

Rachel J Sippy

List of Publications by Year in descending order

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Version: 2024-02-01

24
papers

1,000
citations

758635

12
h-index

610482

24
g-index

30
all docs

30
docs citations

30
times ranked

1539
citing authors

#	ARTICLE	IF	CITATIONS
1	Chronic kidney disease in Ecuador: An epidemiological and health system analysis of an emerging public health crisis. <i>PLoS ONE</i> , 2022, 17, e0265395.	1.1	3
2	The 2018–2019 weak El Niño: Predicting the risk of a dengue outbreak in Machala, Ecuador. <i>International Journal of Climatology</i> , 2021, 41, 3813-3823.	1.5	9
3	Climate predicts geographic and temporal variation in mosquito-borne disease dynamics on two continents. <i>Nature Communications</i> , 2021, 12, 1233.	5.8	49
4	Assessing critical gaps in COVID-19 testing capacity: the case of delayed results in Ecuador. <i>BMC Public Health</i> , 2021, 21, 637.	1.2	32
5	Recommended reporting items for epidemic forecasting and prediction research: The EPIFORGE 2020 guidelines. <i>PLoS Medicine</i> , 2021, 18, e1003793.	3.9	42
6	Household and climate factors influence <i>Aedes aegypti</i> presence in the arid city of Huaquillas, Ecuador. <i>PLoS Neglected Tropical Diseases</i> , 2021, 15, e0009931.	1.3	7
7	A decade of arbovirus emergence in the temperate southern cone of South America: dengue, <i>Aedes aegypti</i> and climate dynamics in Córdoba, Argentina. <i>Heliyon</i> , 2020, 6, e04858.	1.4	8
8	Identification and evaluation of epidemic prediction and forecasting reporting guidelines: A systematic review and a call for action. <i>Epidemics</i> , 2020, 33, 100400.	1.5	10
9	Time to reality check the promises of machine learning-powered precision medicine. <i>The Lancet Digital Health</i> , 2020, 2, e677-e680.	5.9	126
10	Key Findings and Comparisons From Analogous Case-Cluster Studies for Dengue Virus Infection Conducted in Machala, Ecuador, and Kamphaeng Phet, Thailand. <i>Frontiers in Public Health</i> , 2020, 8, 2.	1.3	2
11	Severity Index for Suspected Arbovirus (SISA): Machine learning for accurate prediction of hospitalization in subjects suspected of arboviral infection. <i>PLoS Neglected Tropical Diseases</i> , 2020, 14, e0007969.	1.3	16
12	The origins of dengue and chikungunya viruses in Ecuador following increased migration from Venezuela and Colombia. <i>BMC Evolutionary Biology</i> , 2020, 20, 31.	3.2	15
13	Thermal biology of mosquito-borne disease. <i>Ecology Letters</i> , 2019, 22, 1690-1708.	3.0	349
14	Seasonal and geographic variation in insecticide resistance in <i>Aedes aegypti</i> in southern Ecuador. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007448.	1.3	21
15	Seasonal patterns of dengue fever in rural Ecuador: 2009-2016. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007360.	1.3	12
16	Effects of Political Instability in Venezuela on Malaria Resurgence at Ecuador–Peru Border, 2018. <i>Emerging Infectious Diseases</i> , 2019, 25, 834-836.	2.0	47
17	Prioritization of family member sequencing for the detection of rare variants. <i>BMC Proceedings</i> , 2016, 10, 227-231.	1.8	1
18	DNA Topology and the Initiation of Virus DNA Packaging. <i>PLoS ONE</i> , 2016, 11, e0154785.	1.1	1

#	ARTICLE	IF	CITATIONS
19	Genetics of critical contacts and clashes in the DNA packaging specificities of bacteriophages ϕ and 21. <i>Virology</i> , 2015, 476, 115-123.	1.1	8
20	Genetic Diversity and Antimicrobial Susceptibility of <i>Campylobacter jejuni</i> Isolates Associated with Sheep Abortion in the United States and Great Britain. <i>Journal of Clinical Microbiology</i> , 2014, 52, 1853-1861.	1.8	41
21	Molecular Evidence for Zoonotic Transmission of an Emergent, Highly Pathogenic <i>Campylobacter jejuni</i> Clone in the United States. <i>Journal of Clinical Microbiology</i> , 2012, 50, 680-687.	1.8	98
22	Critical Role of LuxS in the Virulence of <i>Campylobacter jejuni</i> in a Guinea Pig Model of Abortion. <i>Infection and Immunity</i> , 2012, 80, 585-593.	1.0	38
23	Development of a Loop-Mediated Isothermal Amplification Assay for Rapid, Sensitive and Specific Detection of a <i>Campylobacter jejuni</i> Clone. <i>Journal of Veterinary Medical Science</i> , 2012, 74, 591-596.	0.3	13
24	Occurrence and molecular analysis of <i>Campylobacter</i> in wildlife on livestock farms. <i>Veterinary Microbiology</i> , 2012, 157, 369-375.	0.8	45