

Nicolás E Bejerman

List of Publications by Year in descending order

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Version: 2024-02-01

49
papers

1,437
citations

471509

17
h-index

361022

35
g-index

62
all docs

62
docs citations

62
times ranked

1905
citing authors

#	ARTICLE	IF	CITATIONS
1	Taxonomy of the order Mononegavirales: update 2016. <i>Archives of Virology</i> , 2016, 161, 2351-2360.	2.1	407
2	Taxonomy of the order Mononegavirales: update 2017. <i>Archives of Virology</i> , 2017, 162, 2493-2504.	2.1	173
3	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
4	Species Within the Bemisia tabaci (Hemiptera: Aleyrodidae) Complex in Soybean and Bean Crops in Argentina. <i>Journal of Economic Entomology</i> , 2012, 105, 48-53.	1.8	60
5	Diversity and epidemiology of plant rhabdoviruses. <i>Virus Research</i> , 2020, 281, 197942.	2.2	56
6	Complete genome sequence and integrated protein localization and interaction map for alfalfa dwarf virus, which combines properties of both cytoplasmic and nuclear plant rhabdoviruses. <i>Virology</i> , 2015, 483, 275-283.	2.4	54
7	Genome-enabled insights into the biology of thrips as crop pests. <i>BMC Biology</i> , 2020, 18, 142.	3.8	54
8	Illuminating the Plant Rhabdovirus Landscape through Metatranscriptomics Data. <i>Viruses</i> , 2021, 13, 1304.	3.3	45
9	Cytorhabdovirus P3 genes encode 30K-like cell-to-cell movement proteins. <i>Virology</i> , 2016, 489, 20-33.	2.4	32
10	Complete genome sequence of a new enamovirus from Argentina infecting alfalfa plants showing dwarfism symptoms. <i>Archives of Virology</i> , 2016, 161, 2029-2032.	2.1	30
11	Novel birdâ€™s-foot trefoil RNA viruses provide insights into a clade of legume-associated enamoviruses and rhabdoviruses. <i>Archives of Virology</i> , 2019, 164, 1419-1426.	2.1	29
12	The Plant Negative-Sense RNA Virosphere: Virus Discovery Through New Eyes. <i>Frontiers in Microbiology</i> , 2020, 11, 588427.	3.5	29
13	Complete genome sequence and intracellular protein localization of Datura yellow vein nucleorhabdovirus. <i>Virus Research</i> , 2015, 205, 7-11.	2.2	24
14	High-Throughput Sequencing for Deciphering the Virome of Alfalfa (Medicago sativa L.). <i>Frontiers in Microbiology</i> , 2020, 11, 553109.	3.5	24
15	Exploring the tymovirales landscape through metatranscriptomics data. <i>Archives of Virology</i> , 2022, 167, 1785-1803.	2.1	24
16	Complete nucleotide sequence of Alfalfa mosaic virus isolated from alfalfa (Medicago sativa L.) in Argentina. <i>Virus Genes</i> , 2014, 48, 562-565.	1.6	20
17	Viromes of Ten Alfalfa Plants in Australia Reveal Diverse Known Viruses and a Novel RNA Virus. <i>Pathogens</i> , 2020, 9, 214.	2.8	20
18	Molecular characterization of Sunflower chlorotic mottle virus: a member of a distinct species in the genus Potyvirus. <i>Archives of Virology</i> , 2010, 155, 1331-1335.	2.1	19

