## Mirjana Rajilic-Stojanovic

List of Publications by Year in descending order

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54 papers 8,309 citations

186254 28 h-index 206102 48 g-index

54 all docs

54 docs citations

54 times ranked

11435 citing authors

#	Article	IF	CITATIONS
1	The first 1000 cultured species of the human gastrointestinal microbiota. FEMS Microbiology Reviews, 2014, 38, 996-1047.	8.6	923
2	Global and Deep Molecular Analysis of Microbiota Signatures in Fecal Samples From Patients With Irritable Bowel Syndrome. Gastroenterology, 2011, 141, 1792-1801.	1.3	885
3	European consensus conference on faecal microbiota transplantation in clinical practice. Gut, 2017, 66, 569-580.	12.1	793
4	Irritable bowel syndrome. Nature Reviews Disease Primers, 2016, 2, 16014.	30.5	674
5	High-throughput diversity and functionality analysis of the gastrointestinal tract microbiota. Gut, 2008, 57, 1605-1615.	12.1	528
6	Diversity of the human gastrointestinal tract microbiota revisited. Environmental Microbiology, 2007, 9, 2125-2136.	3.8	485
7	Comparative analysis of fecal DNA extraction methods with phylogenetic microarray: Effective recovery of bacterial and archaeal DNA using mechanical cell lysis. Journal of Microbiological Methods, 2010, 81, 127-134.	1.6	480
8	Development and application of the human intestinal tract chip, a phylogenetic microarray: analysis of universally conserved phylotypes in the abundant microbiota of young and elderly adults. Environmental Microbiology, 2009, 11, 1736-1751.	3.8	420
9	Clinical trial: multispecies probiotic supplementation alleviates the symptoms of irritable bowel syndrome and stabilizes intestinal microbiota. Alimentary Pharmacology and Therapeutics, 2008, 27, 48-57.	3.7	309
10	Phylogenetic Analysis of Dysbiosis in Ulcerative Colitis During Remission. Inflammatory Bowel Diseases, 2013, 19, 481-488.	1.9	285
11	Intestinal Microbiota And Diet in IBS: Causes, Consequences, or Epiphenomena?. American Journal of Gastroenterology, 2015, 110, 278-287.	0.4	283
12	High temporal and interâ€individual variation detected in the human ileal microbiota. Environmental Microbiology, 2010, 12, 3213-3227.	3.8	254
13	Colonic Microbiota Signatures across Five Northern European Countries. Applied and Environmental Microbiology, 2005, 71, 4153-4155.	3.1	243
14	The microbial eukaryote <i>Blastocystis</i> is a prevalent and diverse member of the healthy human gut microbiota. FEMS Microbiology Ecology, 2014, 90, 326-330.	2.7	208
15	Longâ€ŧerm monitoring of the human intestinal microbiota composition. Environmental Microbiology, 2013, 15, 1146-1159.	3.8	195
16	Rome Foundation Working Team Report on Post-Infection Irritable Bowel Syndrome. Gastroenterology, 2019, 156, 46-58.e7.	1.3	162
17	Linking phylogenetic identities of bacteria to starch fermentation in an ⟨i⟩in vitro⟨ i⟩ model of the large intestine by RNAâ€based stable isotope probing. Environmental Microbiology, 2009, 11, 914-926.	3.8	157
18	Faecal Microbiota Composition in Adults Is Associated with the FUT2 Gene Determining the Secretor Status. PLoS ONE, 2014, 9, e94863.	2.5	129

