Nikos Vasilakis

List of Publications by Year
in descending order

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147	11,241	36303	33894
papers	citations	51 h-index	99 g-index

153	153	153	12506
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Zika virus: History, emergence, biology, and prospects for control. Antiviral Research, 2016, 130, 69-80.	4.1	571
2	A Screen of FDA-Approved Drugs for Inhibitors of Zika Virus Infection. Cell Host and Microbe, 2016, 20, 259-270.	11.0	420
3	Characterization of a Novel Murine Model to Study Zika Virus. American Journal of Tropical Medicine and Hygiene, 2016, 94, 1362-1369.	1.4	417
4	Taxonomy of the order Mononegavirales: update 2016. Archives of Virology, 2016, 161, 2351-2360.	2.1	407
5	Molecular evolution of dengue viruses: Contributions of phylogenetics to understanding the history and epidemiology of the preeminent arboviral disease. Infection, Genetics and Evolution, 2009, 9, 523-540.	2.3	354
6	The emergence of arthropod-borne viral diseases: A global prospective on dengue, chikungunya and zika fevers. Acta Tropica, 2017, 166, 155-163.	2.0	322
7	Zika, Chikungunya, and Other Emerging Vector-Borne Viral Diseases. Annual Review of Medicine, 2018, 69, 395-408.	12.2	313
8	Zika Virus Infection and Stillbirths: A Case of Hydrops Fetalis, Hydranencephaly and Fetal Demise. PLoS Neglected Tropical Diseases, 2016, 10, e0004517.	3.0	287
9	Fever from the forest: prospects for the continued emergence of sylvatic dengue virus and its impact on public health. Nature Reviews Microbiology, 2011, 9, 532-541.	28.6	274
10	An Infectious cDNA Clone of Zika Virus to Study Viral Virulence, Mosquito Transmission, and Antiviral Inhibitors. Cell Host and Microbe, 2016, 19, 891-900.	11.0	252
11	Divergent Viruses Discovered in Arthropods and Vertebrates Revise the Evolutionary History of the Flaviviridae and Related Viruses. Journal of Virology, 2016, 90, 659-669.	3.4	242
12	Broad-spectrum agents for flaviviral infections: dengue, Zika and beyond. Nature Reviews Drug Discovery, 2017, 16, 565-586.	46.4	227
13	Taxonomy of the order Mononegavirales: update 2019. Archives of Virology, 2019, 164, 1967-1980.	2.1	224
14	Insect-Specific Virus Discovery: Significance for the Arbovirus Community. Viruses, 2015, 7, 4911-4928.	3.3	211
15	Dengue â€" Quo tu et quo vadis?. Viruses, 2011, 3, 1562-1608.	3.3	207
16	ICTV Virus Taxonomy Profile: Rhabdoviridae. Journal of General Virology, 2018, 99, 447-448.	2.9	207
17	Impact of preexisting dengue immunity on Zika virus emergence in a dengue endemic region. Science, 2019, 363, 607-610.	12.6	202
18	The family Rhabdoviridae: mono- and bipartite negative-sense RNA viruses with diverse genome organization and common evolutionary origins. Virus Research, 2017, 227, 158-170.	2.2	200

