

Wonderful T Choga

List of Publications by Citations

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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.

28

papers

320

citations

6

h-index

17

g-index

39

ext. papers

1,092

ext. citations

9.5

avg, IF

1.18

L-index

#	Paper	IF	Citations
28	Rapid epidemic expansion of the SARS-CoV-2 Omicron variant in southern Africa.. <i>Nature</i> , 2022 ,	50.4	205
27	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
26	Rapid epidemic expansion of the SARS-CoV-2 Omicron variant in southern Africa. <i>Nature</i> ,	50.4	20
25	Chronic and Occult Hepatitis B Virus Infection in Pregnant Women in Botswana. <i>Genes</i> , 2018 , 9,	4.2	13
24	Rapid epidemic expansion of the SARS-CoV-2 Omicron variant in southern Africa		12
23	The Impact of Human Pegivirus on CD4 Cell Count in HIV-Positive Persons in Botswana. <i>Open Forum Infectious Diseases</i> , 2017 , 4, ofx222	1	8
22	Molecular Characterization of Near Full-Length Genomes of Hepatitis B Virus Isolated from Predominantly HIV Infected Individuals in Botswana. <i>Genes</i> , 2018 , 9,	4.2	6
21	Molecular characterization of hepatitis B virus in blood donors in Botswana. <i>Virus Genes</i> , 2019 , 55, 33-42.	2.3	5
20	Comparison of an in-house 'home-brew' and commercial ViroSeq integrase genotyping assays on HIV-1 subtype C samples. <i>PLoS ONE</i> , 2019 , 14, e0224292	3.7	4
19	In Silico Analysis of Hepatitis B Virus Occult Associated Mutations in Botswana Using a Novel Algorithm. <i>Genes</i> , 2018 , 9,	4.2	3
18	In Silico Prediction of Human Leukocytes Antigen (HLA) Class II Binding Hepatitis B Virus (HBV) Peptides in Botswana. <i>Viruses</i> , 2020 , 12,	6.2	2
17	The distribution of hepatitis B virus surface antigen polymorphisms at positions associated with vaccine escape. <i>Journal of Medical Virology</i> , 2020 , 92, 3336	19.7	2
16	Incidence of hepatitis B virus infection among human immunodeficiency virus-infected treatment naïve adults in Botswana. <i>Medicine (United States)</i> , 2020 , 99, e19341	1.8	2
15	A25 Impact of polymorphism in the hepatitis B surface gene on human leukocyte antigen (HLA) class II. <i>Virus Evolution</i> , 2019 , 5,	3.7	1
14	Molecular characterization of hepatitis C virus in liver disease patients in Botswana: a retrospective cross-sectional study. <i>BMC Infectious Diseases</i> , 2019 , 19, 875	4	1
13	Hepatitis B virus prevalence and vaccine antibody titers in children HIV exposed but uninfected in Botswana. <i>PLoS ONE</i> , 2020 , 15, e0237252	3.7	1
12	Genetic diversity in L1 ORF of human papillomavirus in women with cervical cancer with and without human immunodeficiency virus in Botswana and Kenya.. <i>BMC Infectious Diseases</i> , 2022 , 22, 95	4	0

- 11 Insilico Functional Analysis of Genome-Wide Dataset From 17,000 Individuals Identifies Candidate Malaria Resistance Genes Enriched in Malaria Pathogenic Pathways. *Frontiers in Genetics*, **2021**, 12, 676960⁴⁵
- 10 Comparison of an in-house home-brew and commercial ViroSeq integrase genotyping assays on HIV-1 subtype C samples **2019**, 14, e0224292
- 9 Comparison of an in-house home-brew and commercial ViroSeq integrase genotyping assays on HIV-1 subtype C samples **2019**, 14, e0224292
- 8 Comparison of an in-house home-brew and commercial ViroSeq integrase genotyping assays on HIV-1 subtype C samples **2019**, 14, e0224292
- 7 Comparison of an in-house home-brew and commercial ViroSeq integrase genotyping assays on HIV-1 subtype C samples **2019**, 14, e0224292
- 6 Hepatitis B virus prevalence and vaccine antibody titers in children HIV exposed but uninfected in Botswana **2020**, 15, e0237252
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