

Christopher Ruis

List of Publications by Year in Descending Order

Source: <https://exaly.com/author-pdf/7649232/christopher-ruis-publications-by-year.pdf>

Version: 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

17
papers

1,754
citations

9
h-index

18
g-index

18
ext. papers

2,924
ext. citations

17.9
avg, IF

5.55
L-index

#	Paper	IF	Citations
17	Context-specific emergence and growth of the SARS-CoV-2 Delta variant. 2021 ,		3
16	Global phylogeny of <i>Treponema pallidum</i> lineages reveals recent expansion and spread of contemporary syphilis. <i>Nature Microbiology</i> , 2021 , 6, 1549-1560	26.6	2
15	Progress and challenges in virus genomic epidemiology. <i>Trends in Parasitology</i> , 2021 , 37, 1038-1049	6.4	4
14	Stepwise pathogenic evolution of. <i>Science</i> , 2021 , 372,	33.3	23
13	Spatiotemporal invasion dynamics of SARS-CoV-2 lineage B.1.1.7 emergence. <i>Science</i> , 2021 , 373, 889-895	33.3	41
12	Addendum: A dynamic nomenclature proposal for SARS-CoV-2 lineages to assist genomic epidemiology. <i>Nature Microbiology</i> , 2021 , 6, 415	26.6	35
11	Establishment and lineage dynamics of the SARS-CoV-2 epidemic in the UK. <i>Science</i> , 2021 , 371, 708-712	33.3	159
10	Dissemination of <i>Mycobacterium abscessus</i> via global transmission networks. <i>Nature Microbiology</i> , 2021 , 6, 1279-1288	26.6	4
9	Context-specific emergence and growth of the SARS-CoV-2 Delta variant. 2021 ,		2
8	Novel Insights Into the Spread of Enteric Pathogens Using Genomics. <i>Journal of Infectious Diseases</i> , 2020 , 221, S319-S330	7	1
7	Preadaptation of pandemic GII.4 noroviruses in unsampled virus reservoirs years before emergence. <i>Virus Evolution</i> , 2020 , 6, veaa067	3.7	9
6	A dynamic nomenclature proposal for SARS-CoV-2 lineages to assist genomic epidemiology. <i>Nature Microbiology</i> , 2020 , 5, 1403-1407	26.6	1250
5	Producing polished prokaryotic pangenomes with the Panaroo pipeline. <i>Genome Biology</i> , 2020 , 21, 180	18.3	108
4	A comprehensive characterization of chronic norovirus infection in immunodeficient hosts. <i>Journal of Allergy and Clinical Immunology</i> , 2019 , 144, 1450-1453	11.5	10
3	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	3.7	52
2	Norovirus Whole-Genome Sequencing by SureSelect Target Enrichment: a Robust and Sensitive Method. <i>Journal of Clinical Microbiology</i> , 2016 , 54, 2530-7	9.7	50
1	Contemporary syphilis is characterised by rapid global spread of pandemic <i>Treponema pallidum</i> lineages		1

