

Michela Chiumenti

List of Publications by Year in descending order

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46
papers

895
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623734

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#	ARTICLE	IF	CITATIONS
1	A New Jasmine Virus C Isolate Identified by Nanopore Sequencing Is Associated to Yellow Mosaic Symptoms of <i>Jasminum officinale</i> in Italy. <i>Plants</i> , 2022, 11, 309.	3.5	5
2	Pest categorisation of <i>Apium virus Y</i> . <i>EFSA Journal</i> , 2022, 20, e06930.	1.8	1
3	Commodity risk assessment of grafted plants of <i>Malus domestica</i> from Moldova. <i>EFSA Journal</i> , 2022, 20, e07201.	1.8	1
4	Commodity risk assessment of <i>Malus domestica</i> plants from Turkey. <i>EFSA Journal</i> , 2022, 20, e07301.	1.8	3
5	Pest categorisation of High Plains wheat mosaic virus. <i>EFSA Journal</i> , 2022, 20, e07302.	1.8	2
6	Sixty Years from the First Disease Description, a Novel Badnavirus Associated with Chestnut Mosaic Disease. <i>Phytopathology</i> , 2021, 111, 1051-1058.	2.2	6
7	Reassessing species demarcation criteria in viroid taxonomy by pairwise identity matrices. <i>Virus Evolution</i> , 2021, 7, veab001.	4.9	13
8	<i>Olea Europaea Geminivirus</i> : A Novel Bipartite Geminivirid Infecting Olive Trees. <i>Viruses</i> , 2021, 13, 481.	3.3	16
9	High throughput sequencing from Angolan citrus accessions discloses the presence of emerging CTV strains. <i>Virology Journal</i> , 2021, 18, 62.	3.4	1
10	Degradome Analysis of Tomato and <i>Nicotiana benthamiana</i> Plants Infected with Potato Spindle Tuber Viroid. <i>International Journal of Molecular Sciences</i> , 2021, 22, 3725.	4.1	13
11	A Primer on the Analysis of High-Throughput Sequencing Data for Detection of Plant Viruses. <i>Microorganisms</i> , 2021, 9, 841.	3.6	36
12	Identification and Characterization of Citrus Concave Gum-Associated Virus Infecting Citrus and Apple Trees by Serological, Molecular and High-Throughput Sequencing Approaches. <i>Plants</i> , 2021, 10, 2390.	3.5	10
13	Commodity risk assessment of <i>Malus domestica</i> plants from Ukraine. <i>EFSA Journal</i> , 2021, 19, e06909.	1.8	0
14	Pest categorisation of carrot thin leaf virus. <i>EFSA Journal</i> , 2021, 19, e06931.	1.8	0
15	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
16	Pest categorisation of non-EU viruses of <i>Rubus L.</i> <i>EFSA Journal</i> , 2020, 18, e05928.	1.8	6
17	List of non-EU phytoplasmas of <i>Cydonia Mill.</i> , <i>Fragaria L.</i> , <i>Malus Mill.</i> , <i>Prunus L.</i> , <i>Pyrus L.</i> , <i>Ribes L.</i> , <i>Rubus L.</i> and <i>Vitis L.</i> <i>EFSA Journal</i> , 2020, 18, e05930.	1.8	1
18	Pest categorisation of the non-EU phytoplasmas of <i>Cydonia Mill.</i> , <i>Fragaria L.</i> , <i>Malus Mill.</i> , <i>Prunus L.</i> , <i>Pyrus L.</i> , <i>Ribes L.</i> , <i>Rubus L.</i> and <i>Vitis L.</i> <i>EFSA Journal</i> , 2020, 18, e05929.	1.8	7

