

# Samuel M Nicholls

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9135084/publications.pdf>

Version: 2024-02-01

10  
papers

2,579  
citations

1039406

9  
h-index

1372195

10  
g-index

19  
all docs

19  
docs citations

19  
times ranked

6226  
citing authors

#	ARTICLE	IF	CITATIONS
1	Future-proofing and maximizing the utility of metadata: The PHA4GE SARS-CoV-2 contextual data specification package. <i>GigaScience</i> , 2022, 11, .	3.3	18
2	Evaluating the Effects of SARS-CoV-2 Spike Mutation D614G on Transmissibility and Pathogenicity. <i>Cell</i> , 2021, 184, 64-75.e11.	13.5	843
3	On the complexity of haplotyping a microbial community. <i>Bioinformatics</i> , 2021, 37, 1360-1366.	1.8	17
4	Establishment and lineage dynamics of the SARS-CoV-2 epidemic in the UK. <i>Science</i> , 2021, 371, 708-712.	6.0	335
5	SARS-CoV-2 within-host diversity and transmission. <i>Science</i> , 2021, 372, .	6.0	278
6	CLIMB-COVID: continuous integration supporting decentralised sequencing for SARS-CoV-2 genomic surveillance. <i>Genome Biology</i> , 2021, 22, 196.	3.8	53
7	Generation and transmission of interlineage recombinants in the SARS-CoV-2 pandemic. <i>Cell</i> , 2021, 184, 5179-5188.e8.	13.5	182
8	Evolution and epidemic spread of SARS-CoV-2 in Brazil. <i>Science</i> , 2020, 369, 1255-1260.	6.0	454
9	Ultra-deep, long-read nanopore sequencing of mock microbial community standards. <i>GigaScience</i> , 2019, 8, .	3.3	209
10	Goldilocks: a tool for identifying genomic regions that are "just right". <i>Bioinformatics</i> , 2016, 32, 2047-2049.	1.8	3