

Filipe Zimmer Dezordi

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8740980/publications.pdf>

Version: 2024-02-01

13
papers

588
citations

933410

10
h-index

1199563

12
g-index

19
all docs

19
docs citations

19
times ranked

1161
citing authors

#	ARTICLE	IF	CITATIONS
1	COVID-19 in Amazonas, Brazil, was driven by the persistence of endemic lineages and P.1 emergence. <i>Nature Medicine</i> , 2021, 27, 1230-1238.	30.7	279
2	Multiple Introductions Followed by Ongoing Community Spread of SARS-CoV-2 at One of the Largest Metropolitan Areas of Northeast Brazil. <i>Viruses</i> , 2020, 12, 1414.	3.3	47
3	Lying in wait: the resurgence of dengue virus after the Zika epidemic in Brazil. <i>Nature Communications</i> , 2021, 12, 2619.	12.8	43
4	ViralFlow: A Versatile Automated Workflow for SARS-CoV-2 Genome Assembly, Lineage Assignment, Mutations and Intra-host Variant Detection. <i>Viruses</i> , 2022, 14, 217.	3.3	40
5	The ongoing evolution of variants of concern and interest of SARS-CoV-2 in Brazil revealed by convergent indels in the amino (N)-terminal domain of the spike protein. <i>Virus Evolution</i> , 2021, 7, veab069.	4.9	31
6	Identification of a novel SARS-CoV-2 P.1 sub-lineage in Brazil provides new insights about the mechanisms of emergence of variants of concern. <i>Virus Evolution</i> , 2021, 7, veab091.	4.9	28
7	In and Outs of Chuviridae Endogenous Viral Elements: Origin of a Potentially New Retrovirus and Signature of Ancient and Ongoing Arms Race in Mosquito Genomes. <i>Frontiers in Genetics</i> , 2020, 11, 542437.	2.3	26
8	Systematic Review of Wolbachia Symbiont Detection in Mosquitoes: An Entangled Topic about Methodological Power and True Symbiosis. <i>Pathogens</i> , 2021, 10, 39.	2.8	25
9	Unusual SARS-CoV-2 intra-host diversity reveals lineage superinfection. <i>Microbial Genomics</i> , 2022, 8, .	2.0	18
10	HTT-DB: new features and updates. <i>Database: the Journal of Biological Databases and Curation</i> , 2018, 2018, .	3.0	16
11	Phylogenetic-based inference reveals distinct transmission dynamics of SARS-CoV-2 lineages Gamma and P.2 in Brazil. <i>iScience</i> , 2022, 25, 104156.	4.1	16
12	Metatranscriptomic analysis identifies different viral-like sequences in two neotropical <i>Mansoniini</i> mosquito species. <i>Virus Research</i> , 2021, 301, 198455.	2.2	4
13	Os 5 passos essenciais para constru�o de �rvores filogen�ticas. , 0, , .		0