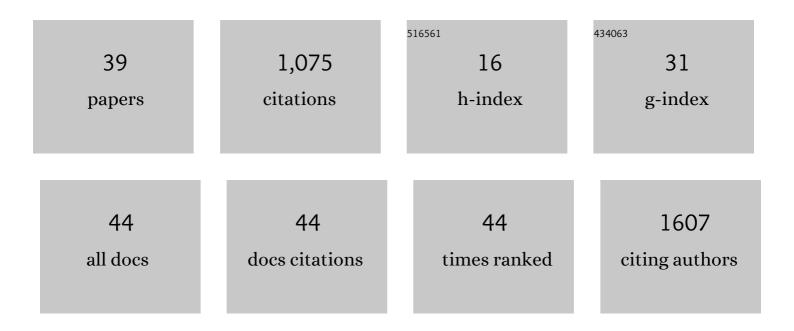
## Henry Kajumbula

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3389135/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Disproportionate Distribution of HBV Genotypes A and D and the Recombinant Genotype D/E in the High and Low HBV Endemic Regions of Uganda: A Wake-Up Call for Regional Specific HBV Management. International Journal of Hepatology, 2022, 2022, 1-15.	0.4	6
2	IL-6 and IL-8 cytokines are associated with elevated prostate-specific antigen levels among patients with adenocarcinoma of the prostate at the Uganda Cancer Institute. Future Oncology, 2022, 18, 661-667.	1.1	7
3	Hepatitis B virus (HBV) serological patterns among the HBsAg negative hospital attendees screened for immunization. Scientific Reports, 2022, 12, 7425.	1.6	5
4	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
5	Implementation of the World Health Organization Global Antimicrobial Resistance Surveillance System in Uganda, 2015-2020: Mixed-Methods Study Using National Surveillance Data. JMIR Public Health and Surveillance, 2021, 7, e29954.	1.2	12
6	Antibiotic resistance profiles and population structure of disease-associated Staphylococcus aureus infecting patients in Fort Portal Regional Referral Hospital, Western Uganda. Microbiology (United) Tj ETQq0 0 C	) rg <b>6.7</b> /0v	erlæck 10 Tf 5
7	Salmonella Bloodstream Infections in Hospitalized Children with Acute Febrile Illness—Uganda, 2016–2019. American Journal of Tropical Medicine and Hygiene, 2021, 105, 37-46.	0.6	12
8	Antibiotic Prevalence Study and Factors Influencing Prescription of WHO Watch Category Antibiotic Ceftriaxone in a Tertiary Care Private Not for Profit Hospital in Uganda. Antibiotics, 2021, 10, 1167.	1.5	3
9	Microbiological Contamination of Mobile Phones and Mobile Phone Hygiene of Final-Year Medical Students in Uganda: A Need for Educational Intervention. Advances in Medical Education and Practice, 2021, Volume 12, 1247-1257.	0.7	2
10	A One Health Approach to Strengthening Antimicrobial Stewardship in Wakiso District, Uganda. Antibiotics, 2020, 9, 764.	1.5	26
11	Whole genome sequences of multi-drug resistant Escherichia coli isolated in a Pastoralist Community of Western Uganda: Phylogenomic changes, virulence and resistant genes. PLoS ONE, 2020, 15, e0231852.	1.1	11
12	Antimicrobial Resistance of Neisseria Gonorrhoeae in a Newly Implemented Surveillance Program in Uganda: Surveillance Report. JMIR Public Health and Surveillance, 2020, 6, e17009.	1.2	14
13	Title is missing!. , 2020, 15, e0231852.		Ο
14	Title is missing!. , 2020, 15, e0231852.		0
15	Title is missing!. , 2020, 15, e0231852.		Ο
16	Title is missing!. , 2020, 15, e0231852.		0
17	Bacteremia in febrile cancer patients in Uganda. BMC Research Notes, 2019, 12, 464.	0.6	28
18	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