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19	Dengue viruses cluster antigenically but not as discrete serotypes. Science, 2015, 349, 1338-1343.	12.6	195
20	2020 taxonomic update for phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. Archives of Virology, 2020, 165, 3023-3072.	2.1	184
21	Arbovirus evolution in vivo is constrained by host alternation. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 6970-6975.	7.1	182
22	Outbreak of Zika Virus Infection, Chiapas State, Mexico, 2015, and First Confirmed Transmission byAedes aegyptiMosquitoes in the Americas. Journal of Infectious Diseases, 2016, 214, 1349-1356.	4.0	173
23	Taxonomy of the order Mononegavirales: update 2017. Archives of Virology, 2017, 162, 2493-2504.	2.1	173
24	Negevirus: a Proposed New Taxon of Insect-Specific Viruses with Wide Geographic Distribution. Journal of Virology, 2013, 87, 2475-2488.	3.4	166
25	Chapter 1 The History and Evolution of Human Dengue Emergence. Advances in Virus Research, 2008, 72, 1-76.	2.1	163
26	Taxonomy of the family Arenaviridae and the order Bunyavirales: update 2018. Archives of Virology, 2018, 163, 2295-2310.	2.1	157
27	Taxonomy of the order Mononegavirales: update 2018. Archives of Virology, 2018, 163, 2283-2294.	2.1	153
28	Fever versus fever: The role of host and vector susceptibility and interspecific competition in shaping the current and future distributions of the sylvatic cycles of dengue virus and yellow fever virus. Infection, Genetics and Evolution, 2013, 19, 292-311.	2.3	152
29	Evolution of Genome Size and Complexity in the Rhabdoviridae. PLoS Pathogens, 2015, 11, e1004664.	4.7	149
30	Variation inAedes aegyptiMosquito Competence for Zika Virus Transmission. Emerging Infectious Diseases, 2017, 23, 625-632.	4.3	147
31	Mosquitoes Put the Brake on Arbovirus Evolution: Experimental Evolution Reveals Slower Mutation Accumulation in Mosquito Than Vertebrate Cells. PLoS Pathogens, 2009, 5, e1000467.	4.7	146
32	Factors shaping the adaptive landscape for arboviruses: implications for the emergence of disease. Future Microbiology, 2013, 8, 155-176.	2.0	124
33	Insect-specific viruses and their potential impact on arbovirus transmission. Current Opinion in Virology, 2015, 15, 69-74.	5.4	122
34	Differential Responses of Human Fetal Brain Neural Stem Cells to Zika Virus Infection. Stem Cell Reports, 2017, 8, 715-727.	4.8	115
35	History and Emergence of Zika Virus. Journal of Infectious Diseases, 2017, 216, S860-S867.	4.0	112
36	Engineered Aedes aegypti JAK/STAT Pathway-Mediated Immunity to Dengue Virus. PLoS Neglected Tropical Diseases, 2017, 11, e0005187.	3.0	110

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37	Zika in the Americas, year 2: What have we learned? What gaps remain? A report from the Global Virus Network. Antiviral Research, 2017, 144, 223-246.	4.1	104
38	Vector-borne transmission and evolution of Zika virus. Nature Ecology and Evolution, 2019, 3, 561-569.	7.8	96
39	Guillainâ€“BarrÃ© Syndrome After Zika Virus Infection in Brazil. American Journal of Tropical Medicine and Hygiene, 2016, 95, 1157-1160.	1.4	92
40	Potential for Zika Virus to Establish a Sylvatic Transmission Cycle in the Americas. PLoS Neglected Tropical Diseases, 2016, 10, e0005055.	3.0	89
41	Flavivirus transmission focusing on Zika. Current Opinion in Virology, 2017, 22, 30-35.	5.4	87
42	Viral Load and Cytokine Response Profile Does Not Support Antibody-Dependent Enhancement in Dengue-Primed Zika Virusâ€“Infected Patients. Clinical Infectious Diseases, 2017, 65, 1260-1265.	5.8	85
43	Potential of ancestral sylvatic dengue-2 viruses to re-emerge. Virology, 2007, 358, 402-412.	2.4	78
44	Zika Virus: Diagnosis, Therapeutics, and Vaccine. ACS Infectious Diseases, 2016, 2, 170-172.	3.8	76
45	Differential Vector Competency of Aedes albopictus Populations from the Americas for Zika Virus. American Journal of Tropical Medicine and Hygiene, 2017, 97, 330-339.	1.4	72
46	Taxonomy of the order Mononegavirales: second update 2018. Archives of Virology, 2019, 164, 1233-1244.	2.1	70
47	Genetic characterization, molecular epidemiology, and phylogenetic relationships of insect-specific viruses in the taxon Negevirus. Virology, 2017, 504, 152-167.	2.4	68
48	Evidence of natural Zika virus infection in neotropical non-human primates in Brazil. Scientific Reports, 2018, 8, 16034.	3.3	68
49	Mesoniviruses are mosquito-specific viruses with extensive geographic distribution and host range. Virology Journal, 2014, 11, 97.	3.4	65
50	Evolutionary Processes among Sylvatic Dengue Type 2 Viruses. Journal of Virology, 2007, 81, 9591-9595.	3.4	64
51	2021 Taxonomic update of phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. Archives of Virology, 2021, 166, 3513-3566.	2.1	62
52	Insect-Specific Viruses Detected in Laboratory Mosquito Colonies and Their Potential Implications for Experiments Evaluating Arbovirus Vector Competence. American Journal of Tropical Medicine and Hygiene, 2015, 92, 422-428.	1.4	58
53	Insect-Specific Viruses. Advances in Virus Research, 2017, 98, 119-146.	2.1	58
54	Sylvatic Dengue Virus Type 2 Activity in Humans, Nigeria, 1966. Emerging Infectious Diseases, 2008, 14, 502-504.	4.3	54

