Shirlee Wohl

List of Publications by Year in descending order

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

Version: 2024-02-01

687363 940533 2,718 22 13 16 citations h-index g-index papers 32 32 32 5340 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Genomic surveillance elucidates Ebola virus origin and transmission during the 2014 outbreak. Science, 2014, 345, 1369-1372.	12.6	1,083
2	Virus genomes reveal factors that spread and sustained the Ebola epidemic. Nature, 2017, 544, 309-315.	27.8	346
3	Zika virus evolution and spread in the Americas. Nature, 2017, 546, 411-415.	27.8	323
4	Ebola Virus Epidemiology, Transmission, and Evolution during Seven Months in Sierra Leone. Cell, 2015, 161, 1516-1526.	28.9	275
5	Repeated Coronavirus Disease 2019 Molecular Testing: Correlation of Severe Acute Respiratory Syndrome Coronavirus 2 Culture With Molecular Assays and Cycle Thresholds. Clinical Infectious Diseases, 2021, 73, e860-e869.	5.8	163
6	Genomic Analysis of Lassa Virus during an Increase in Cases in Nigeria in 2018. New England Journal of Medicine, 2018, 379, 1745-1753.	27.0	135
7	Capturing sequence diversity in metagenomes with comprehensive and scalable probe design. Nature Biotechnology, 2019, 37, 160-168.	17.5	96
8	Evaluation of the Potential Impact of Ebola Virus Genomic Drift on the Efficacy of Sequence-Based Candidate Therapeutics. MBio, 2015, 6, .	4.1	62
9	Genomic Analysis of Viral Outbreaks. Annual Review of Virology, 2016, 3, 173-195.	6.7	61
10	Combining genomics and epidemiology to track mumps virus transmission in the United States. PLoS Biology, 2020, 18, e3000611.	5.6	37
11	Genomic diversity of SARS-CoV-2 during early introduction into the Baltimore–Washington metropolitan area. JCl Insight, 2021, 6, .	5.0	31
12	Evidence of Ebola Virus Replication and High Concentration in Semen of a Patient During Recovery. Clinical Infectious Diseases, 2017, 65, 1400-1403.	5.8	26
13	Ebola Virus Epidemiology and Evolution in Nigeria. Journal of Infectious Diseases, 2016, 214, S102-S109.	4.0	19
14	Successive epidemic waves of cholera in South Sudan between 2014 and 2017: a descriptive epidemiological study. Lancet Planetary Health, The, 2020, 4, e577-e587.	11.4	18
15	Sample size calculation for phylogenetic case linkage. PLoS Computational Biology, 2021, 17, e1009182.	3.2	7
16	Regional sequencing collaboration reveals persistence of the T12 Vibrio cholerae O1 lineage in West Africa. ELife, 2021, 10 , .	6.0	6
17	Combining genomics and epidemiology to track mumps virus transmission in the United States. , 2020, 18, e3000611.		O
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19	Combining genomics and epidemiology to track mumps virus transmission in the United States. , 2020, 18, e3000611.		O
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