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19	List of non-EU phytoplasmas of tuber-forming <i>Solanum</i> spp.. EFSA Journal, 2020, 18, e06355.	1.8	1
20	Pest categorisation of the non-EU phytoplasmas of tuber-forming <i>Solanum</i> spp.. EFSA Journal, 2020, 18, e06356.	1.8	1
21	First Report of Citrus Viroids Infecting Persian (Tahiti) Lime in Greece. Plant Disease, 2020, 104, 998-998.	1.4	6
22	Virus Detection by High-Throughput Sequencing of Small RNAs: Large-Scale Performance Testing of Sequence Analysis Strategies. Phytopathology, 2019, 109, 488-497.	2.2	106
23	Molecular variability of apple hammerhead viroid from Italian apple varieties supports the relevance in vivo of its branched conformation stabilized by a kissing loop interaction. Virus Research, 2019, 270, 197644.	2.2	8
24	Pest categorisation of non-EU viruses of <i>Fragaria</i> L.. EFSA Journal, 2019, 17, e05766.	1.8	3
25	Pest categorisation of non-EU viruses and viroids of <i>Cydonia</i> Mill., <i>Malus</i> Mill. and <i>Pyrus</i> L.. EFSA Journal, 2019, 17, e05590.	1.8	7
26	Pest categorisation of non-EU viruses and viroids of <i>Vitis</i> L.. EFSA Journal, 2019, 17, e05669.	1.8	6
27	List of non-EU viruses and viroids of <i>Cydonia</i> Mill., <i>Fragaria</i> L., <i>Malus</i> Mill., <i>Prunus</i> L., <i>Pyrus</i> L., <i>Ribes</i> L., <i>Rubus</i> L. and <i>Vitis</i> L.. EFSA Journal, 2019, 17, e05501.	1.8	15
28	Pest categorisation of non-EU viruses and viroids of <i>Prunus</i> L.. EFSA Journal, 2019, 17, e05735.	1.8	5
29	Molecular characterisation of a novel gemycircularvirus associated with olive trees in Italy. Virus Research, 2019, 263, 169-172.	2.2	10
30	A new emaravirus discovered in <i>Pistacia</i> from Turkey. Virus Research, 2019, 263, 159-163.	2.2	28
31	How sequence variants of a plastid-replicating viroid with one single nucleotide change initiate disease in its natural host. RNA Biology, 2019, 16, 906-917.	3.1	19
32	Pest categorisation of non-EU viruses of <i>Ribes</i> L.. EFSA Journal, 2019, 17, e05859.	1.8	4
33	Small RNA Isolation from Tissues of Grapevine and Woody Plants. Methods in Molecular Biology, 2018, 1746, 27-36.	0.9	2
34	Molecular and biological characterization of a novel mild strain of citrus tristeza virus in California. Archives of Virology, 2018, 163, 1795-1804.	2.1	31
35	A Short Indel-Lacking-Resistance Gene Triggers Silencing of the Photosynthetic Machinery Components Through TYLCSV-Associated Endogenous siRNAs in Tomato. Frontiers in Plant Science, 2018, 9, 1470.	3.6	15
36	Identification and Characterization of <i>Citrus tristeza virus</i> Isolates Breaking Resistance in Trifoliate Orange in California. Phytopathology, 2017, 107, 901-908.	2.2	33

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37	First Report of <i>Cherry virus A</i> and <i>Plum bark necrosis stem pitting-associated virus</i> in Cherry in Chile. <i>Plant Disease</i> , 2017, 101, 1685-1685.	1.4	5
38	Detection and molecular characterization of a Grapevine Roditis leaf discoloration-associated virus (GRLDaV) variant in an autochthonous grape from Apulia (Italy). <i>Virus Genes</i> , 2016, 52, 428-431.	1.6	9
39	High-throughput-sequencing-based identification of a grapevine fanleaf virus satellite RNA in <i>Vitis vinifera</i> . <i>Archives of Virology</i> , 2016, 161, 1401-1403.	2.1	9
40	Transcriptome profiling of two olive cultivars in response to infection by the CoDiRO strain of <i>Xylella fastidiosa</i> subsp. <i>pauca</i> . <i>BMC Genomics</i> , 2016, 17, 475.	2.8	118
41	Unusual genomic features of a badnavirus infecting mulberry. <i>Journal of General Virology</i> , 2016, 97, 3073-3087.	2.9	19
42	Draft Genome Sequence of CO33, a Coffee-Infecting Isolate of <i>Xylella fastidiosa</i> . <i>Genome Announcements</i> , 2015, 3, .	0.8	10
43	Draft Genome Sequence of the <i>Xylella fastidiosa</i> CoDiRO Strain. <i>Genome Announcements</i> , 2015, 3, .	0.8	51
44	Discovery and molecular characterization of a new cryptovirus dsRNA genome from Japanese persimmon through conventional cloning and high-throughput sequencing. <i>Virus Genes</i> , 2015, 50, 160-164.	1.6	16
45	Identification and characterization of a viroid resembling apple dimple fruit viroid in fig (<i>Ficus carica</i>) Tj ETQq1 1 0.784314 rgBT /Over 2.2 45		
46	Semi-artificial datasets as a resource for validation of bioinformatics pipelines for plant virus detection. , 0, 1, .		8