

Mary Kathryn Grabowski

List of Publications by Year in descending order

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Version: 2024-02-01

38
papers

1,142
citations

471061

17
h-index

454577

30
g-index

42
all docs

42
docs citations

42
times ranked

2243
citing authors

#	ARTICLE	IF	CITATIONS
1	Hypertension and Socioeconomic Status in South Central Uganda: A Population-Based Cohort Study. <i>Global Heart</i> , 2022, 17, 3.	0.9	6
2	A highly virulent variant of HIV-1 circulating in the Netherlands. <i>Science</i> , 2022, 375, 540-545.	6.0	39
3	Phylogenetic estimation of the viral fitness landscape of HIV-1 set-point viral load. <i>Virus Evolution</i> , 2022, 8, veac022.	2.2	1
4	Inferring the Sources of HIV Infection in Africa from Deep-Sequence Data with Semi-Parametric Bayesian Poisson Flow Models. <i>Journal of the Royal Statistical Society Series C: Applied Statistics</i> , 2022, 71, 517-540.	0.5	2
5	The Mirasol Evaluation of Reduction in Infections Trial (MERIT): study protocol for a randomized controlled clinical trial. <i>Trials</i> , 2022, 23, 257.	0.7	7
6	Curating the Evidence About COVID-19 for Frontline Public Health and Clinical Care: The Novel Coronavirus Research Compendium. <i>Public Health Reports</i> , 2022, 137, 197-202.	1.3	2
7	In-person schooling and associated COVID-19 risk in the United States over spring semester 2021. <i>Science Advances</i> , 2022, 8, eabm9128.	4.7	10
8	Short Communication: Validation of the Asante HIV-1 Rapid Recency Assay for Detection of Recent HIV-1 Infections in Uganda. <i>AIDS Research and Human Retroviruses</i> , 2021, 37, 893-896.	0.5	7
9	Household COVID-19 risk and in-person schooling. <i>Science</i> , 2021, 372, 1092-1097.	6.0	162
10	Prescription Antibiotic Use Among the US population 1999â€“2018: National Health and Nutrition Examination Surveys. <i>Open Forum Infectious Diseases</i> , 2021, 8, ofab224.	0.4	3
11	Differential Performance of CoronaCHEK SARS-CoV-2 Lateral Flow Antibody Assay by Geographic Origin of Samples. <i>Journal of Clinical Microbiology</i> , 2021, 59, e0083721.	1.8	10
12	Mobile Ecological Momentary Assessment and Intervention and Health Behavior Change Among Adults in Rakai, Uganda: Pilot Randomized Controlled Trial. <i>JMIR Formative Research</i> , 2021, 5, e22693.	0.7	5
13	<i>Clostridioides difficile</i> Prevalence in the United States: National Inpatient Sample, 2016 to 2018. <i>Open Forum Infectious Diseases</i> , 2021, 8, ofab409.	0.4	3
14	Prevalence and Predictors of Persistent Human Immunodeficiency Virus Viremia and Viral Rebound After Universal Test and Treat: A Population-Based Study. <i>Journal of Infectious Diseases</i> , 2021, 223, 1150-1160.	1.9	16
15	Seroprevalence of <i>Chlamydia trachomatis</i> Among Female Adults in the United States: The National Health and Nutrition Examination Surveys. <i>Clinical Infectious Diseases</i> , 2021, 73, e629-e637.	2.9	10
16	Cytokine and Chemokine Levels in Coronavirus Disease 2019 Convalescent Plasma. <i>Open Forum Infectious Diseases</i> , 2021, 8, ofaa574.	0.4	41
17	Declining HIV incidence in sub-Saharan Africa: a systematic review and meta-analysis of empiric data. <i>Journal of the International AIDS Society</i> , 2021, 24, e25818.	1.2	32
18	Access to and safety of COVID-19 convalescent plasma in the United States Expanded Access Program: A national registry study. <i>PLoS Medicine</i> , 2021, 18, e1003872.	3.9	43

