

# 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	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
2	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
3	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
4	Complete genomic sequences of Venezuelan equine encephalitis virus subtype IIID isolates from mosquitoes. <i>Archives of Virology</i> , 2020, 165, 1715-1717.	2.1	1
5	Lassa virus circulating in Liberia: a retrospective genomic characterisation. <i>Lancet Infectious Diseases</i> , The, 2019, 19, 1371-1378.	9.1	30
6	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> , The, 2019, 19, 648-657.	9.1	62
7	Taxonomy of the order Mononegavirales: update 2018. <i>Archives of Virology</i> , 2018, 163, 2283-2294.	2.1	153
8	Active Ebola Virus Replication and Heterogeneous Evolutionary Rates in EVD Survivors. <i>Cell Reports</i> , 2018, 22, 1159-1168.	6.4	37
9	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
10	Genomic epidemiology reveals multiple introductions of Zika virus into the United States. <i>Nature</i> , 2017, 546, 401-405.	27.8	298
11	Beatrice Hill Virus Represents a Novel Species in the Genus Tibrovirus ( Mononegavirales : ) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	0.8	6
12	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
13	Genomic Characterization of the Genus Nairovirus (Family Bunyaviridae). <i>Viruses</i> , 2016, 8, 164.	3.3	57
14	Reduced evolutionary rate in reemerged Ebola virus transmission chains. <i>Science Advances</i> , 2016, 2, e1600378.	10.3	62
15	Complete Genome Sequences of Five Zika Virus Isolates. <i>Genome Announcements</i> , 2016, 4, .	0.8	40
16	A Multicomponent Animal Virus Isolated from Mosquitoes. <i>Cell Host and Microbe</i> , 2016, 20, 357-367.	11.0	123
17	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
18	Evolution and Spread of Ebola Virus in Liberia, 2014â€“2015. <i>Cell Host and Microbe</i> , 2015, 18, 659-669.	11.0	87

#	ARTICLE	IF	CITATIONS
19	Molecular Evidence of Sexual Transmission of Ebola Virus. New England Journal of Medicine, 2015, 373, 2448-2454.	27.0	380