

# Carlos SÃ¡nchez Casimiro-Soriguer

## List of Publications by Year in descending order

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Version: 2024-02-01

19  
papers

252  
citations

1478505

6  
h-index

1058476

14  
g-index

22  
all docs

22  
docs citations

22  
times ranked

401  
citing authors

#	ARTICLE	IF	CITATIONS
1	Towards a metagenomics machine learning interpretable model for understanding the transition from adenoma to colorectal cancer. <i>Scientific Reports</i> , 2022, 12, 450.	3.3	11
2	Genome-wide analysis of DNA methylation in Hirschsprung enteric precursor cells: unraveling the epigenetic landscape of enteric nervous system development. <i>Clinical Epigenetics</i> , 2021, 13, 51.	4.1	4
3	Taxonomic variations in the gut microbiome of gout patients with and without tophi might have a functional impact on urate metabolism. <i>Molecular Medicine</i> , 2021, 27, 50.	4.4	31
4	Phylogenetic Analysis of the 2020 West Nile Virus (WNV) Outbreak in Andalusia (Spain). <i>Viruses</i> , 2021, 13, 836.	3.3	13
5	Highly accurate whole-genome imputation of SARS-CoV-2 from partial or low-quality sequences. <i>GigaScience</i> , 2021, 10, .	6.4	2
6	Virulence role of the outer membrane protein CarO in carbapenem-resistant <i>Acinetobacter baumannii</i> . <i>Virulence</i> , 2020, 11, 1727-1737.	4.4	16
7	Ancient evolutionary signals of protein-coding sequences allow the discovery of new genes in the <i>Drosophila melanogaster</i> genome. <i>BMC Genomics</i> , 2020, 21, 210.	2.8	3
8	Using AnAblast for intergenic sORF prediction in the <i>Caenorhabditis elegans</i> genome. <i>Bioinformatics</i> , 2020, 36, 4827-4832.	4.1	6
9	Antibiotic resistance and metabolic profiles as functional biomarkers that accurately predict the geographic origin of city metagenomics samples. <i>Biology Direct</i> , 2019, 14, 15.	4.6	17
10	AnAblast: Re-searching for Protein-Coding Sequences in Genomic Regions. <i>Methods in Molecular Biology</i> , 2019, 1962, 207-214.	0.9	4
11	Multiple Genome Sequences of <i>Lactobacillus pentosus</i> Strains Isolated from Biofilms on the Skin of Fermented Green Table Olives. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.6	5
12	Draft Genome Sequences of Six <i>Lactobacillus pentosus</i> Strains Isolated from Brines of Traditionally Fermented Spanish-Style Green Table Olives. <i>Genome Announcements</i> , 2018, 6, .	0.8	7
13	Sma3s: A universal tool for easy functional annotation of proteomes and transcriptomes. <i>Proteomics</i> , 2017, 17, 1700071.	2.2	71
14	Insight into Potential Probiotic Markers Predicted in <i>Lactobacillus pentosus</i> MP-10 Genome Sequence. <i>Frontiers in Microbiology</i> , 2017, 8, 891.	3.5	47
15	PlantFuncSSR: Integrating First and Next Generation Transcriptomics for Mining of SSR-Functional Domains Markers. <i>Frontiers in Plant Science</i> , 2016, 7, 878.	3.6	5
16	<i>Crepis</i> sect. <i>Lepidoseris</i> sensu Babcock en la Península Ibérica y Baleares. <i>Crepis</i> sect. <i>Lepidoseris</i> sensu Babcock in the Iberian Peninsula and Balearic Islands. <i>Acta Botanica Malacitana</i> , 0, 38, 231-240.	0.0	4
17	Tipificación de <i>Crepis bursifolia</i> y su distribución en Marruecos. Typification of <i>Crepis bursifolia</i> and chorology in Morocco. <i>Acta Botanica Malacitana</i> , 0, 38, 222-224.	0.0	1
18	Reencuentro de <i>Crepis balliana</i> Babc. en Marruecos. On the presence of <i>Crepis balliana</i> Babc. in Morocco. <i>Acta Botanica Malacitana</i> , 0, 38, 219-221.	0.0	0

#	ARTICLE	IF	CITATIONS
19	Crepis alpina en España. Crepis alpina in Spain. Acta Botanica Malacitana, 0, 38, 229-231.	0.0	0