

# Alfred Ssekagiri

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5529907/publications.pdf>

Version: 2024-02-01

10  
papers

50  
citations

1936888

4  
h-index

1719596

7  
g-index

10  
all docs

10  
docs citations

10  
times ranked

103  
citing authors

#	ARTICLE	IF	CITATIONS
1	African Genomic Medicine Portal: A Web Portal for Biomedical Applications. <i>Journal of Personalized Medicine</i> , 2022, 12, 265.	1.1	0
2	H3ABioNet genomic medicine and microbiome data portals hackathon proceedings. <i>Database: the Journal of Biological Databases and Curation</i> , 2021, 2021, .	1.4	7
3	Phylogenetic Networks and Parameters Inferred from HIV Nucleotide Sequences of High-Risk and General Population Groups in Uganda: Implications for Epidemic Control. <i>Viruses</i> , 2021, 13, 970.	1.5	5
4	Circulating levels of angiogenic factors and their association with preeclampsia among pregnant women at Mulago National Referral Hospital in Uganda. <i>PLoS ONE</i> , 2021, 16, e0251227.	1.1	5
5	Mosquito-borne arboviruses in Uganda: history, transmission and burden. <i>Journal of General Virology</i> , 2021, 102, .	1.3	2
6	Mosquito-borne arboviruses in Uganda: history, transmission and burden. <i>Journal of General Virology</i> , 2021, 102, .	1.3	0
7	Employing phylogenetic tree shape statistics to resolve the underlying host population structure. <i>BMC Bioinformatics</i> , 2021, 22, 546.	1.2	3
8	The Molecular Epidemiology and Transmission Dynamics of HIV Type 1 in a General Population Cohort in Uganda. <i>Viruses</i> , 2020, 12, 1283.	1.5	4
9	Diverse vaginal microbiome was associated with pro-inflammatory vaginal milieu among pregnant women in Uganda. <i>Human Microbiome Journal</i> , 2020, 18, 100076.	3.8	7
10	Phylogenetic and Demographic Characterization of Directed HIV-1 Transmission Using Deep Sequences from High-Risk and General Population Cohorts/Groups in Uganda. <i>Viruses</i> , 2020, 12, 331.	1.5	17