

# Daniel Jañ©n-Luchoro

## List of Publications by Year in descending order

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Version: 2024-02-01

26  
papers

278  
citations

1162367

8  
h-index

1058022

14  
g-index

28  
all docs

28  
docs citations

28  
times ranked

332  
citing authors

#	ARTICLE	IF	CITATIONS
1	Everything Is Everywhere: Physiological Responses of the Mediterranean Sea and Eastern Pacific Ocean Epiphyte <i>Cobetia</i> Sp. to Varying Nutrient Concentration. <i>Microbial Ecology</i> , 2022, 83, 296-313.	1.4	3
2	Identification of Antibiotic Resistance Proteins via MiCIdâ€™s Augmented Workflow. A Mass Spectrometry-Based Proteomics Approach. <i>Journal of the American Society for Mass Spectrometry</i> , 2022, 33, 917-931.	1.2	3
3	Identity of blaCTX-M Carrying Plasmids in Sequential ESBL- <i>E. coli</i> Isolates from Patients with Recurrent Urinary Tract Infections. <i>Microorganisms</i> , 2021, 9, 1138.	1.6	7
4	Analyses of Virulence Genes of <i>Clavibacter michiganensis</i> subsp. <i>michiganensis</i> Strains Reveal Heterogeneity and Deletions That Correlate with Pathogenicity. <i>Microorganisms</i> , 2021, 9, 1530.	1.6	4
5	Comparative Genomics of Clinical Isolates of the Emerging Tick-Borne Pathogen <i>Neoehrlichia mikurensis</i> . <i>Microorganisms</i> , 2021, 9, 1488.	1.6	9
6	Mass Spectrometry Proteotyping-Based Detection and Identification of <i>Staphylococcus aureus</i> , <i>Escherichia coli</i> , and <i>Candida albicans</i> in Blood. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021, 11, 634215.	1.8	5
7	<i>Corynebacterium sanguinis</i> sp. nov., a clinical and environmental associated corynebacterium. <i>Systematic and Applied Microbiology</i> , 2020, 43, 126039.	1.2	20
8	Complete genome sequences of <i>Streptococcus pyogenes</i> type strain reveal 100%-match between PacBio-solo and Illumina-Oxford Nanopore hybrid assemblies. <i>Scientific Reports</i> , 2020, 10, 11656.	1.6	5
9	Comparative Genomics of Pathogenic <i>Clavibacter michiganensis</i> subsp. <i>michiganensis</i> Strains from Chile Reveals Potential Virulence Features for Tomato Plants. <i>Microorganisms</i> , 2020, 8, 1679.	1.6	14
10	Discovery of Species-unique Peptide Biomarkers of Bacterial Pathogens by Tandem Mass Spectrometry-based Proteotyping. <i>Molecular and Cellular Proteomics</i> , 2020, 19, 518-528.	2.5	22
11	Genomic and Proteomic Characterization of the Extended-Spectrum Î²-Lactamase (ESBL)-Producing <i>Escherichia coli</i> Strain CCUG 73778: A Virulent, Nosocomial Outbreak Strain. <i>Microorganisms</i> , 2020, 8, 893.	1.6	6
12	A Pangenome Approach for Discerning Species-Unique Gene Markers for Identifications of <i>Streptococcus pneumoniae</i> and <i>Streptococcus pseudopneumoniae</i> . <i>Frontiers in Cellular and Infection Microbiology</i> , 2020, 10, 222.	1.8	18
13	<i>Staphylococcus borealis</i> sp. nov., isolated from human skin and blood. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 6067-6078.	0.8	23
14	Beware of False "Type Strain" Genome Sequences. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.3	3
15	Complete Genome Sequence of the Marine Hydrocarbon Degradator <i>Alcaligenes aquatilis</i> QD168, Isolated from Crude Oil-Polluted Sediment of Quintero Bay, Central Chile. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.3	17
16	Complete Genome Sequence of Hydrocarbon-Degrading Halotolerant <i>Acinetobacter radioresistens</i> DD78, Isolated from the Aconcagua River Mouth in Central Chile. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.3	5
17	Complete genome sequence of the marine <i>Rhodococcus</i> sp. H-CA8f isolated from Comau fjord in Northern Patagonia, Chile. <i>Marine Genomics</i> , 2018, 40, 13-17.	0.4	9
18	Complete Genome Sequence of the Hydrocarbon-Degrading Strain <i>Achromobacter</i> sp. B7, Isolated during Petroleum Hydrocarbon Bioremediation in the Valparaíso Region, Chile. <i>Microbiology Resource Announcements</i> , 2018, 7, .	0.3	4

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19	Proteotyping bacteria: Characterization, differentiation and identification of pneumococcus and other species within the Mitis Group of the genus <i>Streptococcus</i> by tandem mass spectrometry proteomics. <i>PLoS ONE</i> , 2018, 13, e0208804.	1.1	47
20	Genome Sequences of Two Naphthalene-Degrading Strains of <i>Pseudomonas balearica</i> , Isolated from Polluted Marine Sediment and from an Oil Refinery Site. <i>Genome Announcements</i> , 2017, 5, .	0.8	3
21	First insights into a type II toxin-antitoxin system from the clinical isolate <i>Mycobacterium</i> sp. MHSD3, similar to epsilon/zeta systems. <i>PLoS ONE</i> , 2017, 12, e0189459.	1.1	6
22	Draft Genome Sequence of <i>Streptococcus gordonii</i> Type Strain CCUG 33482 T. <i>Genome Announcements</i> , 2016, 4, .	0.8	1
23	Complete Genome Sequence of <i>Pseudomonas balearica</i> DSM 6083 T. <i>Genome Announcements</i> , 2016, 4, .	0.8	6
24	Complete Genome Sequence of <i>Mycobacterium chelonae</i> Type Strain CCUG 47445, a Rapidly Growing Species of Nontuberculous Mycobacteria. <i>Genome Announcements</i> , 2016, 4, .	0.8	1
25	Complete Genome Sequence of the <i>Mycobacterium immunogenum</i> Type Strain CCUG 47286. <i>Genome Announcements</i> , 2016, 4, .	0.8	2
26	A protocol for extraction and purification of high-quality and quantity bacterial DNA applicable for genome sequencing: a modified version of the Marmur procedure.. <i>Protocol Exchange</i> , 0, , .	0.3	34