

John R Giles

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2851952/publications.pdf>

Version: 2024-02-01

11
papers

2,677
citations

1162889

8
h-index

1281743

11
g-index

14
all docs

14
docs citations

14
times ranked

1891
citing authors

#	ARTICLE	IF	CITATIONS
1	Global prevalence and burden of depressive and anxiety disorders in 204 countries and territories in 2020 due to the COVID-19 pandemic. <i>Lancet, The</i> , 2021, 398, 1700-1712.	6.3	2,234
2	The use of mobile phone data to inform analysis of COVID-19 pandemic epidemiology. <i>Nature Communications</i> , 2020, 11, 4961.	5.8	246
3	Synchronous shedding of multiple bat paramyxoviruses coincides with peak periods of Hendra virus spillover. <i>Emerging Microbes and Infections</i> , 2019, 8, 1314-1323.	3.0	49
4	Environmental drivers of spatiotemporal foraging intensity in fruit bats and implications for Hendra virus ecology. <i>Scientific Reports</i> , 2018, 8, 9555.	1.6	33
5	Models of Eucalypt phenology predict bat population flux. <i>Ecology and Evolution</i> , 2016, 6, 7230-7245.	0.8	30
6	The duration of travel impacts the spatial dynamics of infectious diseases. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 22572-22579.	3.3	27
7	Ecology and Geography of Plague Transmission Areas in Northeastern Brazil. <i>PLoS Neglected Tropical Diseases</i> , 2011, 5, e925.	1.3	22
8	Optimizing noninvasive sampling of a zoonotic bat virus. <i>Ecology and Evolution</i> , 2021, 11, 12307-12321.	0.8	13
9	Characterizing human mobility patterns in rural settings of sub-Saharan Africa. <i>ELife</i> , 2021, 10, .	2.8	8
10	Sample size calculation for phylogenetic case linkage. <i>PLoS Computational Biology</i> , 2021, 17, e1009182.	1.5	7
11	Trip duration drives shift in travel network structure with implications for the predictability of spatial disease spread. <i>PLoS Computational Biology</i> , 2021, 17, e1009127.	1.5	4