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19	Microarray analysis reveals marked intestinal microbiota aberrancy in infants having eczema compared to healthy children in at-risk for atopic disease. BMC Microbiology, 2013, 13, 12.	3.3	127
20	Systematic review: gastric microbiota in health and disease. Alimentary Pharmacology and Therapeutics, 2020, 51, 582-602.	3.7	113
21	Evaluating the microbial diversity of an in vitro model of the human large intestine by phylogenetic microarray analysis. Microbiology (United Kingdom), 2010, 156, 3270-3281.	1.8	84
22	Function of the microbiota. Bailliere's Best Practice and Research in Clinical Gastroenterology, 2013, 27, 5-16.	2.4	81
23	Optimisation of microwave-assisted extraction parameters for antioxidants from waste Achillea millefolium dust. Industrial Crops and Products, 2015, 77, 333-341.	5.2	55
24	Improvement of mechanical properties and antibacterial activity of crosslinked electrospun chitosan/poly (ethylene oxide) nanofibers. Composites Part B: Engineering, 2017, 121, 58-67.	12.0	49
25	Cefazolin-loaded polycaprolactone fibers produced via different electrospinning methods: Characterization, drug release and antibacterial effect. European Journal of Pharmaceutical Sciences, 2018, 124, 26-36.	4.0	45
26	Plant Extracts Rich in Polyphenols as Potent Modulators in the Growth of Probiotic and Pathogenic Intestinal Microorganisms. Frontiers in Nutrition, 2021, 8, 688843.	3.7	40
27	Microwave-assisted extraction for the recovery of antioxidants from waste Equisetum arvense. Industrial Crops and Products, 2014, 61, 388-397.	5.2	34
28	From Agricultural Waste to Biofuel: Enzymatic Potential of a Bacterial Isolate Streptomyces fulvissimus CKS7 for Bioethanol Production. Waste and Biomass Valorization, 2021, 12, 165-174.	3.4	34
29	Water Kefir grain as a source of potent dextran producing lactic acid bacteria. Hemijska Industrija, 2015, 69, 595-604.	0.7	26
30	Molecular methods for the analysis of gut microbiota. Microbial Ecology in Health and Disease, 2004, 16, 71-85.	3.5	25
31	Lignocellulosic waste material as substrate for Avicelase production by a new strain of Paenibacillus chitinolyticus CKS1. International Biodeterioration and Biodegradation, 2015, 104, 426-434.	3.9	20
32	Biocontrol and plant stimulating potential of novel strain Bacillus sp. PPM3 isolated from marine sediment. Microbial Pathogenesis, 2018, 120, 71-78.	2.9	18
33	Considerations for the design and conduct of human gut microbiota intervention studies relating to foods. European Journal of Nutrition, 2020, 59, 3347-3368.	3.9	17
34	The Interrelationship Among Non-Alcoholic Fatty Liver Disease, Colonic Diverticulosis and Metabolic Syndrome. Journal of Gastrointestinal and Liver Diseases, 2021, 30, 274-282.	0.9	17
35	Helicobacter Pylori Eradication Therapy is Not Associated with the Onset of Inflammatory Bowel Diseases. A Case-Control Study. Journal of Gastrointestinal and Liver Diseases, 2019, 27, 119-125.	0.9	15
36	Enzymatic hydrolysis of waste bread by newly isolated Hymenobacter sp. CKS3: Statistical optimization and bioethanol production. Renewable Energy, 2020, 152, 627-633.	8.9	13

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37	Does Day-to-Day Variability in Stool Consistency Link to the Fecal Microbiota Composition?. Frontiers in Cellular and Infection Microbiology, 2021, 11, 639667.	3.9	11
38	Valorization of corn stover and molasses for enzyme synthesis, lignocellulosic hydrolysis and bioethanol production by Hymenobacter sp. CKS3. Environmental Technology and Innovation, 2021, 23, 101627.	6.1	9
39	Carboxymethyl cellulase production from a Paenibacillus sp Hemijska Industrija, 2016, 70, 329-338.	0.7	9
40	Development of an Environmentally Acceptable Detergent Formulation for Fatty Soils Based on the Lipase from the Indigenous Extremophile <i>Pseudomonas aeruginosa</i> Surfactants and Detergents, 2015, 18, 383-395.	2.1	8
41	Sugar Beet Pulp as Leuconostoc mesenteroides T3 Support for Enhanced Dextransucrase Production on Molasses. Applied Biochemistry and Biotechnology, 2016, 180, 1016-1027.	2.9	7
42	Multiscale characterization of antimicrobial poly(vinyl butyral)/titania nanofibrous composites. Polymers for Advanced Technologies, 2017, 28, 909-914.	3.2	7
43	Enhanced fertilization effect of a compost obtained from mixed herbs waste inoculated with novel strains of mesophilic bacteria. Hemijska Industrija, 2017, 71, 503-513.	0.7	7
44	Vitamin B Complex and Experimental Autoimmune Encephalomyelitis –Attenuation of the Clinical Signs and Gut Microbiota Dysbiosis. Nutrients, 2022, 14, 1273.	4.1	7
45	Characterization of dextransucrase from Leuconostoc mesenteroides T3, water kefir grains isolate. Hemijska Industrija, 2017, 71, 351-360.	0.7	5
46	Valorization of lignocellulosic wastes for extracellular enzyme production by novel Basidiomycetes: screening, hydrolysis, and bioethanol production. Biomass Conversion and Biorefinery, 0, , 1.	4.6	4
47	$\hat{l}^2$ -Amylase production from packaging-industry wastewater using a novel strain Paenibacillus chitinolyticus CKS 1. RSC Advances, 2015, 5, 90895-90903.	3.6	3
48	Recovery of bioactive molecules from Hypericum perforatum L. dust using microwave-assisted extraction. Biomass Conversion and Biorefinery, 2024, 14, 7111-7123.	4.6	3
49	Gut Microbiota and its Role in Human Health. Psihologijske Teme, 2018, 27, 17-32.	0.2	1
50	Microbiota Changes Throughout Life - An Overview. , 2022, , 1-12.		1
51	The Human Intestinal Microbiota and Its Impact on Health. , 0, , 11-32.		1
52	Utilization of agro-industrial by-products as substrates for dextransucrase production by Leuconostoc mesenteroides T3: Process optimization using response surface methodology. Hemijska Industrija, 2021, 75, 135-146.	0.7	0
53	Lactic Acid Bacteria in the Gut. , 2019, , 383-408.		O
54	Optimization of spray drying conditions for production of Achillea millefolium extract powder. Hemijska Industrija, 2021, 75, 353-363.	0.7	0