses upon interaction with human cells. <i>Food Research International</i> , 2012 , 46, 99-107	7	86
82	Diversity of bifidobacteria within the infant gut microbiota. <i>PLoS ONE</i> , 2012 , 7, e36957	3.7	415
81	Toward improving technological and functional properties of probiotics in foods. <i>Trends in Food Science and Technology</i> , 2012 , 26, 56-63	15.3	34
80	Development of probiotic products for nutritional requirements of specific human populations. <i>Engineering in Life Sciences</i> , 2012 , 12, 368-376	3.4	14
79	Establishment and development of intestinal microbiota in preterm neonates. <i>FEMS Microbiology Ecology</i> , 2012 , 79, 763-72	4.3	268
78	Deep 16S rRNA metagenomics and quantitative PCR analyses of the premature infant fecal microbiota. <i>Anaerobe</i> , 2012 , 18, 378-80	2.8	50
77	Enhancing probiotic stability in industrial processes. <i>Microbial Ecology in Health and Disease</i> , 2012 , 23,		17
76	Controlled gene expression in bifidobacteria by use of a bile-responsive element. <i>Applied and Environmental Microbiology</i> , 2012 , 78, 581-5	4.8	16

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75	Genome sequence of the Antarctic psychrophile bacterium Planococcus antarcticus DSM 14505. Journal of Bacteriology, 2012 , 194, 4465	3.5	11	
74	Molecular clues to understand the aerotolerance phenotype of Bifidobacterium animalis subsp. lactis. <i>Applied and Environmental Microbiology</i> , 2012 , 78, 644-50	4.8	31	
73	Genome sequence of Parascardovia denticolens IPLA 20019, isolated from human breast milk. Journal of Bacteriology, 2012 , 194, 4776-7	3.5	9	
72	Role of extracellular transaldolase from Bifidobacterium bifidum in mucin adhesion and aggregation. <i>Applied and Environmental Microbiology</i> , 2012 , 78, 3992-8	4.8	76	
71	Facultative to strict anaerobes ratio in the preterm infant microbiota: a target for intervention?. <i>Gut Microbes</i> , 2012 , 3, 583-8	8.8	47	
70	Interaction of Bifidobacterium bifidum LMG13195 with HT29 cells influences regulatory-T-cell-associated chemokine receptor expression. <i>Applied and Environmental Microbiology</i> , 2012 , 78, 2850-7	4.8	46	
69	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. <i>Research in Microbiology</i> , 2011 , 162, 514-9	4	36	
68	Immune response to Bifidobacterium bifidum strains support Treg/Th17 plasticity. <i>PLoS ONE</i> , 2011 , 6, e24776	3.7	94	
67	Current and Future Applications of Probiotics. Current Nutrition and Food Science, 2011, 7, 170-180	0.7	2	
66	Evaluation of the ability of Bifidobacterium longum to metabolize human intestinal mucus. <i>FEMS Microbiology Letters</i> , 2011 , 314, 125-30	2.9	20	
65	Assessment of intestinal microbiota of full-term breast-fed infants from two different geographical locations. <i>Early Human Development</i> , 2011 , 87, 511-3	2.2	31	
64	Safety and intestinal microbiota modulation by the exopolysaccharide-producing strains Bifidobacterium animalis IPLA R1 and Bifidobacterium longum IPLA E44 orally administered to Wistar rats. <i>International Journal of Food Microbiology</i> , 2011 , 144, 342-51	5.8	55	
63	Characterization and in vitro properties of potentially probiotic Bifidobacterium strains isolated from breast-milk. <i>International Journal of Food Microbiology</i> , 2011 , 149, 28-36	5.8	92	
62	How do bifidobacteria counteract environmental challenges? Mechanisms involved and physiological consequences. <i>Genes and Nutrition</i> , 2011 , 6, 307-18	4.3	76	
61	Manufacturing process influences properties of probiotic bacteria. <i>British Journal of Nutrition</i> , 2011 , 105, 887-94	3.6	88	
60	Probiotics and Health Claims: How to be Met by SMEs? 2010 , 263-270			
59	Probiotics in Adhesion of Pathogens 2010 , 353-370		19	
58	Fecal Bifidobacterium Levels in Elderly Nursing Home Patients. <i>Bioscience and Microflora</i> , 2010 , 29, 111-	113	12	

57	Genetic basis of tetracycline resistance in Bifidobacterium animalis subsp. lactis. <i>Applied and Environmental Microbiology</i> , 2010 , 76, 3364-9	4.8	57
56	Establishment and development of lactic acid bacteria and bifidobacteria microbiota in breast-milk and the infant gut. <i>Anaerobe</i> , 2010 , 16, 307-10	2.8	219
55	Distinct Bifidobacterium strains drive different immune responses in vitro. <i>International Journal of Food Microbiology</i> , 2010 , 138, 157-65	5.8	122
54	Bile affects the synthesis of exopolysaccharides by Bifidobacterium animalis. <i>Applied and Environmental Microbiology</i> , 2009 , 75, 1204-7	4.8	81
53	Microbial-host interactions: selecting the right probiotics and prebiotics for infants. <i>Nestle Nutrition Workshop Series Paediatric Programme</i> , 2009 , 64, 201-13; discussion 213-7, 251-7		15
52	Bile-inducible efflux transporter from Bifidobacterium longum NCC2705, conferring bile resistance. <i>Applied and Environmental Microbiology</i> , 2009 , 75, 3153-60	4.8	58
51	Coculture of Bifidobacterium longum and Bifidobacterium breve alters their protein expression profiles and enzymatic activities. <i>International Journal of Food Microbiology</i> , 2009 , 133, 148-53	5.8	35
50	Microbiomic analysis of the bifidobacterial population in the human distal gut. <i>ISME Journal</i> , 2009 , 3, 745-51	11.9	111
49	Probiotic fermented milks: Present and future. International Journal of Dairy Technology, 2009, 62, 472-	4 8 3	44
48	Effect of clinical and probiotic Lactobacillus rhamnosus strains on intestinal permeability and bacterial translocation in healthy and colitic rats. <i>Food Research International</i> , 2009 , 42, 636-640	7	8
47	Inflammation markers and malnutrition as risk factors for infections and impaired health-related quality of life among older nursing home residents. <i>Journal of the American Medical Directors Association</i> , 2009 , 10, 348-53	5.9	30
46	Safety Assessment of Probiotics 2009 , 1193-1235		10
45	Degradation of 16S rRNA and attributes of viability of viable but nonculturable probiotic bacteria. <i>Letters in Applied Microbiology</i> , 2008 , 46, 693-8	2.9	69
44	Probiotic Microorganisms 2008 , 1-176		O
43	Two different tetracycline resistance mechanisms, plasmid-carried tet(L) and chromosomally located transposon-associated tet(M), coexist in Lactobacillus sakei Rits 9. <i>Applied and Environmental Microbiology</i> , 2008 , 74, 1394-401	4.8	63
42	Mucin degradation by Bifidobacterium strains isolated from the human intestinal microbiota. <i>Applied and Environmental Microbiology</i> , 2008 , 74, 1936-40	4.8	159
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