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19	Bean leafroll virus (BLRV) in Argentina: molecular characterization and detection in alfalfa fields. <i>European Journal of Plant Pathology</i> , 2016, 146, 207-212.	1.7	17
20	Genome characterization of an Argentinean isolate of alfalfa leaf curl virus. <i>Archives of Virology</i> , 2018, 163, 799-803.	2.1	16
21	Distribution and genetic variability of alfalfa dwarf virus, a cytorhabdovirus associated with alfalfa dwarf disease in Argentina. <i>Virus Genes</i> , 2018, 54, 612-615.	1.6	16
22	Alfalfa dwarf cytorhabdovirus P protein is a local and systemic RNA silencing suppressor which inhibits programmed RISC activity and prevents transitive amplification of RNA silencing. <i>Virus Research</i> , 2016, 224, 19-28.	2.2	15
23	Complete nucleotide sequence of an Argentinean isolate of sweet potato virus G. <i>Virus Genes</i> , 2012, 45, 593-595.	1.6	14
24	Molecular characterization of a novel cytorhabdovirus with a unique genomic organization infecting yerba mate (<i>Ilex paraguariensis</i>) in Argentina. <i>Archives of Virology</i> , 2020, 165, 1475-1479.	2.1	14
25	Complete genome sequence of Colocasia bobone disease-associated virus, a putative cytorhabdovirus infecting taro. <i>Archives of Virology</i> , 2016, 161, 745-748.	2.1	13
26	Molecular characterization of yerba mate chlorosis-associated virus, a putative cytorhabdovirus infecting yerba mate (<i>Ilex paraguariensis</i>). <i>Archives of Virology</i> , 2017, 162, 2481-2484.	2.1	13
27	Bean Yellow Mosaic Virus in Soybean from Argentina. <i>Journal of Phytopathology</i> , 2014, 162, 322-325.	1.0	12
28	The Westward Journey of Alfalfa Leaf Curl Virus. <i>Viruses</i> , 2018, 10, 542.	3.3	12
29	Redefining the medicago sativa alphapartitiviruses genome sequences. <i>Virus Research</i> , 2019, 265, 156-161.	2.2	12
30	Identification and Characterization of a New Strain of Sunflower chlorotic mottle virus, a Potyvirus Infecting Asteraceae in Argentina. <i>Journal of Phytopathology</i> , 2010, 158, 536-541.	1.0	9
31	Letter to the Editor: Bean-Associated Cytorhabdovirus and Papaya Cytorhabdovirus are Strains of the Same Virus. <i>Viruses</i> , 2019, 11, 230.	3.3	9
32	First Report of Orchid fleck virus in Lilyturf (<i>Liriope spicata</i>) in Australia. <i>Plant Disease</i> , 2016, 100, 1028-1028.	1.4	8
33	<i>Dipsacus fullonum</i> : an Alternative Host of Sunflower chlorotic mottle virus in Argentina. <i>Journal of Phytopathology</i> , 2009, 157, 325-328.	1.0	7
34	Complete genome sequence of sunflower ring blotch virus, a new potyvirus infecting sunflower in Argentina. <i>Archives of Virology</i> , 2017, 162, 1787-1790.	2.1	7
35	Sequencing of two Sunflower chlorotic mottle virus isolates obtained from different natural hosts shed light on its evolutionary history. <i>Virus Genes</i> , 2013, 46, 105-110.	1.6	5
36	Development and validation of PCR assays for detection of alfalfa dwarf disease-associated viruses in Australian lucerne pastures. <i>Australasian Plant Pathology</i> , 2018, 47, 215-225.	1.0	5

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37	First Report of Tobacco Streak Virus Infecting Sunflower in Argentina. <i>Plant Disease</i> , 2019, 103, 3290.	1.4	5
38	Geminivirus-Vector Relationship. , 2019, , 137-145.		4
39	Identification and molecular characterization of a novel circular single-stranded DNA virus associated with yerba mate in Argentina. <i>Archives of Virology</i> , 2018, 163, 2811-2815.	2.1	3
40	Biological and molecular characterization of bean bushy stunt virus, a novel bipartite begomovirus infecting common bean in northwestern Argentina. <i>Archives of Virology</i> , 2021, 166, 1409-1414.	2.1	3
41	Joã yellow blotch-associated virus, a new alphanucleorhabdovirus from a wild solanaceous plant in Brazil. <i>Archives of Virology</i> , 2021, 166, 1615-1622.	2.1	3
42	First Report of <i>Bean common mosaic virus</i>, Peanut Strain, Infecting Peanut in Argentina. <i>Plant Disease</i> , 2015, 99, 735-735.	1.4	3
43	Development of a full-length infectious clone of sunflower chlorotic mottle virus (SuCMoV). <i>Archives of Virology</i> , 2013, 158, 485-490.	2.1	2
44	Use of 454-sequencing for the Characterization of <i>Sweet Potato Virus C</i> and <i>Sweet Potato Feathery Mottle Virus</i> Isolates from Argentina and Development of a Multiplex One-Step RT-PCR for Their Simultaneous Detection. <i>Journal of Phytopathology</i> , 2016, 164, 386-394.	1.0	2
45	Analysis of the coding-complete genomic sequence of groundnut ringspot virus suggests a common ancestor with tomato chlorotic spot virus. <i>Archives of Virology</i> , 2016, 161, 2311-2316.	2.1	2
46	Phylogenetics of sunflower chlorotic mottle virus, an emerging pathosystem. <i>Virology</i> , 2020, 545, 33-39.	2.4	2
47	Exploring species composition and population dynamics of thrips (Thysanoptera: Thripidae) in peanut crops in Argentina. <i>Phytoparasitica</i> , 2021, 49, 785-792.	1.2	2
48	Orthotospovirus disease epidemic: molecular characterization and incidence in peanut crops. <i>Journal of Plant Pathology</i> , 2021, 103, 305-309.	1.2	1
49	Papaya ringspot virus W infecting <i>Luffa aegyptiaca</i> in Cuba. <i>Australasian Plant Disease Notes</i> , 2017, 12, 1.	0.7	0