

Theocharis Tsoleridis

List of Publications by Year in descending order

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Version: 2024-02-01

21
papers

309
citations

1464605

7
h-index

1113639

15
g-index

28
all docs

28
docs citations

28
times ranked

956
citing authors

#	ARTICLE	IF	CITATIONS
1	The Impact of Real-Time Whole-Genome Sequencing in Controlling Healthcare-Associated SARS-CoV-2 Outbreaks. <i>Journal of Infectious Diseases</i> , 2022, 225, 10-18.	1.9	25
2	Enterovirus D68 epidemic, UK, 2018, was caused by subclades B3 and D1, predominantly in children and adults, respectively, with both subclades exhibiting extensive genetic diversity. <i>Microbial Genomics</i> , 2022, 8, .	1.0	2
3	Coronavirus (CoV) proteins in GtoPdb v.2022.2. <i>IUPHAR/BPS Guide To Pharmacology CITE</i> , 2022, 2022, .	0.2	0
4	Retrospective screening of routine respiratory samples revealed undetected community transmission and missed intervention opportunities for SARS-CoV-2 in the United Kingdom. <i>Journal of General Virology</i> , 2021, 102, .	1.3	10
5	Immunogenicity of a new gorilla adenovirus vaccine candidate for COVID-19. <i>Molecular Therapy</i> , 2021, 29, 2412-2423.	3.7	41
6	Two doses of the SARS-CoV-2 BNT162b2 vaccine enhance antibody responses to variants in individuals with prior SARS-CoV-2 infection. <i>Science Translational Medicine</i> , 2021, 13, eabj0847.	5.8	40
7	Coronavirus (CoV) proteins in GtoPdb v.2021.3. <i>IUPHAR/BPS Guide To Pharmacology CITE</i> , 2021, 2021, .	0.2	0
8	Polymer microarrays rapidly identify competitive adsorbents of virus-like particles. <i>Biointerphases</i> , 2020, 15, 061005.	0.6	5
9	Discovery and Prevalence of Divergent RNA Viruses in European Field Voles and Rabbits. <i>Viruses</i> , 2020, 12, 47.	1.5	9
10	Retrieval of the Complete Coding Sequence of the UK-Endemic Tatenale Orthohantavirus Reveals Extensive Strain Variation and Supports Its Classification as a Novel Species. <i>Viruses</i> , 2020, 12, 454.	1.5	4
11	SARS-CoV-2 proteins (version 2020.2) in the IUPHAR/BPS Guide to Pharmacology Database. <i>IUPHAR/BPS Guide To Pharmacology CITE</i> , 2020, 2020, .	0.2	4
12	Discovery of Novel Coronaviruses in Rodents. <i>Methods in Molecular Biology</i> , 2020, 2203, 33-40.	0.4	4
13	Coronavirus (CoV) proteins (version 2020.4) in the IUPHAR/BPS Guide to Pharmacology Database. <i>IUPHAR/BPS Guide To Pharmacology CITE</i> , 2020, 2020, .	0.2	1
14	Shared Common Ancestry of Rodent Alphacoronaviruses Sampled Globally. <i>Viruses</i> , 2019, 11, 125.	1.5	35
15	Identification of Infectious Agents in High-Throughput Sequencing Data Sets Is Easily Achievable Using Free, Cloud-Based Bioinformatics Platforms. <i>Journal of Clinical Microbiology</i> , 2019, 57, .	1.8	3
16	Wombling surplus diagnostic nucleic acid for novel pathogenesis and genetic epidemiology of viral infections. <i>Access Microbiology</i> , 2019, 1, .	0.2	0
17	Discovery of novel highly divergent RNA viruses in European rodents and rabbits. <i>Access Microbiology</i> , 2019, 1, .	0.2	0
18	Investigation and genome characterisation of tatenale hantavirus in wild rodent populations in the United Kingdom. <i>Access Microbiology</i> , 2019, 1, .	0.2	0

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19	High-throughput sequencing of patients with symptoms of unknown etiology. <i>Access Microbiology</i> , 2019, 1, .	0.2	0
20	Discovery of Novel Alphacoronaviruses in European Rodents and Shrews. <i>Viruses</i> , 2016, 8, 84.	1.5	45
21	A Diverse Panel of Hepatitis C Virus Glycoproteins for Use in Vaccine Research Reveals Extremes of Monoclonal Antibody Neutralization Resistance. <i>Journal of Virology</i> , 2016, 90, 3288-3301.	1.5	62