

Simon J Anthony

List of Publications by Year in descending order

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Version: 2024-02-01

28
papers

2,657
citations

361413

20
h-index

454955

30
g-index

35
all docs

35
docs citations

35
times ranked

4634
citing authors

#	ARTICLE	IF	CITATIONS
1	Coronavirus and Paramyxovirus Shedding by Bats in a Cave and Buildings in Ethiopia. <i>EcoHealth</i> , 2022, 19, 216-232.	2.0	3
2	Classification of new morbillivirus and jeilongvirus sequences from bats sampled in Brazil and Malaysia. <i>Archives of Virology</i> , 2022, 167, 1977-1987.	2.1	11
3	Trypsin Treatment Unlocks Barrier for Zoonotic Bat Coronavirus Infection. <i>Journal of Virology</i> , 2020, 94, .	3.4	162
4	Spillover of ebolaviruses into people in eastern Democratic Republic of Congo prior to the 2018 Ebola virus disease outbreak. <i>One Health Outlook</i> , 2020, 2, 21.	3.4	5
5	2020 taxonomic update for phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. <i>Archives of Virology</i> , 2020, 165, 3023-3072.	2.1	184
6	Nipah virus dynamics in bats and implications for spillover to humans. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 29190-29201.	7.1	119
7	A Virion-Based Assay for Glycoprotein Thermostability Reveals Key Determinants of Filovirus Entry and Its Inhibition. <i>Journal of Virology</i> , 2020, 94, .	3.4	7
8	Serological Evidence for Henipa-like and Filo-like Viruses in Trinidad Bats. <i>Journal of Infectious Diseases</i> , 2020, 221, S375-S382.	4.0	20
9	Hospital-based zoonotic disease surveillance in Bangladesh: design, field data and difficulties. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2019, 374, 20190019.	4.0	8
10	Development of a Human Antibody Cocktail that Deploys Multiple Functions to Confer Pan-Ebolavirus Protection. <i>Cell Host and Microbe</i> , 2019, 25, 39-48.e5.	11.0	83
11	The discovery of Bombali virus adds further support for bats as hosts of ebolaviruses. <i>Nature Microbiology</i> , 2018, 3, 1084-1089.	13.3	283
12	One Health proof of concept: Bringing a transdisciplinary approach to surveillance for zoonotic viruses at the human-wild animal interface. <i>Preventive Veterinary Medicine</i> , 2017, 137, 112-118.	1.9	112
13	Global patterns in coronavirus diversity. <i>Virus Evolution</i> , 2017, 3, vex012.	4.9	310
14	Viral Diversity, Prey Preference, and Bartonella Prevalence in <i>Desmodus rotundus</i> in Guatemala. <i>EcoHealth</i> , 2016, 13, 761-774.	2.0	37
15	Adenovirus and Herpesvirus Diversity in Free-Ranging Great Apes in the Sangha Region of the Republic of Congo. <i>PLoS ONE</i> , 2015, 10, e0118543.	2.5	27
16	Virus hunting. <i>Virology</i> , 2015, 479-480, 194-199.	2.4	38
17	Non-random patterns in viral diversity. <i>Nature Communications</i> , 2015, 6, 8147.	12.8	65
18	A Strategy To Estimate Unknown Viral Diversity in Mammals. <i>MBio</i> , 2013, 4, e00598-13.	4.1	320

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19	Ebola Virus Antibodies in Fruit Bats, Bangladesh. <i>Emerging Infectious Diseases</i> , 2013, 19, 270-273.	4.3	129
20	Bats are a major natural reservoir for hepaciviruses and pegiviruses. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 8194-8199.	7.1	251
21	Surveillance theory applied to virus detection: a case for targeted discovery. <i>Future Virology</i> , 2013, 8, 1201-1206.	1.8	3
22	Pandemic H1N1 Influenza Isolated from Free-Ranging Northern Elephant Seals in 2010 off the Central California Coast. <i>PLoS ONE</i> , 2013, 8, e62259.	2.5	46
23	Identification of a Novel Cetacean Polyomavirus from a Common Dolphin (<i>Delphinus delphis</i>) with Tracheobronchitis. <i>PLoS ONE</i> , 2013, 8, e68239.	2.5	18
24	Zoonotic Viruses Associated with Illegally Imported Wildlife Products. <i>PLoS ONE</i> , 2012, 7, e29505.	2.5	122
25	Saliva Proteins of Vector <i>Culicoides</i> Modify Structure and Infectivity of Bluetongue Virus Particles. <i>PLoS ONE</i> , 2011, 6, e17545.	2.5	28
26	RT-PCR Assays for Seven Serotypes of Epizootic Haemorrhagic Disease Virus & Their Use to Type Strains from the Mediterranean Region and North America. <i>PLoS ONE</i> , 2010, 5, e12782.	2.5	42
27	Evolution and Phylogenetic Analysis of Full-Length VP3 Genes of Eastern Mediterranean Bluetongue Virus Isolates. <i>PLoS ONE</i> , 2009, 4, e6437.	2.5	33
28	Sequence analysis of bluetongue virus serotype 8 from the Netherlands 2006 and comparison to other European strains. <i>Virology</i> , 2008, 377, 308-318.	2.4	172