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55	A Zika virus envelope mutation preceding the 2015 epidemic enhances virulence and fitness for transmission. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 20190-20197.	7.1	53
56	Genetic and phenotypic characterization of sylvatic dengue virus type 2 strains. Virology, 2008, 377, 296-307.	2.4	51
57	Population bottlenecks and founder effects: implications for mosquito-borne arboviral emergence. Nature Reviews Microbiology, 2021, 19, 184-195.	28.6	51
58	Lineage II of Southeast Asian/American DENV-2 is Associated with a Severe Dengue Outbreak in the Peruvian Amazon. American Journal of Tropical Medicine and Hygiene, 2014, 91, 611-620.	1.4	50
59	A Newly Isolated Reovirus Has the Simplest Genomic and Structural Organization of Any Reovirus. Journal of Virology, 2015, 89, 676-687.	3.4	50
60	ICTV Virus Taxonomy Profile: Rhabdoviridae 2022. Journal of General Virology, 2022, 103, .	2.9	46
61	Characterization of Three New Insect-Specific Flaviviruses: Their Relationship to the Mosquito-Borne Flavivirus Pathogens. American Journal of Tropical Medicine and Hygiene, 2018, 98, 410-419.	1.4	45
62	Did Zika Virus Mutate to Cause Severe Outbreaks?. Trends in Microbiology, 2018, 26, 877-885.	7.7	43
63	Abundance and distribution of sylvatic dengue virus vectors in three different land cover types in Sarawak, Malaysian Borneo. Parasites and Vectors, 2017, 10, 406.	2.5	42
64	Genomes of viral isolates derived from different mosquitos species. Virus Research, 2017, 242, 49-57.	2.2	40
65	Zika, dengue and yellow fever viruses induce differential anti-viral immune responses in human monocytic and first trimester trophoblast cells. Antiviral Research, 2018, 151, 55-62.	4.1	40
66	Arboretum and Puerto Almendras viruses: two novel rhabdoviruses isolated from mosquitoes in Peru. Journal of General Virology, 2014, 95, 787-792.	2.9	39
67	Into the woods: Changes in mosquito community composition and presence of key vectors at increasing distances from the urban edge in urban forest parks in Manaus, Brazil. Acta Tropica, 2020, 206, 105441.	2.0	39
68	Seroprevalence of Neutralizing Antibodies Against Dengue Virus in Two Localities in the State of Morelos, Mexico. American Journal of Tropical Medicine and Hygiene, 2014, 91, 1057-1065.	1.4	38
69	Experimental Zika Virus Infection of Neotropical Primates. American Journal of Tropical Medicine and Hygiene, 2018, 98, 173-177.	1.4	38
70	Genetic and phenotypic characterization of sylvatic dengue virus type 4 strains. Virology, 2012, 423, 58-67.	2.4	37
71	Genomic Characterization of Yogue, Kasokero, Issyk-Kul, Keterah, Gossas, and Thiafora Viruses: Nairoviruses Naturally Infecting Bats, Shrews, and Ticks. American Journal of Tropical Medicine and Hygiene, 2015, 93, 1041-1051.	1.4	36
72	Impact of SARS-CoV-2 Gamma lineage introduction and COVID-19 vaccination on the epidemiological landscape of a Brazilian city. Communications Medicine, 2022, 2, .	4.2	32

