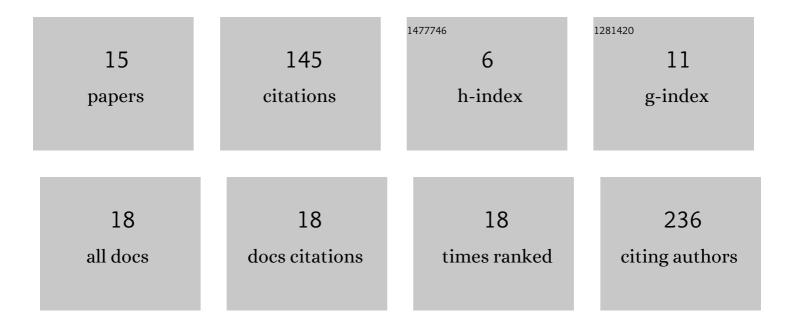
Ivan Sserwadda

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1334153/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	blaVIM- and blaOXA-mediated carbapenem resistance among Acinetobacter baumannii and Pseudomonas aeruginosa isolates from the Mulago hospital intensive care unit in Kampala, Uganda. BMC Infectious Diseases, 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. BMC Infectious Diseases, 2018, 18, 68.	1.3	27
3	Role of genomics literacy in reducing the burden of common genetic diseases in Africa. Molecular Genetics & Genomic Medicine, 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. Canadian Journal of Infectious Diseases and Medical Microbiology, 2018, 2018, 1-9.	0.7	10
5	Investigating colistin drug resistance: The role of high-throughput sequencing and bioinformatics. F1000Research, 2019, 8, 150.	0.8	9
6	Genomics and bioinformatics capacity in Africa: no continent is left behind. Genome, 2021, 64, 503-513.	0.9	8
7	Bioinformatics mentorship in a resource limited setting. Briefings in Bioinformatics, 2022, 23, .	3.2	8
8	First report of whole-genome analysis of an extensively drug-resistant Mycobacterium tuberculosis clinical isolate with bedaquiline, linezolid and clofazimine resistance from Uganda. Antimicrobial Resistance and Infection Control, 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. Infection and Drug Resistance, 2021, Volume 14, 1733-1745.	1.1	6
10	rMAP: the Rapid Microbial Analysis Pipeline for ESKAPE bacterial group whole-genome sequence data. Microbial Genomics, 2021, 7, .	1.0	6
11	Transmission Dynamics of Antimicrobial Resistance at a National Referral Hospital in Uganda. American Journal of Tropical Medicine and Hygiene, 2021, 105, 498-506.	0.6	6
12	Investigating colistin drug resistance: The role of high-throughput sequencing and bioinformatics. F1000Research, 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?. Tropical Medicine and Infectious Disease, 2021, 6, 82.	0.9	4
14	Whole-genome sequence analysis ofÂVibrio cholerae from three outbreaks in Uganda, 2014 - 2016. F1000Research, 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. F1000Research, 0, 9, 878.	0.8	0