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List of Publications by Year in descending order

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21
papers

934
citations

687363

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794594

19
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29
all docs

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docs citations

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times ranked

2023
citing authors

#	ARTICLE	IF	CITATIONS
1	Characteristics of SARS-CoV-2 variants of concern B.1.1.7, B.1.351 or P.1: data from seven EU/EEA countries, weeks 38/2020 to 10/2021. <i>Eurosurveillance</i> , 2021, 26, .	7.0	216
2	Tracking the international spread of SARS-CoV-2 lineages B.1.1.7 and B.1.351/501Y-V2 with grinch. <i>Wellcome Open Research</i> , 2021, 6, 121.	1.8	129
3	Mutation rate of SARS-CoV-2 and emergence of mutators during experimental evolution. <i>Evolution, Medicine and Public Health</i> , 2022, 10, 142-155.	2.5	101
4	Tracking SARS-CoV-2 lineage B.1.1.7 dissemination: insights from nationwide spike gene target failure (SGTF) and spike gene late detection (SGTL) data, Portugal, week 49 2020 to week 3 2021. <i>Eurosurveillance</i> , 2021, 26, .	7.0	64
5	Long-Term Evolution of SARS-CoV-2 in an Immunocompromised Patient with Non-Hodgkin Lymphoma. <i>MSphere</i> , 2021, 6, e0024421.	2.9	63
6	INNUENDO: A cross-sectoral platform for the integration of genomics in the surveillance of food-borne pathogens. <i>EFSA Supporting Publications</i> , 2018, 15, 1498E.	0.7	56
7	Nosocomial Outbreak of SARS-CoV-2 in a "Non-COVID-19" Hospital Ward: Virus Genome Sequencing as a Key Tool to Understand Cryptic Transmission. <i>Viruses</i> , 2021, 13, 604.	3.3	45
8	Screening of Bacteriocinogenic Lactic Acid Bacteria and Their Characterization as Potential Probiotics. <i>Microorganisms</i> , 2020, 8, 393.	3.6	40
9	Virulence and antibiotic resistance plasticity of <i>Arcobacter butzleri</i> : Insights on the genomic diversity of an emerging human pathogen. <i>Infection, Genetics and Evolution</i> , 2020, 80, 104213.	2.3	38
10	Genomic Study of a <i>Clostridium difficile</i> Multidrug Resistant Outbreak-Related Clone Reveals Novel Determinants of Resistance. <i>Frontiers in Microbiology</i> , 2018, 9, 2994.	3.5	25
11	Imipenem Resistance in <i>Clostridium difficile</i> Ribotype 017, Portugal. <i>Emerging Infectious Diseases</i> , 2018, 24, 741-745.	4.3	24
12	Massive dissemination of a SARS-CoV-2 Spike Y839 variant in Portugal. <i>Emerging Microbes and Infections</i> , 2020, 9, 2488-2496.	6.5	20
13	<i>Chlamydia trachomatis</i> : when the virulence-associated genome backbone imports a prevalence-associated major antigen signature. <i>Microbial Genomics</i> , 2019, 5, .	2.0	18
14	Characterization of Multidrug-Resistant Isolates of <i>Salmonella enterica</i> Serovars Heidelberg and Minnesota from Fresh Poultry Meat Imported to Portugal. <i>Microbial Drug Resistance</i> , 2021, 27, 87-98.	2.0	13
15	Transcontinental Dissemination of the L2b/D-Da Recombinant <i>Chlamydia trachomatis</i> Lymphogranuloma venereum (LGV) Strain: Need of Broad Multi-Country Molecular Surveillance. <i>Clinical Infectious Diseases</i> , 2021, 73, e1004-e1007.	5.8	10
16	Overview of <i>Clostridium difficile</i> Infection: Life Cycle, Epidemiology, Antimicrobial Resistance and Treatment. , 0, , .		9
17	Multidrug-Resistant <i>Salmonella enterica</i> Serovar Rissen Clusters Detected in Azores Archipelago, Portugal. <i>International Journal of Genomics</i> , 2019, 2019, 1-9.	1.6	9
18	<i>Neisseria gonorrhoeae</i> clustering to reveal major European whole-genome-sequencing-based genogroups in association with antimicrobial resistance. <i>Microbial Genomics</i> , 2021, 7, .	2.0	9

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19	SARS-CoV-2 introductions and early dynamics of the epidemic in Portugal. <i>Communications Medicine</i> , 2022, 2, .	4.2	5
20	Potential recurrence of COVID-19 in a healthcare professional: SARS-CoV-2 genome sequencing confirms contagiousness after re-positivity. <i>International Journal of Infectious Diseases</i> , 2021, 112, 318-320.	3.3	2
21	Unraveling the hurdles of a large COVID-19 epidemiological investigation by viral genomics. <i>Journal of Infection</i> , 2022, 85, 64-74.	3.3	0