David Patrick Kateete

List of Publications by Year in descending order

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567144 610775 47 720 15 24 citations h-index g-index papers 63 63 63 1286 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Prevalence and Antimicrobial Susceptibility Patterns of Bacteria from Milkmen and Cows with Clinical Mastitis in and around Kampala, Uganda. PLoS ONE, 2013, 8, e63413.	1.1	71
2	Diversity of vaginal microbiota in sub-Saharan Africa and its effects on HIV transmission and prevention. American Journal of Obstetrics and Gynecology, 2019, 220, 155-166.	0.7	58
3	CA-MRSA and HA-MRSA coexist in community and hospital settings in Uganda. Antimicrobial Resistance and Infection Control, 2019, 8, 94.	1.5	51
4	Characteristics and outcomes of admitted patients infected with SARS-CoV-2 in Uganda. BMJ Open Respiratory Research, 2020, 7, e000646.	1.2	42
5	Molecular Characterization of Staphylococcus aureus from Patients with Surgical Site Infections at Mulago Hospital in Kampala, Uganda. PLoS ONE, 2013, 8, e66153.	1.1	38
6	Whole-Exome Sequencing Reveals Uncaptured Variation and Distinct Ancestry in the Southern African Population of Botswana. American Journal of Human Genetics, 2018, 102, 731-743.	2.6	38
7	Genotypic diversity among multidrug resistant Pseudomonas aeruginosa and Acinetobacter species at Mulago Hospital in Kampala, Uganda. BMC Research Notes, 2017, 10, 284.	0.6	32
8	Prevalence and patterns of rifampicin and isoniazid resistance conferring mutations in Mycobacterium tuberculosis isolates from Uganda. PLoS ONE, 2018, 13, e0198091.	1.1	31
9	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
10	Determination of circulating Mycobacterium tuberculosis strains and transmission patterns among pulmonary TB patients in Kawempe municipality, Uganda, using MIRU-VNTR. BMC Research Notes, 2011, 4, 280.	0.6	24
11	Prevalence of virulence determinants in Staphylococcus epidermidis from ICU patients in Kampala, Uganda. Journal of Infection in Developing Countries, 2012, 6, 242-250.	0.5	23
12	The T2 Mycobacterium tuberculosis Genotype, Predominant in Kampala, Uganda, Shows Negative Correlation with Antituberculosis Drug Resistance. Antimicrobial Agents and Chemotherapy, 2014, 58, 3853-3859.	1.4	21
13	Comparison of transformation frequencies among selected Streptococcus pneumoniae serotypes. International Journal of Antimicrobial Agents, 2010, 36, 124-128.	1.1	17
14	Isolation of Mycobacterium avium subspecies paratuberculosis from Ugandan cattle and strain differentiation using optimised DNA typing techniques. BMC Veterinary Research, 2012, 8, 99.	0.7	17
15	Rates of HIV-1 virological suppression and patterns of acquired drug resistance among fisherfolk on first-line antiretroviral therapy in Uganda. Journal of Antimicrobial Chemotherapy, 2019, 74, 3021-3029.	1.3	16
16	Rhomboids of Mycobacteria: Characterization Using an aarA Mutant of Providencia stuartii and Gene Deletion in Mycobacterium smegmatis. PLoS ONE, 2012, 7, e45741.	1.1	16
17	Nasopharyngeal carriage, spa types and antibiotic susceptibility profiles of Staphylococcus aureus from healthy children less than 5 years in Eastern Uganda. BMC Infectious Diseases, 2019, 19, 1023.	1.3	15
18	The Collaborative African Genomics Network (CAfGEN): Applying Genomic technologies to probe host factors important to the progression of HIV and HIV-tuberculosis infection in sub-Saharan Africa. AAS Open Research, 2018, 1, 3.	1.5	15

