

List of Publications by Citations

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Version: 2024-04-09

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

15 papers	143 citations	4 h-index	11 g-index
19 ext. papers	297 ext. citations	8.5 avg, IF	1.3 L-index

#	Paper	IF	Citations
15	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	59.2	82
14	A year of genomic surveillance reveals how the SARS-CoV-2 pandemic unfolded in Africa. <i>Science</i> , 2021 , 374, 423-431	33.3	35
13	Real-time Metagenomic Analysis of Undiagnosed Fever Cases Unveils a Yellow Fever Outbreak in Edo State, Nigeria. <i>Scientific Reports</i> , 2020 , 10, 3180	4.9	10
12	Polymorphisms in Plasmodium falciparum chloroquine resistance transporter (Pfcr1) and multidrug-resistant gene 1 (Pfmdr-1) in Nigerian children 10 years post-adoption of artemisinin-based combination treatments. <i>International Journal for Parasitology</i> , 2021 , 51, 301-310	4.3	4
11	First report of Wolbachia from field populations of Culex mosquitoes in south-western Nigeria. <i>African Zoology</i> , 2019 , 54, 181-185	1.1	3
10	HIV-1 drug resistance and genetic diversity in a cohort of people with HIV-1 in Nigeria. <i>Aids</i> , 2022 , 36, 137-146	3.5	2
9	Future-proofing and maximizing the utility of metadata: The PHA4GE SARS-CoV-2 contextual data specification package.. <i>GigaScience</i> , 2022 , 11,	7.6	2
8	Lassa Virus Genetics. <i>Current Topics in Microbiology and Immunology</i> , 2020 , 1	3.3	1
7	Microbial Metagenomic Approach Uncovers the First Rabbit Haemorrhagic Disease Virus genome in Sub-Saharan Africa		1
6	Genetic diversity and population structure of Plasmodium falciparum in Nigeria: insights from microsatellite loci analysis. <i>Malaria Journal</i> , 2021 , 20, 236	3.6	1
5	Microbial metagenomic approach uncovers the first rabbit haemorrhagic disease virus genome in Sub-Saharan Africa. <i>Scientific Reports</i> , 2021 , 11, 13689	4.9	1
4	VGEA: an RNA viral assembly toolkit. <i>PeerJ</i> , 2021 , 9, e12129	3.1	1
3	Polymorphisms in Plasmodium falciparum dihydropteroate synthetase and dihydrofolate reductase genes in Nigerian children with uncomplicated malaria using high-resolution melting technique. <i>Scientific Reports</i> , 2021 , 11, 471	4.9	0
2	Molecular profiling of the artemisinin resistance Kelch 13 gene in Plasmodium falciparum from Nigeria.. <i>PLoS ONE</i> , 2022 , 17, e0264548	3.7	0
1	Metagenomic sequencing characterizes a wide diversity of viruses in field mosquito samples in Nigeria.. <i>Scientific Reports</i> , 2022 , 12, 7616	4.9	0