

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
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361022

35
g-index

62
all docs

62
docs citations

62
times ranked

1905
citing authors

#	ARTICLE	IF	CITATIONS
1	Exploring the tymovirales landscape through metatranscriptomics data. Archives of Virology, 2022, 167, 1785-1803.	2.1	24
2	Orthotospovirus disease epidemic: molecular characterization and incidence in peanut crops. Journal of Plant Pathology, 2021, 103, 305-309.	1.2	1
3	Biological and molecular characterization of bean bushy stunt virus, a novel bipartite begomovirus infecting common bean in northwestern Argentina. Archives of Virology, 2021, 166, 1409-1414.	2.1	3
4	Joã yellow blotch-associated virus, a new alphanucleorhabdovirus from a wild solanaceous plant in Brazil. Archives of Virology, 2021, 166, 1615-1622.	2.1	3
5	Exploring species composition and population dynamics of thrips (Thysanoptera: Thripidae) in peanut crops in Argentina. Phytoparasitica, 2021, 49, 785-792.	1.2	2
6	Illuminating the Plant Rhabdovirus Landscape through Metatranscriptomics Data. Viruses, 2021, 13, 1304.	3.3	45
7	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
8	High-Throughput Sequencing for Deciphering the Virome of Alfalfa (Medicago sativa L.). Frontiers in Microbiology, 2020, 11, 553109.	3.5	24
9	Genome-enabled insights into the biology of thrips as crop pests. BMC Biology, 2020, 18, 142.	3.8	54
10	The Plant Negative-Sense RNA Viroisphere: Virus Discovery Through New Eyes. Frontiers in Microbiology, 2020, 11, 588427.	3.5	29
11	Viromes of Ten Alfalfa Plants in Australia Reveal Diverse Known Viruses and a Novel RNA Virus. Pathogens, 2020, 9, 214.	2.8	20
12	Molecular characterization of a novel cytorhabdovirus with a unique genomic organization infecting yerba mate (Ilex paraguariensis) in Argentina. Archives of Virology, 2020, 165, 1475-1479.	2.1	14
13	Phylogenetics of sunflower chlorotic mottle virus, an emerging pathosystem. Virology, 2020, 545, 33-39.	2.4	2
14	Diversity and epidemiology of plant rhabdoviruses. Virus Research, 2020, 281, 197942.	2.2	56
15	Geminivirus-Vector Relationship. , 2019, , 137-145.		4
16	Novel bird-foam trefoil RNA viruses provide insights into a clade of legume-associated enamoviruses and rhabdoviruses. Archives of Virology, 2019, 164, 1419-1426.	2.1	29
17	Letter to the Editor: Bean-Associated Cytorhabdovirus and Papaya Cytorhabdovirus are Strains of the Same Virus. Viruses, 2019, 11, 230.	3.3	9
18	Redefining the medicago sativa alphapartitiviruses genome sequences. Virus Research, 2019, 265, 156-161.	2.2	12

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19	First Report of Tobacco Streak Virus Infecting Sunflower in Argentina. <i>Plant Disease</i> , 2019, 103, 3290.	1.4	5
20	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
21	Genome characterization of an Argentinean isolate of alfalfa leaf curl virus. <i>Archives of Virology</i> , 2018, 163, 799-803.	2.1	16
22	The Westward Journey of Alfalfa Leaf Curl Virus. <i>Viruses</i> , 2018, 10, 542.	3.3	12
23	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
24	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
25	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
26	Taxonomy of the order Mononegavirales: update 2017. <i>Archives of Virology</i> , 2017, 162, 2493-2504.	2.1	173
27	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
28	Papaya ringspot virus W infecting <i>Luffa aegyptiaca</i> in Cuba. <i>Australasian Plant Disease Notes</i> , 2017, 12, 1.	0.7	0
29	Taxonomy of the order Mononegavirales: update 2016. <i>Archives of Virology</i> , 2016, 161, 2351-2360.	2.1	407
30	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
31	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
32	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
33	Use of 454 Pyrosequencing 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
34	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
35	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
36	Cytorhabdovirus P3 genes encode 30K-like cell-to-cell movement proteins. <i>Virology</i> , 2016, 489, 20-33.	2.4	32

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37	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
38	Complete genome sequence and intracellular protein localization of <i>Datura</i> yellow vein nucleorhabdovirus. <i>Virus Research</i> , 2015, 205, 7-11.	2.2	24
39	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
40	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
41	<i>Bean Yellow Mosaic Virus</i> in Soybean from Argentina. <i>Journal of Phytopathology</i> , 2014, 162, 322-325.	1.0	12
42	Complete nucleotide sequence of Alfalfa mosaic virus isolated from alfalfa (<i>Medicago sativa</i> L.) in Argentina. <i>Virus Genes</i> , 2014, 48, 562-565.	1.6	20
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	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
45	Complete nucleotide sequence of an Argentinean isolate of sweet potato virus G. <i>Virus Genes</i> , 2012, 45, 593-595.	1.6	14
46	Species Within the <i>Bemisia tabaci</i> (Hemiptera: Aleyrodidae) Complex in Soybean and Bean Crops in Argentina. <i>Journal of Economic Entomology</i> , 2012, 105, 48-53.	1.8	60
47	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
48	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
49	<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