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73	Modeling Zika Virus Infection in Mice. Cell Stem Cell, 2016, 19, 4-6.	11.1	30
74	Role of mutational reversions and fitness restoration in Zika virus spread to the Americas. Nature Communications, 2021, 12, 595.	12.8	29
75	Antigenic Relationships between Sylvatic and Endemic Dengue Viruses. American Journal of Tropical Medicine and Hygiene, 2008, 79, 128-132.	1.4	29
76	Kolente virus, a rhabdovirus species isolated from ticks and bats in the Republic of Guinea. Journal of General Virology, 2013, 94, 2609-2615.	2.9	28
77	Ledantevirus: A Proposed New Genus in the Rhabdoviridae has a Strong Ecological Association with Bats. American Journal of Tropical Medicine and Hygiene, 2015, 92, 405-410.	1.4	27
78	Almendravirus : A Proposed New Genus of Rhabdoviruses Isolated from Mosquitoes in Tropical Regions of the Americas. American Journal of Tropical Medicine and Hygiene, 2017, 96, 100-109.	1.4	27
79	A Tale of Two Viruses: Does Heterologous Flavivirus Immunity Enhance Zika Disease?. Trends in Microbiology, 2018, 26, 186-190.	7.7	27
80	The vertical stratification of potential bridge vectors of mosquito-borne viruses in a central Amazonian forest bordering Manaus, Brazil. Scientific Reports, 2020, 10, 18254.	3.3	27
81	Niakha virus: A novel member of the family Rhabdoviridae isolated from phlebotomine sandflies in Senegal. Virology, 2013, 444, 80-89.	2.4	26
82	Potential for sylvatic and urban Aedes mosquitoes from Senegal to transmit the new emerging dengue serotypes 1, 3 and 4 in West Africa. PLoS Neglected Tropical Diseases, 2019, 13, e0007043.	3.0	26
83	Antigenic relationships between sylvatic and endemic dengue viruses. American Journal of Tropical Medicine and Hygiene, 2008, 79, 128-32.	1.4	26
84	Itaya virus, a NovelOrthobunyavirusAssociated with Human Febrile Illness, Peru. Emerging Infectious Diseases, 2015, 21, 781-8.	4.3	25
85	Sylvatic Dengue Viruses Share the Pathogenic Potential of Urban/Endemic Dengue Viruses. Journal of Virology, 2010, 84, 3726-3728.	3.4	24
86	Emergence potential of sylvatic dengue virus type 4 in the urban transmission cycle is restrained by vaccination and homotypic immunity. Virology, 2013, 439, 34-41.	2.4	24
87	Lack of evidence for Zika virus transmission by Culex mosquitoes. Emerging Microbes and Infections, 2017, 6, 1-2.	6.5	24
88	Unusual clinical manifestations of dengue disease â€“ Real or imagined?. Acta Tropica, 2019, 199, 105134.	2.0	24
89	Experimental Infection with and Maintenance of Cell Fusing Agent Virus (Flavivirus) in Aedes aegypti. American Journal of Tropical Medicine and Hygiene, 2017, 97, 299-304.	1.4	24
90	Transient Hearing Loss in Adults Associated with Zika Virus Infection. Clinical Infectious Diseases, 2016, 64, ciw770.	5.8	23

