

Narciso M Quijada

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3332566/publications.pdf>

Version: 2024-02-01

20
papers

765
citations

567281

15
h-index

713466

21
g-index

21
all docs

21
docs citations

21
times ranked

1163
citing authors

#	ARTICLE	IF	CITATIONS
1	Identifying Beneficial Qualities of <i>Trichoderma parareesei</i> for Plants. Applied and Environmental Microbiology, 2014, 80, 1864-1873.	3.1	124
2	Co-occurrence of colistin-resistance genes mcr-1 and mcr-3 among multidrug-resistant Escherichia coli isolated from cattle, Spain, September 2015. Eurosurveillance, 2017, 22, .	7.0	100
3	TORMES: an automated pipeline for whole bacterial genome analysis. Bioinformatics, 2019, 35, 4207-4212.	4.1	82
4	Microbiota of newborn calves and their mothers reveals possible transfer routes for newborn calves' gastrointestinal microbiota. PLoS ONE, 2019, 14, e0220554.	2.5	61
5	Dynamics of the oral microbiota as a tool to estimate time since death. Molecular Oral Microbiology, 2017, 32, 511-516.	2.7	52
6	Daily thanatomicrobiome changes in soil as an approach of postmortem interval estimation: An ecological perspective. Forensic Science International, 2017, 278, 388-395.	2.2	47
7	Different Lactobacillus populations dominate in 'Chorizo de Le ³ ' manufacturing performed in different production plants. Food Microbiology, 2018, 70, 94-102.	4.2	41
8	Autochthonous facility-specific microbiota dominates washed-rind Austrian hard cheese surfaces and its production environment. International Journal of Food Microbiology, 2018, 267, 54-61.	4.7	31
9	Fecal Microbiota of Toxigenic Clostridioides difficile-Associated Diarrhea. Frontiers in Microbiology, 2018, 9, 3331.	3.5	30
10	Virulence characterization and comparative genomics of Listeria monocytogenes sequence type 155 strains. BMC Genomics, 2020, 21, 847.	2.8	26
11	Co-Occurrence of Listeria spp. and Spoilage Associated Microbiota During Meat Processing Due to Cross-Contamination Events. Frontiers in Microbiology, 2021, 12, 632935.	3.5	26
12	Escherichia coli ST167 carrying plasmid mobilisable mcr-1 and blaCTX-M-15 resistance determinants isolated from a human respiratory infection. International Journal of Antimicrobial Agents, 2017, 50, 285-286.	2.5	24
13	Oxacillin-susceptible mecA-positive Staphylococcus aureus associated with processed food in Europe. Food Microbiology, 2019, 82, 107-110.	4.2	21
14	High-throughput sequencing and food microbiology. Advances in Food and Nutrition Research, 2020, 91, 275-300.	3.0	21
15	Propidium Monoazide Integrated with qPCR Enables the Detection and Enumeration of Infectious Enteric RNA and DNA Viruses in Clam and Fermented Sausages. Frontiers in Microbiology, 2016, 7, 2008.	3.5	20
16	Co-infection of Chicken Layers With Histomonas meleagridis and Avian Pathogenic Escherichia coli Is Associated With Dysbiosis, Cecal Colonization and Translocation of the Bacteria From the Gut Lumen. Frontiers in Microbiology, 2020, 11, 586437.	3.5	16
17	The Core Human Microbiome: Does It Exist and How Can We Find It? A Critical Review of the Concept. Nutrients, 2022, 14, 2872.	4.1	16
18	Austrian Raw-Milk Hard-Cheese Ripening Involves Successional Dynamics of Non-Inoculated Bacteria and Fungi. Foods, 2020, 9, 1851.	4.3	13

#	ARTICLE	IF	CITATIONS
19	Dietary Supplementation with Sugar Beet Fructooligosaccharides and Garlic Residues Promotes Growth of Beneficial Bacteria and Increases Weight Gain in Neonatal Lambs. <i>Biomolecules</i> , 2020, 10, 1179.	4.0	4
20	Infrequent isolation of extensively drug-resistant (XDR) <i>Klebsiella pneumoniae</i> resistant to colistin in Spain. <i>International Journal of Antimicrobial Agents</i> , 2018, 51, 531-533.	2.5	2