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19	A cholera outbreak caused by drinking contaminated river water, Bulambuli District, Eastern Uganda, March 2016. BMC Infectious Diseases, 2019, 19, 516.	1.3	14
20	Burden of fungal asthma in Africa: A systematic review and meta-analysis. PLoS ONE, 2019, 14, e0216568.	1.1	43
21	A Cross-Cutting Approach to Surveillance and Laboratory Capacity as a Platform to Improve Health Security in Uganda. Health Security, 2018, 16, S-76-S-86.	0.9	21
22	Antimicrobial Drug Resistance in Blood Culture Isolates at a Tertiary Hospital, Uganda. Emerging Infectious Diseases, 2018, 24, 174-175.	2.0	35
23	Multidrug resistance among Escherichia coli and Klebsiella pneumoniae carried in the gut of out-patients from pastoralist communities of Kasese district, Uganda. PLoS ONE, 2018, 13, e0200093.	1.1	27
24	A large and persistent outbreak of typhoid fever caused by consuming contaminated water and street-vended beverages: Kampala, Uganda, January – June 2015. BMC Public Health, 2017, 17, 23.	1.2	56
25	Platelet transfusion therapy in subâ€Saharan Africa: bacterial contamination, recipient characteristics, and acute transfusion reactions. Transfusion, 2016, 56, 1951-1959.	0.8	22
26	Human Immune Response Varies by the Degree of Relative Cryptococcal Antigen Shedding. Open Forum Infectious Diseases, 2016, 3, ofv194.	0.4	18
27	Characteristics of pyogenic odontogenic infection in patients attending Mulago Hospital, Uganda: a cross-sectional study. BMC Microbiology, 2015, 15, 46.	1.3	20
28	Regulation of medical diagnostics and medical devices in the East African community partner states. BMC Health Services Research, 2014, 14, 524.	0.9	50
29	Antimicrobial resistance in hospitalized surgical patients: a silently emerging public health concern in Uganda. BMC Research Notes, 2013, 6, 298.	0.6	90
30	External ocular surface bacterial isolates and their antimicrobial susceptibility patterns among pre-operative cataract patients at Mulago National Hospital in Kampala, Uganda. BMC Ophthalmology, 2013, 13, 71.	0.6	41
31	Cryptococcal Genotype Influences Immunologic Response and Human Clinical Outcome after Meningitis. MBio, 2012, 3, .	1.8	79
32	Targeting wild-type Erythrocyte receptors for Plasmodium falciparum and vivax Merozoites by Zinc Finger Nucleases In- silico: Towards a Genetic Vaccine against Malaria. Genetic Vaccines and Therapy, 2012, 10, 8.	1.5	2
33	Nasopharyngeal carriage rate of Streptococcus pneumoniae in Ugandan children with Sickle Cell Disease. BMC Research Notes, 2012, 5, 28.	0.6	14
34	Country Leadership and Policy Are Critical Factors for Implementing Laboratory Accreditation in Developing Countries. American Journal of Clinical Pathology, 2010, 134, 381-387.	0.4	9
35	In silico evidence for the species-specific conservation of mosquito retroposons: implications as a molecular biomarker. Theoretical Biology and Medical Modelling, 2009, 6, 14.	2.1	3
36	Identification of restriction endonuclease with potential ability to cleave the HSV-2 genome: Inherent potential for biosynthetic versus live recombinant microbicides. Theoretical Biology and Medical Modelling, 2008, 5, 18.	2.1	7

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37	Evaluation of global clustering patterns and strain variation over an extended ORF26 gene locus from Kaposi's sarcoma herpesvirus. Journal of Clinical Virology, 2007, 40, 19-25.	1.6	15
38	Ugandan Kaposi's Sarcoma-Associated Herpesvirus Phylogeny: Evidence for Cross-Ethnic Transmission of Viral Subtypes. Intervirology, 2006, 49, 133-143.	1.2	43
39	A novel viral mechanism for dysregulation of β-catenin in Kaposi's sarcoma–associated herpesvirus latency. Nature Medicine, 2003, 9, 300-306.	15.2	301