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19	Evaluation of Capilia TB assay for rapid identification of Mycobacterium tuberculosis complex in BACTEC MGIT 960 and BACTEC 9120 blood cultures. BMC Research Notes, 2012, 5, 44.	0.6	14
20	Species, antibiotic susceptibility profiles and van gene frequencies among enterococci isolated from patients at Mulago National Referral Hospital in Kampala, Uganda. BMC Infectious Diseases, 2019, 19, 486.	1.3	14
21	Frequency and patterns of second-line resistance conferring mutations among MDR-TB isolates resistant to a second-line drug from eSwatini, Somalia and Uganda (2014–2016). BMC Pulmonary Medicine, 2019, 19, 124.	0.8	13
22	Impact of vitamin D status and cathelicidin antimicrobial peptide on adults with active pulmonary TB globally: A systematic review and meta-analysis. PLoS ONE, 2021, 16, e0252762.	1.1	13
23	High prevalence of phenotypic pyrazinamide resistance and its association with pncA gene mutations in Mycobacterium tuberculosis isolates from Uganda. PLoS ONE, 2020, 15, e0232543.	1.1	12
24	The Collaborative African Genomics Network (CAfGEN): Applying Genomic technologies to probe host factors important to the progression of HIV and HIV-tuberculosis infection in sub-Saharan Africa. AAS Open Research, 2018, 1, 3.	1.5	10
25	Species and drug susceptibility profiles of staphylococci isolated from healthy children in Eastern Uganda. PLoS ONE, 2020, 15, e0229026.	1.1	9
26	Tuberculosis resistance-conferring mutations with fitness cost among HIV-positive individuals in Uganda. International Journal of Tuberculosis and Lung Disease, 2017, 21, 531-536.	0.6	8
27	Biobanking: Strengthening Uganda's Rapid Response to COVID-19 and Other Epidemics. Biopreservation and Biobanking, 2022, 20, 238-243.	0.5	8
28	Incremental Yield of Serial Sputum Cultures for Diagnosis of Tuberculosis among HIV Infected Smear Negative Pulmonary TB Suspects in Kampala, Uganda. PLoS ONE, 2012, 7, e37650.	1.1	7
29	Performance and cost-effectiveness of a pooled testing strategy for SARS-CoV-2 using real-time polymerase chain reaction in Uganda. International Journal of Infectious Diseases, 2021, 113, 355-358.	1.5	7
30	Acute hypoxaemic respiratory failure in a low-income country: a prospective observational study of hospital prevalence and mortality. BMJ Open Respiratory Research, 2020, 7, e000719.	1.2	7
31	Diarrhoeagenic Escherichia coli isolated from children with acute diarrhoea at Rakai hospital, Southern Uganda. African Health Sciences, 2022, 22, 581-8.	0.3	6
32	The Mycobacterium tuberculosisUganda II family and resistance to first-line anti-tuberculosis drugs in Uganda. BMC Infectious Diseases, 2014, 14, 703.	1.3	5
33	Application of antibodies to recombinant heat shock protein 70 in immunohistochemical diagnosis of mycobacterium avium subspecies paratuberculosis in tissues of naturally infected cattle. Irish Veterinary Journal, 2017, 70, 10.	0.8	5
34	Sputum microbiota profiles of treatment-na \tilde{A} -ve TB patients in Uganda before and during first-line therapy. Scientific Reports, 2021, 11, 24486.	1.6	5
35	Mycobacterium tuberculosis Uganda II is more susceptible to rifampicin and isoniazid compared to Beijing and Delhi/CAS families. BMC Infectious Diseases, 2016, 16, 173.	1.3	4
36	Whole-genome analysis to determine the rate and patterns of intra-subtype reassortment among influenza type-A viruses in Africa. Virus Evolution, 2022, 8, veac005.	2,2	4

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37	Hypovitaminosis D among newly diagnosed pulmonary TB patients and their household contacts in Uganda. Scientific Reports, 2022, 12, 5296.	1.6	4
38	Detection of Mycobacterium tuberculosis DNA in CD34+ peripheral blood mononuclear cells of Ugandan adults with latent infection: a cross-sectional and nested prospective study. AAS Open Research, 2020, 3, 34.	1.5	3
39	Prevalence of plasmid-mediated AmpC beta-lactamases in Enterobacteria isolated from urban and rural folks in Uganda. AAS Open Research, 2020, 3, 62.	1.5	3
40	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 (Org ®.7 √Ov	verloack 10 Tf 5
41	Unique circulating microRNA profiles in epidemic Kaposi's sarcoma. Non-coding RNA Research, 2022, 7, 114-122.	2.4	1
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