

# Paul E Oluniyi

## List of Publications by Year in Descending Order

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This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.  
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	HIV-1 drug resistance and genetic diversity in a cohort of people with HIV-1 in Nigeria. <i>Aids</i> , <b>2022</b> , 36, 137-146	3.5	2
14	Future-proofing and maximizing the utility of metadata: The PHA4GE SARS-CoV-2 contextual data specification package.. <i>GigaScience</i> , <b>2022</b> , 11,	7.6	2
13	Molecular profiling of the artemisinin resistance Kelch 13 gene in <i>Plasmodium falciparum</i> from Nigeria.. <i>PLoS ONE</i> , <b>2022</b> , 17, e0264548	3.7	0
12	Metagenomic sequencing characterizes a wide diversity of viruses in field mosquito samples in Nigeria.. <i>Scientific Reports</i> , <b>2022</b> , 12, 7616	4.9	0
11	A year of genomic surveillance reveals how the SARS-CoV-2 pandemic unfolded in Africa. <i>Science</i> , <b>2021</b> , 374, 423-431	33.3	35
10	Polymorphisms in <i>Plasmodium falciparum</i> 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> , <b>2021</b> , 51, 301-310	4.3	4
9	Genetic diversity and population structure of <i>Plasmodium falciparum</i> in Nigeria: insights from microsatellite loci analysis. <i>Malaria Journal</i> , <b>2021</b> , 20, 236	3.6	1
8	Polymorphisms in <i>Plasmodium falciparum</i> dihydropteroate synthetase and dihydrofolate reductase genes in Nigerian children with uncomplicated malaria using high-resolution melting technique. <i>Scientific Reports</i> , <b>2021</b> , 11, 471	4.9	0
7	Microbial metagenomic approach uncovers the first rabbit haemorrhagic disease virus genome in Sub-Saharan Africa. <i>Scientific Reports</i> , <b>2021</b> , 11, 13689	4.9	1
6	VGEA: an RNA viral assembly toolkit. <i>PeerJ</i> , <b>2021</b> , 9, e12129	3.1	1
5	Lassa Virus Genetics. <i>Current Topics in Microbiology and Immunology</i> , <b>2020</b> , 1	3.3	1
4	Real-time Metagenomic Analysis of Undiagnosed Fever Cases Unveils a Yellow Fever Outbreak in Edo State, Nigeria. <i>Scientific Reports</i> , <b>2020</b> , 10, 3180	4.9	10
3	First report of <i>Wolbachia</i> from field populations of <i>Culex</i> mosquitoes in south-western Nigeria. <i>African Zoology</i> , <b>2019</b> , 54, 181-185	1.1	3
2	Genomic Analysis of Lassa Virus during an Increase in Cases in Nigeria in 2018. <i>New England Journal of Medicine</i> , <b>2018</b> , 379, 1745-1753	59.2	82
1	Microbial Metagenomic Approach Uncovers the First Rabbit Haemorrhagic Disease Virus genome in Sub-Saharan Africa		1