

Nicholas Di Paola

List of Publications by Year in descending order

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Version: 2024-02-01

27
papers

826
citations

623734

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552781

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docs citations

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times ranked

1529
citing authors

#	ARTICLE	IF	CITATIONS
1	Delayed viral clearance despite high number of activated T cells during the acute phase in Argentinean patients with hantavirus pulmonary syndrome. <i>EBioMedicine</i> , 2022, 75, 103765.	6.1	3
2	<i>Jingchuvirales</i> : a New Taxonomical Framework for a Rapidly Expanding Order of Unusual Monjiviricete Viruses Broadly Distributed among Arthropod Subphyla. <i>Applied and Environmental Microbiology</i> , 2022, 88, AEM0195421.	3.1	16
3	Rabies surveillance in Senegal 2001 to 2015 uncovers first infection of a honey badger. <i>Transboundary and Emerging Diseases</i> , 2022, , .	3.0	1
4	History and classification of Aigai virus (formerly Crimean-Congo haemorrhagic fever virus genotype) Tj ETQq0 0,0rgBT /Overlock 10	2.9	11
5	Integration of genomic sequencing into the response to the Ebola virus outbreak in Nord Kivu, Democratic Republic of the Congo. <i>Nature Medicine</i> , 2021, 27, 710-716.	30.7	35
6	Molecular analysis of the 2012 Bundibugyo virus disease outbreak. <i>Cell Reports Medicine</i> , 2021, 2, 100351.	6.5	4
7	2021 Taxonomic update of phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. <i>Archives of Virology</i> , 2021, 166, 3513-3566.	2.1	62
8	On-Demand Patient-Specific Phenotype-to-Genotype Ebola Virus Characterization. <i>Viruses</i> , 2021, 13, 2010.	3.3	1
9	Novel Filoviruses, Hantavirus, and Rhabdovirus in Freshwater Fish, Switzerland, 2017. <i>Emerging Infectious Diseases</i> , 2021, 27, 3082-3091.	4.3	16
10	Shedding of Marburg Virus in Naturally Infected Egyptian Rousette Bats, South Africa, 2017. <i>Emerging Infectious Diseases</i> , 2020, 26, 3051-3055.	4.3	23
11	“Super-Spreaders” and Person-to-Person Transmission of Andes Virus in Argentina. <i>New England Journal of Medicine</i> , 2020, 383, 2230-2241.	27.0	52
12	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
13	A Model for the Production of Regulatory Grade Viral Hemorrhagic Fever Exposure Stocks: From Field Surveillance to Advanced Characterization of SFTSV. <i>Viruses</i> , 2020, 12, 958.	3.3	5
14	Viral genomics in Ebola virus research. <i>Nature Reviews Microbiology</i> , 2020, 18, 365-378.	28.6	30
15	ICTV Virus Taxonomy Profile: Nairoviridae. <i>Journal of General Virology</i> , 2020, 101, 798-799.	2.9	56
16	Lassa virus circulating in Liberia: a retrospective genomic characterisation. <i>Lancet Infectious Diseases</i> , The, 2019, 19, 1371-1378.	9.1	30
17	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
18	2018 Ebola virus disease outbreak in “quateur Province, Democratic Republic of the Congo: a retrospective genomic characterisation. <i>Lancet Infectious Diseases</i> , The, 2019, 19, 641-647.	9.1	27

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19	Usutu Virus Isolated from Rodents in Senegal. <i>Viruses</i> , 2019, 11, 181.	3.3	37
20	An Outbreak of Human Parvovirus B19 Hidden by Dengue Fever. <i>Clinical Infectious Diseases</i> , 2019, 68, 810-817.	5.8	11
21	Complete Genome Sequences of Five Human Respiratory Syncytial Virus Isolates Collected in Brazil. <i>Genome Announcements</i> , 2018, 6, .	0.8	0
22	Complete Genome Sequence of a Human Metapneumovirus Isolate Collected in Brazil. <i>Genome Announcements</i> , 2018, 6, .	0.8	1
23	Molecular epidemiological study on Infectious Pancreatic Necrosis Virus isolates from aquafarms in Scotland over three decades. <i>Journal of General Virology</i> , 2018, 99, 1567-1581.	2.9	18
24	Does adaptation to vertebrate codon usage relate to flavivirus emergence potential?. <i>PLoS ONE</i> , 2018, 13, e0191652.	2.5	11
25	Complete Genome Sequences of Two Human Parainfluenza Virus Type 3 Isolates Collected in Brazil. <i>Genome Announcements</i> , 2017, 5, .	0.8	1
26	Biological and phylogenetic characteristics of West African lineages of West Nile virus. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0006078.	3.0	83
27	Where there is no overlap, there is a gap. <i>Revista Da Sociedade Brasileira De Medicina Tropical</i> , 2014, 47, 675-676.	0.9	0