Lynette Isabella Ochola-Oyier

List of Publications by Year in descending order

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Lynette Isabella

#	Article	IF	CITATIONS
1	Seroprevalence of Antibodies to Severe Acute Respiratory Syndrome Coronavirus 2 Among Healthcare Workers in Kenya. Clinical Infectious Diseases, 2022, 74, 288-293.	5.8	36
2	Comparative performance of WANTAI ELISA for total immunoglobulin to receptor binding protein and an ELISA for IgG to spike protein in detecting SARS-CoV-2 antibodies in Kenyan populations. Journal of Clinical Virology, 2022, 146, 105061.	3.1	14
3	Epidemiology of COVID-19 infections on routine polymerase chain reaction (PCR) and serology testing in Coastal Kenya. Wellcome Open Research, 2022, 7, 69.	1.8	12
4	Targeted Amplicon Deep Sequencing for Monitoring Antimalarial Resistance Markers in Western Kenya. Antimicrobial Agents and Chemotherapy, 2022, 66, e0194521.	3.2	4
5	Genomic Epidemiology of SARS-CoV-2 in Seychelles, 2020–2021. Viruses, 2022, 14, 1318.	3.3	3
6	Maintenance of high temporal Plasmodium falciparum genetic diversity and complexity of infection in asymptomatic and symptomatic infections in Kilifi, Kenya from 2007 to 2018. Malaria Journal, 2022, 21, .	2.3	4
7	Seroprevalence of anti–SARS-CoV-2 IgG antibodies in Kenyan blood donors. Science, 2021, 371, 79-82.	12.6	247
8	An open dataset of Plasmodium falciparum genome variation in 7,000 worldwide samples. Wellcome Open Research, 2021, 6, 42.	1.8	97
9	The Design and Evaluation of a Mobile System for Rapid Diagnostic Test Interpretation. , 2021, 5, 1-26.		12
10	Anti-Severe Acute Respiratory Syndrome Coronavirus 2 Immunoglobulin G Antibody Seroprevalence Among Truck Drivers and Assistants in Kenya. Open Forum Infectious Diseases, 2021, 8, ofab314.	0.9	12
11	Temporal trends of SARS-CoV-2 seroprevalence during the first wave of the COVID-19 epidemic in Kenya. Nature Communications, 2021, 12, 3966.	12.8	40
12	The Plasmodium falciparum Rh5 invasion protein complex reveals an excess of rare variant mutations. Malaria Journal, 2021, 20, 278.	2.3	10
13	Tracking the introduction and spread of SARS-CoV-2 in coastal Kenya. Nature Communications, 2021, 12, 4809.	12.8	32
14	A review of the frequencies of Plasmodium falciparum Kelch 13 artemisinin resistance mutations in Africa. International Journal for Parasitology: Drugs and Drug Resistance, 2021, 16, 155-161.	3.4	42
15	COVID-19 transmission dynamics underlying epidemic waves in Kenya. Science, 2021, 374, 989-994.	12.6	62
16	An optimisation of four SARS-CoV-2 qRT-PCR assays in a Kenyan laboratory to support the national COVID-19 rapid response teams. Wellcome Open Research, 2020, 5, 162.	1.8	13
17	Pooled testing conserves SARS-CoV-2 laboratory resources and improves test turn-around time: experience on the Kenyan Coast. Wellcome Open Research, 2020, 5, 186.	1.8	13
18	Understanding P. falciparum Asymptomatic Infections: A Proposition for a Transcriptomic Approach. Frontiers in Immunology, 2019, 10, 2398.	4.8	27

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19	Targeted Next Generation Sequencing for malaria research in Africa: current status and outlook. Malaria Journal, 2019, 18, 324.	2.3	9
20	Few Plasmodium falciparum merozoite ligand and erythrocyte receptor pairs show evidence of balancing selection. Infection, Genetics and Evolution, 2019, 69, 235-245.	2.3	7
21	No Evidence of Plasmodium falciparum <i>k13</i> Artemisinin Resistance-Conferring Mutations over a 24-Year Analysis in Coastal Kenya but a Near Complete Reversion to Chloroquine-Sensitive Parasites. Antimicrobial Agents and Chemotherapy, 2019, 63, .	3.2	26
22	Peripheral blood mononuclear cell transcriptomes reveal an over-representation of down-regulated genes associated with immunity in HIV-exposed uninfected infants. Scientific Reports, 2019, 9, 18124.	3.3	8
23	Transmission and Age Impact the Risk of Developing Febrile Malaria in Children with Asymptomatic Plasmodium falciparum Parasitemia. Journal of Infectious Diseases, 2019, 219, 936-944.	4.0	20
24	Polymorphisms in the K13 Gene in Plasmodium falciparum from Different Malaria Transmission Areas of Kenya. American Journal of Tropical Medicine and Hygiene, 2018, 98, 1360-1366.	1.4	30
25	Geographic-genetic analysis of Plasmodium falciparum parasite populations from surveys of primary school children in Western Kenya. Wellcome Open Research, 2017, 2, 29.	1.8	14
26	Comparison of allele frequencies of Plasmodium falciparum merozoite antigens in malaria infections sampled in different years in a Kenyan population. Malaria Journal, 2016, 15, 261.	2.3	7
27	Implications from predicted B-cell and T-cell epitopes of Plasmodium falciparum merozoite proteins EBA175-RII and Rh5. Bioinformation, 2016, 12, 82-91.	0.5	2
28	The MSPDBL2 Codon 591 Polymorphism Is Associated with Lumefantrine <i>In Vitro</i> Drug Responses in Plasmodium falciparum Isolates from Kilifi, Kenya. Antimicrobial Agents and Chemotherapy, 2015, 59, 1770-1775.	3.2	2
29	Temporal trends in prevalence of Plasmodium falciparum drug resistance alleles over two decades of changing antimalarial policy in coastal Kenya. International Journal for Parasitology: Drugs and Drug Resistance, 2014, 4, 152-163.	3.4	34
30	Piperaquine and Lumefantrine resistance in Plasmodium berghei ANKA associated with increased expression of Ca2+/H+ antiporter and glutathione associated enzymes. Experimental Parasitology, 2014, 147, 23-32.	1.2	8
31	The Polymorphic Linker Domain ofpfmdr1Is Associated with Resistance-Conferring Mutations in Plasmodium falciparum Populations from East and West Africa. Antimicrobial Agents and Chemotherapy, 2013, 57, 4595-4598.	3.2	3
32	Repeat Polymorphisms in the Low-Complexity Regions of Plasmodium falciparum ABC Transporters and Associations with <i>In Vitro</i> Antimalarial Responses. Antimicrobial Agents and Chemotherapy, 2013, 57, 6196-6204.	3.2	16
33	Revealing the extent of the first wave of the COVID-19 pandemic in Kenya based on serological and PCR-test data. Wellcome Open Research, 0, 6, 127.	1.8	8
34	Maintaining laboratory quality assurance and safety in a pandemic: Experiences from the KEMRI-Wellcome Trust Research Programme laboratory's COVID-19 response. Wellcome Open Research, 0, 6, 205.	1.8	0
35	Geographic-genetic analysis of Plasmodium falciparum parasite populations from surveys of primary school children in Western Kenya. Wellcome Open Research, 0, 2, 29.	1.8	10
36	Spatio-temporal distribution of antimalarial drug resistant gene mutations in a Plasmodium falciparum parasite population from Kilifi, Kenya: A 25-year retrospective study. Wellcome Open Research, 0, 7, 45.	1.8	8