

Christopher Ruis

List of Publications by Year in descending order

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Version: 2024-02-01

14
papers

3,685
citations

687363

13
h-index

1058476

14
g-index

18
all docs

18
docs citations

18
times ranked

7491
citing authors

#	ARTICLE	IF	CITATIONS
1	A dynamic nomenclature proposal for SARS-CoV-2 lineages to assist genomic epidemiology. <i>Nature Microbiology</i> , 2020, 5, 1403-1407.	13.3	2,291
2	Producing polished prokaryotic pangenomes with the Panaroo pipeline. <i>Genome Biology</i> , 2020, 21, 180.	8.8	419
3	Establishment and lineage dynamics of the SARS-CoV-2 epidemic in the UK. <i>Science</i> , 2021, 371, 708-712.	12.6	335
4	Spatiotemporal invasion dynamics of SARS-CoV-2 lineage B.1.1.7 emergence. <i>Science</i> , 2021, 373, 889-895.	12.6	142
5	Stepwise pathogenic evolution of <i>Mycobacterium abscessus</i> . <i>Science</i> , 2021, 372, .	12.6	91
6	Norovirus Whole-Genome Sequencing by SureSelect Target Enrichment: a Robust and Sensitive Method. <i>Journal of Clinical Microbiology</i> , 2016, 54, 2530-2537.	3.9	67
7	Addendum: A dynamic nomenclature proposal for SARS-CoV-2 lineages to assist genomic epidemiology. <i>Nature Microbiology</i> , 2021, 6, 415-415.	13.3	65
8	The emerging GII.P16-GII.4 Sydney 2012 norovirus lineage is circulating worldwide, arose by late-2014 and contains polymerase changes that may increase virus transmission. <i>PLoS ONE</i> , 2017, 12, e0179572.	2.5	63
9	Global phylogeny of <i>Treponema pallidum</i> lineages reveals recent expansion and spread of contemporary syphilis. <i>Nature Microbiology</i> , 2021, 6, 1549-1560.	13.3	51
10	Dissemination of <i>Mycobacterium abscessus</i> via global transmission networks. <i>Nature Microbiology</i> , 2021, 6, 1279-1288.	13.3	47
11	Progress and challenges in virus genomic epidemiology. <i>Trends in Parasitology</i> , 2021, 37, 1038-1049.	3.3	45
12	A comprehensive characterization of chronic norovirus infection in immunodeficient hosts. <i>Journal of Allergy and Clinical Immunology</i> , 2019, 144, 1450-1453.	2.9	24
13	Preadaptation of pandemic GII.4 noroviruses in unsampled virus reservoirs years before emergence. <i>Virus Evolution</i> , 2020, 6, veaa067.	4.9	22
14	Novel Insights Into the Spread of Enteric Pathogens Using Genomics. <i>Journal of Infectious Diseases</i> , 2019, 221, S319-S330.	4.0	2