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19	Phylogenetic Analysis of Human Immunodeficiency Virus from People Who Inject Drugs in Indonesia, Ukraine, and Vietnam: HPTN 074. <i>Clinical Infectious Diseases</i> , 2020, 71, 1836-1846.	2.9	6
20	The engines of SARS-CoV-2 spread. <i>Science</i> , 2020, 370, 406-407.	6.0	100
21	Impact of male circumcision on risk of HIV infection in men in a changing epidemic context – systematic review and meta-analysis. <i>Journal of the International AIDS Society</i> , 2020, 23, e25490.	1.2	20
22	Quantifying HIV transmission flow between high-prevalence hotspots and surrounding communities: a population-based study in Rakai, Uganda. <i>Lancet HIV</i> , 2020, 7, e173-e183.	2.1	59
23	HIV-1 Subtype Distribution and Diversity Over 18 Years in Rakai, Uganda. <i>AIDS Research and Human Retroviruses</i> , 2020, 36, 522-526.	0.5	3
24	Impact of combination HIV interventions on HIV incidence in hyperendemic fishing communities in Uganda: a prospective cohort study. <i>Lancet HIV</i> , 2019, 6, e680-e687.	2.1	52
25	Characteristics Associated With Human Immunodeficiency Virus Transmission Networks Involving Adolescent Girls and Young Women in Human Immunodeficiency Virus Prevention Trials Network 068 Study. <i>Sexually Transmitted Diseases</i> , 2019, 46, e46-e49.	0.8	1
26	Inferring HIV-1 transmission networks and sources of epidemic spread in Africa with deep-sequence phylogenetic analysis. <i>Nature Communications</i> , 2019, 10, 1411.	5.8	50
27	Effect of HIV Subtype and Antiretroviral Therapy on HIV-Associated Neurocognitive Disorder Stage in Rakai, Uganda. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2019, 81, 216-223.	0.9	21
28	Short Communication: The Interaction of HIV Set Point Viral Load and Subtype on Disease Progression. <i>AIDS Research and Human Retroviruses</i> , 2019, 35, 49-51.	0.5	11
29	Increases in Human Papillomavirus Vaccination Among Adolescent and Young Adult Males in the United States, 2011–2016. <i>Journal of Infectious Diseases</i> , 2018, 218, 109-113.	1.9	17
30	Easy and accurate reconstruction of whole HIV genomes from short-read sequence data with shiver. <i>Virus Evolution</i> , 2018, 4, vey007.	2.2	64
31	Ethical considerations in global HIV phylogenetic research. <i>Lancet HIV</i> , 2018, 5, e656-e666.	2.1	39
32	HIV-1 diversity among young women in rural South Africa: HPTN 068. <i>PLoS ONE</i> , 2018, 13, e0198999.	1.1	12
33	HIV-1 Full-Genome Phylogenetics of Generalized Epidemics in Sub-Saharan Africa: Impact of Missing Nucleotide Characters in Next-Generation Sequences. <i>AIDS Research and Human Retroviruses</i> , 2017, 33, 1083-1098.	0.5	18
34	Phylogenetic insights into age-disparate partnerships and HIV. <i>Lancet HIV</i> , 2017, 4, e8-e9.	2.1	8
35	HIV Prevention Efforts and Incidence of HIV in Uganda. <i>New England Journal of Medicine</i> , 2017, 377, 2154-2166.	13.9	163
36	Viral genetic variation accounts for a third of variability in HIV-1 set-point viral load in Europe. <i>PLoS Biology</i> , 2017, 15, e2001855.	2.6	38

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37	Trends in the Mechanistic and Dynamic Modeling of Infectious Diseases. Current Epidemiology Reports, 2016, 3, 212-222.	1.1	27
38	Measuring Spatial Dependence for Infectious Disease Epidemiology. PLoS ONE, 2016, 11, e0155249.	1.1	29