

# Ivan Sserwadda

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1334153/publications.pdf>

Version: 2024-02-01

15  
papers

145  
citations

1477746

6  
h-index

1281420

11  
g-index

18  
all docs

18  
docs citations

18  
times ranked

236  
citing authors

#	ARTICLE	IF	CITATIONS
1	blaVIM- and blaOXA-mediated carbapenem resistance among <i>Acinetobacter baumannii</i> and <i>Pseudomonas aeruginosa</i> isolates from the Mulago hospital intensive care unit in Kampala, Uganda. <i>BMC Infectious Diseases</i> , 2019, 19, 853.	1.3	29
2	Microbial contaminants isolated from items and work surfaces in the post- operative ward at Kawolo general hospital, Uganda. <i>BMC Infectious Diseases</i> , 2018, 18, 68.	1.3	27
3	Role of genomics literacy in reducing the burden of common genetic diseases in Africa. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2019, 7, e00776.	0.6	16
4	Human Genomic Loci Important in Common Infectious Diseases: Role of High-Throughput Sequencing and Genome-Wide Association Studies. <i>Canadian Journal of Infectious Diseases and Medical Microbiology</i> , 2018, 2018, 1-9.	0.7	10
5	Investigating colistin drug resistance: The role of high-throughput sequencing and bioinformatics. <i>F1000Research</i> , 2019, 8, 150.	0.8	9
6	Genomics and bioinformatics capacity in Africa: no continent is left behind. <i>Genome</i> , 2021, 64, 503-513.	0.9	8
7	Bioinformatics mentorship in a resource limited setting. <i>Briefings in Bioinformatics</i> , 2022, 23, .	3.2	8
8	First report of whole-genome analysis of an extensively drug-resistant <i>Mycobacterium tuberculosis</i> clinical isolate with bedaquiline, linezolid and clofazimine resistance from Uganda. <i>Antimicrobial Resistance and Infection Control</i> , 2022, 11, 68.	1.5	8
9	Gastrointestinal Tract Colonization Rate of Extended-Spectrum Beta-Lactamase-Producing Gram-Negative Bacteria and Associated Factors Among Orthopaedic Patients in a Tertiary Hospital in Tanzania: Implications for Infection Prevention. <i>Infection and Drug Resistance</i> , 2021, Volume 14, 1733-1745.	1.1	6
10	rMAP: the Rapid Microbial Analysis Pipeline for ESKAPE bacterial group whole-genome sequence data. <i>Microbial Genomics</i> , 2021, 7, .	1.0	6
11	Transmission Dynamics of Antimicrobial Resistance at a National Referral Hospital in Uganda. <i>American Journal of Tropical Medicine and Hygiene</i> , 2021, 105, 498-506.	0.6	6
12	Investigating colistin drug resistance: The role of high-throughput sequencing and bioinformatics. <i>F1000Research</i> , 2019, 8, 150.	0.8	6
13	Increasing Antimicrobial Resistance in Surgical Wards at Mulago National Referral Hospital, Uganda, from 2014 to 2018—Cause for Concern?. <i>Tropical Medicine and Infectious Disease</i> , 2021, 6, 82.	0.9	4
14	Whole-genome sequence analysis of <i>Vibrio cholerae</i> from three outbreaks in Uganda, 2014 - 2016. <i>F1000Research</i> , 0, 8, 1340.	0.8	1
15	Re-thinking antimicrobial resistance transmission dynamics: a meta-analysis of cross-sectional studies at referral hospitals in Uganda. <i>F1000Research</i> , 0, 9, 878.	0.8	0