

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 | Bioinformatics mentorship in a resource limited setting. <i>Briefings in Bioinformatics</i> , 2022, 23, . | 3.2 | 8 |
| 2 | 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 |
| 3 | Genomics and bioinformatics capacity in Africa: no continent is left behind. <i>Genome</i> , 2021, 64, 503-513. | 0.9 | 8 |
| 4 | 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 |
| 5 | Increasing Antimicrobial Resistance in Surgical Wards at Mulago National Referral Hospital, Uganda, from 2014 to 2018â€”Cause for Concern?. <i>Tropical Medicine and Infectious Disease</i> , 2021, 6, 82. | 0.9 | 4 |
| 6 | rMAP: the Rapid Microbial Analysis Pipeline for ESKAPE bacterial group whole-genome sequence data. <i>Microbial Genomics</i> , 2021, 7, . | 1.0 | 6 |
| 7 | 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 |
| 8 | 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 |
| 9 | Role of genomics literacy in reducing the burden of common genetic diseases in Africa. <i>Molecular Genetics & Genomic Medicine</i> , 2019, 7, e00776. | 0.6 | 16 |
| 10 | Investigating colistin drug resistance: The role of high-throughput sequencing and bioinformatics. <i>F1000Research</i> , 2019, 8, 150. | 0.8 | 9 |
| 11 | Investigating colistin drug resistance: The role of high-throughput sequencing and bioinformatics. <i>F1000Research</i> , 2019, 8, 150. | 0.8 | 6 |
| 12 | 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 |
| 13 | 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 |
| 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 |