

# Ilaria Patuzzi

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4690357/publications.pdf>

Version: 2024-02-01

17  
papers

386  
citations

933447

10  
h-index

888059

17  
g-index

17  
all docs

17  
docs citations

17  
times ranked

758  
citing authors

#	ARTICLE	IF	CITATIONS
1	Assessing the Influence of Vegan, Vegetarian and Omnivore Oriented Westernized Dietary Styles on Human Gut Microbiota: A Cross Sectional Study. <i>Frontiers in Microbiology</i> , 2018, 9, 317.	3.5	78
2	Testing nano-silver food packaging to evaluate silver migration and food spoilage bacteria on chicken meat. <i>Food Additives and Contaminants - Part A Chemistry, Analysis, Control, Exposure and Risk Assessment</i> , 2016, 33, 1063-1071.	2.3	49
3	Silver As Antibacterial toward <i>Listeria monocytogenes</i> . <i>Frontiers in Microbiology</i> , 2016, 7, 307.	3.5	42
4	SPARSim single cell: a count data simulator for scRNA-seq data. <i>Bioinformatics</i> , 2020, 36, 1468-1475.	4.1	33
5	Bovine meat versus pork in <i>Toxoplasma gondii</i> transmission in Italy: A quantitative risk assessment model. <i>International Journal of Food Microbiology</i> , 2018, 269, 1-11.	4.7	31
6	Effectiveness of Washing Procedures in Reducing <i>Salmonella enterica</i> and <i>Listeria monocytogenes</i> on a Raw Leafy Green Vegetable ( <i>Eruca vesicaria</i> ). <i>Frontiers in Microbiology</i> , 2016, 7, 1663.	3.5	30
7	A specific microbiota signature is associated to various degrees of ulcerative colitis as assessed by a machine learning approach. <i>Gut Microbes</i> , 2022, 14, 2028366.	9.8	26
8	Assessing antimicrobial resistance gene load in vegan, vegetarian and omnivore human gut microbiota. <i>International Journal of Antimicrobial Agents</i> , 2018, 52, 702-705.	2.5	18
9	Evaluation and quantification of antimicrobial residues and antimicrobial resistance genes in two Italian swine farms. <i>Environmental Pollution</i> , 2019, 255, 113183.	7.5	17
10	metaSPARSim: a 16S rRNA gene sequencing count data simulator. <i>BMC Bioinformatics</i> , 2019, 20, 416.	2.6	16
11	Faecal Microbiome Transplantation as a Solution to Chronic Enteropathies in Dogs: A Case Study of Beneficial Microbial Evolution. <i>Animals</i> , 2021, 11, 1433.	2.3	12
12	The Interplay between <i>Campylobacter</i> and the Caecal Microbial Community of Commercial Broiler Chickens over Time. <i>Microorganisms</i> , 2021, 9, 221.	3.6	10
13	Effect of sterilization on the canine vaginal microbiota: a pilot study. <i>BMC Veterinary Research</i> , 2020, 16, 455.	1.9	8
14	Possible Influence of Natural Events on Heavy Metals Exposure from Shellfish Consumption: A Case Study in the North-East of Italy. <i>Frontiers in Public Health</i> , 2015, 3, 21.	2.7	7
15	Characterization of intestinal microbiota in normal weight and overweight Border Collie and Labrador Retriever dogs. <i>Scientific Reports</i> , 2022, 12, .	3.3	4
16	Beware to ignore the rare: how imputing zero-values can improve the quality of 16S rRNA gene studies results. <i>BMC Bioinformatics</i> , 2021, 22, 618.	2.6	3
17	Modeling Microbial Community Networks: Methods and Tools. <i>Current Genomics</i> , 2021, 22, 267-290.	1.6	2