

Gularte, Js

List of Publications by Year in descending order

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Version: 2024-02-01

17
papers

261
citations

1307594

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h-index

996975

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21
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21
docs citations

21
times ranked

552
citing authors

#	ARTICLE	IF	CITATIONS
1	Y380Q novel mutation in receptor-binding domain of SARS-CoV-2 spike protein together with C379W interfere in the neutralizing antibodies interaction. <i>Diagnostic Microbiology and Infectious Disease</i> , 2022, 102, 115636.	1.8	2
2	Early introduction, dispersal and evolution of Delta SARS-CoV-2 in Southern Brazil, late predominance of AY.99.2 and AY.101 related lineages. <i>Virus Research</i> , 2022, 311, 198702.	2.2	15
3	Hepatitis E virus genotype 3 in bovine livers slaughtered in the state of Rio Grande do Sul, Brazil. <i>Brazilian Journal of Microbiology</i> , 2022, 53, 1115-1120.	2.0	6
4	Viral isolation allows characterization of early samples of SARS-CoV-2 lineage B.1.1.33 with unique mutations (S: H655Y and T63N) circulating in Southern Brazil in 2020. <i>Brazilian Journal of Microbiology</i> , 2022, 53, 1313-1319.	2.0	2
5	SARS-CoV-2 and COVID-19: A perspective from environmental virology. <i>Genetics and Molecular Biology</i> , 2021, 44, e20200228.	1.3	2
6	Microbial Source Tracking in Small Farms: Use of Different Methods for Adenovirus Detection. <i>Water, Air, and Soil Pollution</i> , 2021, 232, 1.	2.4	3
7	Pervasive transmission of E484K and emergence of VUI-NP13L with evidence of SARS-CoV-2 co-infection events by two different lineages in Rio Grande do Sul, Brazil. <i>Virus Research</i> , 2021, 296, 198345.	2.2	105
8	Low circulation of Influenza A and coinfection with SARS-CoV-2 among other respiratory viruses during the COVID-19 pandemic in a region of southern Brazil. <i>Journal of Medical Virology</i> , 2021, 93, 4392-4398.	5.0	22
9	Genomic epidemiology of SARS-CoV-2 in Esteio, Rio Grande do Sul, Brazil. <i>BMC Genomics</i> , 2021, 22, 371.	2.8	22
10	Reinfection cases by closely related SARS-CoV-2 lineages in Southern Brazil. <i>Brazilian Journal of Microbiology</i> , 2021, 52, 1881-1885.	2.0	2
11	Functionalized Surfaces as a Tool for Virus Sensing: A Demonstration of Human mastadenovirus Detection in Environmental Waters. <i>Chemosensors</i> , 2021, 9, 19.	3.6	1
12	Early detection of SARS-CoV-2 P.1 variant in Southern Brazil and reinfection of the same patient by P.2. <i>Revista Do Instituto De Medicina Tropical De Sao Paulo</i> , 2021, 63, e58.	1.1	31
13	Predominance of SARS-CoV-2 P.1 (Gamma) lineage inducing the recent COVID-19 wave in southern Brazil and the finding of an additional S: D614A mutation. <i>Infection, Genetics and Evolution</i> , 2021, 96, 105134.	2.3	11
14	Temporal dynamics of Human mastadenovirus species in cases of respiratory illness in southern Brazil. <i>Brazilian Journal of Microbiology</i> , 2019, 50, 677-684.	2.0	3
15	Human mastadenovirus in water, sediment, sea surface microlayer, and bivalve mollusk from southern Brazilian beaches. <i>Marine Pollution Bulletin</i> , 2019, 142, 335-349.	5.0	18
16	Evaluation of the Gravata-River sediment quality (Rio Grande do Sul- Brazil) using <i>Daphnia magna</i> (Straus, 1820) as the test-organism for toxicity assays. <i>Acta Limnologica Brasiliensia</i> , 2010, 22, 367-377.	0.4	2
17	Bioassay using <i>Daphnia magna</i> Straus, 1820 to evaluate the sediment of Ca-River (Rio Grande do Sul,) Tj ETQq1 1.0,784314,rgBT /Ore	0.4	3