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## List of Publications by Year in descending order

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687363 794594 21 934 13 19 h-index citations g-index papers 29 29 29 2023 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	SARS-CoV-2 introductions and early dynamics of the epidemic in Portugal. Communications Medicine, 2022, 2, .	4.2	5
2	Mutation rate of SARS-CoV-2 and emergence of mutators during experimental evolution. Evolution, Medicine and Public Health, 2022, 10, 142-155.	2.5	101
3	Unraveling the hurdles of a large COVID-19 epidemiological investigation by viral genomics. Journal of Infection, 2022, 85, 64-74.	3.3	O
4	Characterization of Multidrug-Resistant Isolates of <i>Salmonella enterica</i> Serovars Heidelberg and Minnesota from Fresh Poultry Meat Imported to Portugal. Microbial Drug Resistance, 2021, 27, 87-98.	2.0	13
5	Neisseria gonorrhoeae clustering to reveal major European whole-genome-sequencing-based genogroups in association with antimicrobial resistance. Microbial Genomics, 2021, 7, .	2.0	9
6	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. Eurosurveillance, 2021, 26, .	7.0	64
7	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. Eurosurveillance, 2021, 26, .	7.0	216
8	Nosocomial Outbreak of SARS-CoV-2 in a "Non-COVID-19―Hospital Ward: Virus Genome Sequencing as a Key Tool to Understand Cryptic Transmission. Viruses, 2021, 13, 604.	3.3	45
9	Long-Term Evolution of SARS-CoV-2 in an Immunocompromised Patient with Non-Hodgkin Lymphoma. MSphere, 2021, 6, e0024421.	2.9	63
10	Potential recurrence of COVID-19 in a healthcare professional: SARS-CoV-2 genome sequencing confirms contagiousness after re-positivity. International Journal of Infectious Diseases, 2021, 112, 318-320.	3.3	2
11	Tracking the international spread of SARS-CoV-2 lineages B.1.1.7 and B.1.351/501Y-V2 with grinch. Wellcome Open Research, 2021, 6, 121.	1.8	129
12	Transcontinental Dissemination of the L2b/D-Da Recombinant <i>Chlamydia trachomatis</i> Lymphogranuloma venereum (LGV) Strain: Need of Broad Multi-Country Molecular Surveillance. Clinical Infectious Diseases, 2021, 73, e1004-e1007.	5.8	10
13	Massive dissemination of a SARS-CoV-2 Spike Y839 variant in Portugal. Emerging Microbes and Infections, 2020, 9, 2488-2496.	6.5	20
14	Screening of Bacteriocinogenic Lactic Acid Bacteria and Their Characterization as Potential Probiotics. Microorganisms, 2020, 8, 393.	3.6	40
15	Virulence and antibiotic resistance plasticity of Arcobacter butzleri: Insights on the genomic diversity of an emerging human pathogen. Infection, Genetics and Evolution, 2020, 80, 104213.	2.3	38
16	Multidrug-Resistant <i>Salmonella enterica </i> Serovar Rissen Clusters Detected in Azores Archipelago, Portugal. International Journal of Genomics, 2019, 2019, 1-9.	1.6	9
17	Chlamydia trachomatis: when the virulence-associated genome backbone imports a prevalence-associated major antigen signature. Microbial Genomics, 2019, 5, .	2.0	18
18	INNUENDO: A crossâ€sectoral platform for the integration of genomics in the surveillance of foodâ€borne pathogens. EFSA Supporting Publications, 2018, 15, 1498E.	0.7	56

#	Article	IF	CITATION
19	Genomic Study of a Clostridium difficile Multidrug Resistant Outbreak-Related Clone Reveals Novel Determinants of Resistance. Frontiers in Microbiology, 2018, 9, 2994.	3.5	25
20	Imipenem Resistance in <i>Clostridium difficile</i> Ribotype 017, Portugal. Emerging Infectious Diseases, 2018, 24, 741-745.	4.3	24
21	Overview of Clostridium difficile Infection: Life Cycle, Epidemiology, Antimicrobial Resistance and Treatment. , 0, , .		9