

Karla Prieto

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1280325/publications.pdf>

Version: 2024-02-01

19
papers

1,520
citations

623734

14
h-index

794594

19
g-index

19
all docs

19
docs citations

19
times ranked

3299
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular Evidence of Sexual Transmission of Ebola Virus. <i>New England Journal of Medicine</i> , 2015, 373, 2448-2454.	27.0	380
2	Genomic epidemiology reveals multiple introductions of Zika virus into the United States. <i>Nature</i> , 2017, 546, 401-405.	27.8	298
3	Taxonomy of the order Mononegavirales: update 2018. <i>Archives of Virology</i> , 2018, 163, 2283-2294.	2.1	153
4	A Multicomponent Animal Virus Isolated from Mosquitoes. <i>Cell Host and Microbe</i> , 2016, 20, 357-367.	11.0	123
5	Evolution and Spread of Ebola Virus in Liberia, 2014–2015. <i>Cell Host and Microbe</i> , 2015, 18, 659-669.	11.0	87
6	Monitoring of Ebola Virus Makona Evolution through Establishment of Advanced Genomic Capability in Liberia. <i>Emerging Infectious Diseases</i> , 2015, 21, 1135-1143.	4.3	79
7	Reduced evolutionary rate in reemerged Ebola virus transmission chains. <i>Science Advances</i> , 2016, 2, e1600378.	10.3	62
8	Medical countermeasures during the 2018 Ebola virus disease outbreak in the North Kivu and Ituri Provinces of the Democratic Republic of the Congo: a rapid genomic assessment. <i>Lancet Infectious Diseases</i> , 2019, 19, 648-657.	9.1	62
9	First report of naturally infected <i>Aedes aegypti</i> with chikungunya virus genotype ECSA in the Americas. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0005630.	3.0	59
10	Genomic Characterization of the Genus Nairovirus (Family Bunyaviridae). <i>Viruses</i> , 2016, 8, 164.	3.3	57
11	Complete Genome Sequences of Five Zika Virus Isolates. <i>Genome Announcements</i> , 2016, 4, .	0.8	40
12	Active Ebola Virus Replication and Heterogeneous Evolutionary Rates in EVD Survivors. <i>Cell Reports</i> , 2018, 22, 1159-1168.	6.4	37
13	Persistence and Intra-Host Genetic Evolution of Zika Virus Infection in Symptomatic Adults: A Special View in the Male Reproductive System. <i>Viruses</i> , 2018, 10, 615.	3.3	30
14	Lassa virus circulating in Liberia: a retrospective genomic characterisation. <i>Lancet Infectious Diseases</i> , 2019, 19, 1371-1378.	9.1	30
15	Chikungunya viruses containing the A226V mutation detected retrospectively in Cameroon form a new geographical subclade. <i>International Journal of Infectious Diseases</i> , 2021, 113, 65-73.	3.3	11
16	Beatrice Hill Virus Represents a Novel Species in the Genus Tibrovirus (Mononegavirales :) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 142 Td	0.8	6
17	Novel viruses in hard ticks collected in the Republic of Korea unveiled by metagenomic high-throughput sequencing analysis. <i>Ticks and Tick-borne Diseases</i> , 2021, 12, 101820.	2.7	4
18	A Department of Defense Laboratory Consortium Approach to Next Generation Sequencing and Bioinformatics Training for Infectious Disease Surveillance in Kenya. <i>Frontiers in Genetics</i> , 2020, 11, 577563.	2.3	1

#	ARTICLE	IF	CITATIONS
19	Complete genomic sequences of Venezuelan equine encephalitis virus subtype IIIID isolates from mosquitoes. Archives of Virology, 2020, 165, 1715-1717.	2.1	1