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91	Colonized Sabethes cyaneus, a Sylvatic New World Mosquito Species, Shows a Low Vector Competence for Zika Virus Relative to Aedes aegypti. Viruses, 2018, 10, 434.	3.3	23
92	Case Study of Two Post Vaccination SARS-CoV-2 Infections with P1 Variants in CoronaVac Vaccinees in Brazil. Viruses, 2021, 13, 1237.	3.3	23
93	Sinu virus, a novel and divergent orthomyxovirus related to members of the genus Thogotovirus isolated from mosquitoes in Colombia. Virology, 2017, 501, 166-175.	2.4	22
94	Re-emergence of yellow fever in the neotropics â€" quo vadis?. Emerging Topics in Life Sciences, 2020, 4, 411-422.	2.6	22
95	Mercadeo Virus: A Novel Mosquito-Specific Flavivirus from Panama. American Journal of Tropical Medicine and Hygiene, 2015, 93, 1014-1019.	1.4	21
96	Bunyavirus Taxonomy: Limitations and Misconceptions Associated with the Current ICTV Criteria Used for Species Demarcation. American Journal of Tropical Medicine and Hygiene, 2018, 99, 11-16.	1.4	21
97	Infection Dynamics of Sylvatic Dengue Virus in a Natural Primate Host, the African Green Monkey. American Journal of Tropical Medicine and Hygiene, 2014, 91, 672-676.	1.4	20
98	Evaluation of Aptima Zika Virus Assay. Journal of Clinical Microbiology, 2017, 55, 2198-2203.	3.9	19
99	Molecular Epidemiology of Dengue in Panama: 25 Years of Circulation. Viruses, 2019, 11, 764.	3.3	18
100	Possibility and Challenges of Conversion of Current Virus Species Names to Linnaean Binomials. Systematic Biology, 2016, 66, syw096.	5.6	17
101	Viral immunogenicity determines epidemiological fitness in a cohort of DENV-1 infection in Brazil. PLoS Neglected Tropical Diseases, 2018, 12, e0006525.	3.0	17
102	Exploiting the Legacy of the Arbovirus Hunters. Viruses, 2019, 11, 471.	3.3	17
103	Fatal Outcome of Ilheus Virus in the Cerebrospinal Fluid of a Patient Diagnosed with Encephalitis. Viruses, 2020, 12, 957.	3.3	17
104	Presentation of fatal stroke due to SARSâ€CoVâ€2 and dengue virus coinfection. Journal of Medical Virology, 2021, 93, 1770-1775.	5.0	16
105	Characterization of Farmington virus, a novel virus from birds that is distantly related to members of the family Rhabdoviridae. Virology Journal, 2013, 10, 219.	3.4	14
106	Whole Genome Analysis of Sierra Nevada Virus, a Novel Mononegavirus in the Family Nyamiviridae. American Journal of Tropical Medicine and Hygiene, 2014, 91, 159-164.	1.4	14
107	Molecular classification of outcomes from dengue virus -3 infections. Journal of Clinical Virology, 2015, 64, 97-106.	3.1	14
108	Dengue virus surveillance: Detection of DENV-4 in the city of SÃ£o JosÃ© do Rio Preto, SP, Brazil. Acta Tropica, 2016, 164, 84-89.	2.0	14

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109	Characterization of five unclassified orthobunyaviruses (Bunyaviridae) from Africa and the Americas. Journal of General Virology, 2017, 98, 2258-2266.	2.9	13
110	Rocio Virus: An Updated View on an Elusive Flavivirus. Viruses, 2021, 13, 2293.	3.3	13
111	Malpais spring virus is a new species in the genus vesiculovirus. Virology Journal, 2013, 10, 69.	3.4	11
112	Support for the Transmission-Clearance Trade-Off Hypothesis from a Study of Zika Virus Delivered by Mosquito Bite to Mice. Viruses, 2019, 11, 1072.	3.3	11
113	Strengthening the Interaction of the Virology Community with the International Committee on Taxonomy of Viruses (ICTV) by Linking Virus Names and Their Abbreviations to Virus Species. Systematic Biology, 2019, 68, 828-839.	5.6	11
114	Shifts in mosquito diversity and abundance along a gradient from oil palm plantations to conterminous forests in Borneo. Ecosphere, 2021, 12, e03463.	2.2	11
115	Identification of a new newcastle disease virus isolate from Indonesia represents an ancestral lineage of class II genotype XIII. Virus Genes, 2013, 47, 168-172.	1.6	10
116	Koolpinyah and Yata viruses: Two newly recognised ephemeroviruses from tropical regions of Australia and Africa. Veterinary Microbiology, 2014, 174, 547-553.	1.9	10
117	Electron Microscopy in Discovery of Novel and Emerging Viruses from the Collection of the World Reference Center for Emerging Viruses and Arboviruses (WRCEVA). Viruses, 2019, 11, 477.	3.3	10
118	Genomic characterisation of Cuiaba and Charleville viruses: arboviruses (family Rhabdoviridae, genus) Tj ETQq0 0 0 rgBT /Overlock 10 Tf	1.6	10
119	ZIKV Demonstrates Minimal Pathologic Effects and Mosquito Infectivity in Viremic Cynomolgus Macaques. Viruses, 2018, 10, 661.	3.3	9
120	Evolution of resistance to fluoroquinolones by dengue virus serotype 4 provides insight into mechanism of action and consequences for viral fitness. Virology, 2021, 552, 94-106.	2.4	9
121	Characterization of the Gamboa Virus Serogroup (Orthobunyavirus Genus, Peribunyaviridae Family). American Journal of Tropical Medicine and Hygiene, 2018, 98, 1502-1511.	1.4	9
122	Emergence potential of mosquito-borne arboviruses from the Florida Everglades. PLoS ONE, 2021, 16, e0259419.	2.5	9
123	Dianke virus: A new mesonivirus species isolated from mosquitoes in Eastern Senegal. Virus Research, 2020, 275, 197802.	2.2	8
124	Identification of Mosquito Bloodmeals Collected in Diverse Habitats in Malaysian Borneo Using COI Barcoding. Tropical Medicine and Infectious Disease, 2020, 5, 51.	2.3	7
125	Characterization of Port Bolivar Virus, a Novel Entomobirnavirus (Birnaviridae) Isolated from Mosquitoes Collected in East Texas, USA. Viruses, 2020, 12, 390.	3.3	7
126	Lack of Evidence of Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) Spillover in Free-Living Neotropical Non-Human Primates, Brazil. Viruses, 2021, 13, 1933.	3.3	7

