

# Christopher Ruis

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7649232/publications.pdf>

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