

Shirlee Wohl

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2543571/publications.pdf>

Version: 2024-02-01

22
papers

2,718
citations

687363

13
h-index

940533

16
g-index

32
all docs

32
docs citations

32
times ranked

5340
citing authors

#	ARTICLE	IF	CITATIONS
1	Repeated Coronavirus Disease 2019 Molecular Testing: Correlation of Severe Acute Respiratory Syndrome Coronavirus 2 Culture With Molecular Assays and Cycle Thresholds. <i>Clinical Infectious Diseases</i> , 2021, 73, e860-e869.	5.8	163
2	Genomic diversity of SARS-CoV-2 during early introduction into the Baltimore-Washington metropolitan area. <i>JCI Insight</i> , 2021, 6, .	5.0	31
3	Regional sequencing collaboration reveals persistence of the T12 <i>Vibrio cholerae</i> O1 lineage in West Africa. <i>ELife</i> , 2021, 10, .	6.0	6
4	Sample size calculation for phylogenetic case linkage. <i>PLoS Computational Biology</i> , 2021, 17, e1009182.	3.2	7
5	Combining genomics and epidemiology to track mumps virus transmission in the United States. <i>PLoS Biology</i> , 2020, 18, e3000611.	5.6	37
6	Successive epidemic waves of cholera in South Sudan between 2014 and 2017: a descriptive epidemiological study. <i>Lancet Planetary Health</i> , The, 2020, 4, e577-e587.	11.4	18
7	Combining genomics and epidemiology to track mumps virus transmission in the United States. , 2020, 18, e3000611.		0
8	Combining genomics and epidemiology to track mumps virus transmission in the United States. , 2020, 18, e3000611.		0
9	Combining genomics and epidemiology to track mumps virus transmission in the United States. , 2020, 18, e3000611.		0
10	Combining genomics and epidemiology to track mumps virus transmission in the United States. , 2020, 18, e3000611.		0
11	Combining genomics and epidemiology to track mumps virus transmission in the United States. , 2020, 18, e3000611.		0
12	Combining genomics and epidemiology to track mumps virus transmission in the United States. , 2020, 18, e3000611.		0
13	Capturing sequence diversity in metagenomes with comprehensive and scalable probe design. <i>Nature Biotechnology</i> , 2019, 37, 160-168.	17.5	96
14	Genomic Analysis of Lassa Virus during an Increase in Cases in Nigeria in 2018. <i>New England Journal of Medicine</i> , 2018, 379, 1745-1753.	27.0	135
15	Virus genomes reveal factors that spread and sustained the Ebola epidemic. <i>Nature</i> , 2017, 544, 309-315.	27.8	346
16	Zika virus evolution and spread in the Americas. <i>Nature</i> , 2017, 546, 411-415.	27.8	323
17	Evidence of Ebola Virus Replication and High Concentration in Semen of a Patient During Recovery. <i>Clinical Infectious Diseases</i> , 2017, 65, 1400-1403.	5.8	26
18	Ebola Virus Epidemiology and Evolution in Nigeria. <i>Journal of Infectious Diseases</i> , 2016, 214, S102-S109.	4.0	19

#	ARTICLE	IF	CITATIONS
19	Genomic Analysis of Viral Outbreaks. Annual Review of Virology, 2016, 3, 173-195.	6.7	61
20	Evaluation of the Potential Impact of Ebola Virus Genomic Drift on the Efficacy of Sequence-Based Candidate Therapeutics. MBio, 2015, 6, .	4.1	62
21	Ebola Virus Epidemiology, Transmission, and Evolution during Seven Months in Sierra Leone. Cell, 2015, 161, 1516-1526.	28.9	275
22	Genomic surveillance elucidates Ebola virus origin and transmission during the 2014 outbreak. Science, 2014, 345, 1369-1372.	12.6	1,083