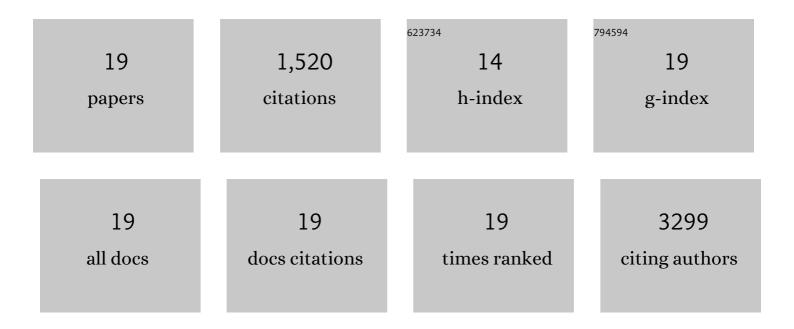
## Karla Prieto

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1280325/publications.pdf Version: 2024-02-01



KADIA DDIETO

#	Article	IF	CITATIONS
1	Molecular Evidence of Sexual Transmission of Ebola Virus. New England Journal of Medicine, 2015, 373, 2448-2454.	27.0	380
2	Genomic epidemiology reveals multiple introductions of Zika virus into the United States. Nature, 2017, 546, 401-405.	27.8	298
3	Taxonomy of the order Mononegavirales: update 2018. Archives of Virology, 2018, 163, 2283-2294.	2.1	153
4	A Multicomponent Animal Virus Isolated from Mosquitoes. Cell Host and Microbe, 2016, 20, 357-367.	11.0	123
5	Evolution and Spread of Ebola Virus in Liberia, 2014–2015. Cell Host and Microbe, 2015, 18, 659-669.	11.0	87
6	Monitoring of Ebola Virus Makona Evolution through Establishment of Advanced Genomic Capability in Liberia. Emerging Infectious Diseases, 2015, 21, 1135-1143.	4.3	79
7	Reduced evolutionary rate in reemerged Ebola virus transmission chains. Science Advances, 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. Lancet Infectious Diseases, The, 2019, 19, 648-657.	9.1	62
9	First report of naturally infected Aedes aegypti with chikungunya virus genotype ECSA in the Americas. PLoS Neglected Tropical Diseases, 2017, 11, e0005630.	3.0	59
10	Genomic Characterization of the Genus Nairovirus (Family Bunyaviridae). Viruses, 2016, 8, 164.	3.3	57
11	Complete Genome Sequences of Five Zika Virus Isolates. Genome Announcements, 2016, 4, .	0.8	40
12	Active Ebola Virus Replication and Heterogeneous Evolutionary Rates in EVD Survivors. Cell Reports, 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. Viruses, 2018, 10, 615.	3.3	30
14	Lassa virus circulating in Liberia: a retrospective genomic characterisation. Lancet Infectious Diseases, The, 2019, 19, 1371-1378.	9.1	30
15	Chikungunya viruses containing the A226V mutation detected retrospectively in Cameroon form a new geographical subclade. International Journal of Infectious Diseases, 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 $\mid$	Overlock 10	) Tf 50 142 To
17	Novel viruses in hard ticks collected in the Republic of Korea unveiled by metagenomic high-throughput sequencing analysis. Ticks and Tick-borne Diseases, 2021, 12, 101820.	2.7	4

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