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127	Inhibition of innate immune response ameliorates Zika virus-induced neurogenesis deficit in human neural stem cells. PLoS Neglected Tropical Diseases, 2021, 15, e0009183.	3.0	6
128	Characterization of Triniti virus supports its reclassification in the family Peribunyaviridae. Journal of General Virology, 2019, 100, 137-144.	2.9	6
129	Microclimate and the vertical stratification of potential bridge vectors of mosquitoâ€'borne viruses captured by nets and ovitraps in a central Amazonian forest bordering Manaus, Brazil. Scientific Reports, 2021, 11, 21129.	3.3	6
130	GeneSV â€“ an Approach to Help Characterize Possible Variations in Genomic and Protein Sequences. Bioinformatics and Biology Insights, 2014, 8, BBI.S13076.	2.0	5
131	Characterization of Three Novel Viruses from the Families Nyamiviridae, Orthomyxoviridae, and Peribunyaviridae, Isolated from Dead Birds Collected during West Nile Virus Surveillance in Harris County, Texas. Viruses, 2019, 11, 927.	3.3	5
132	Flavivirus Infection Associated with Cerebrovascular Events. Viruses, 2020, 12, 671.	3.3	5
133	Why Did ZIKV Perinatal Outcomes Differ in Distinct Regions of Brazil? An Exploratory Study of Two Cohorts. Viruses, 2021, 13, 736.	3.3	5
134	ICTV Virus Taxonomy Profile: Nyamiviridae. Journal of General Virology, 2017, 98, 2914-2915.	2.9	5
135	Implications of a highly divergent dengue virus strain for cross-neutralization, protection, and vaccine immunity. Cell Host and Microbe, 2021, 29, 1634-1648.e5.	11.0	5
136	The Arboviruses: Quo Vadis?. , 2016, , 1-6.		4
137	The reintroduction of DENV-2 in 2011 in Panama and subsequent outbreak characteristic. Acta Tropica, 2018, 177, 58-65.	2.0	3
138	Aedes aegypti Shows Increased Susceptibility to Zika Virus via Both In Vitro and In Vivo Models of Type II Diabetes. Viruses, 2022, 14, 665.	3.3	3
139	Arboviral Infections in Neurological Disorders in Hospitalized Patients in SÃ£o JosÃ© do Rio Preto, SÃ£o Paulo, Brazil. Viruses, 2022, 14, 1488.	3.3	3
140	Experimental Zika Virus Infection in a Neotropical Primate Model. Open Forum Infectious Diseases, 2016, 3, .	0.9	2
141	Age and Sex in the Zika Pandemic Era. Journal of Infectious Diseases, 2018, 217, 1675-1677.	4.0	2
142	Genome Sequence of Chiqui Virus, a Novel Reovirus Isolated from Mosquitoes Collected in Colombia. Microbiology Resource Announcements, 2018, 7, .	0.6	2
143	Reversible sensory polyneuropathy during an arboviral outbreak in Salvador, Bahia, Brazil. Journal of the Neurological Sciences, 2018, 391, 3-4.	0.6	1
144	Seek and You Shall Find â€" Unknown Pathogens?. New England Journal of Medicine, 2019, 380, 2174-2175.	27.0	1

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145	ICTV Virus Taxonomy Profile: Artoviridae. Journal of General Virology, 2019, 100, 1202-1203.	2.9	1
146	ICTV Virus Taxonomy Profile: Nyamiviridae 2021. Journal of General Virology, 2021, 102, .	2.9	1
147	Zika Virus (Flaviviridae). , 2021, , 899-909.		0