

Giuliana Loconsole

List of Publications by Year in descending order

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Version: 2024-02-01

21
papers

883
citations

687363

13
h-index

752698

20
g-index

21
all docs

21
docs citations

21
times ranked

938
citing authors

#	ARTICLE	IF	CITATIONS
1	Infectivity and Transmission of <i>Xylella fastidiosa</i> by <i>Philaenus spumarius</i> (Hemiptera: Aphrophoridae) in Apulia, Italy. <i>Journal of Economic Entomology</i> , 2014, 107, 1316-1319.	1.8	152
2	Spittlebugs as vectors of <i>Xylella fastidiosa</i> in olive orchards in Italy. <i>Journal of Pest Science</i> , 2017, 90, 521-530.	3.7	131
3	Identification of a single-stranded DNA virus associated with citrus chlorotic dwarf disease, a new member in the family Geminiviridae. <i>Virology</i> , 2012, 432, 162-172.	2.4	130
4	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
5	Genome-Wide Analysis Provides Evidence on the Genetic Relatedness of the Emergent <i>Xylella fastidiosa</i> Genotype in Italy to Isolates from Central America. <i>Phytopathology</i> , 2017, 107, 816-827.	2.2	61
6	Draft Genome Sequence of the <i>Xylella fastidiosa</i> CoDiRO Strain. <i>Genome Announcements</i> , 2015, 3, .	0.8	51
7	Complete Genome Sequence of the Olive-Infecting Strain <i>Xylella fastidiosa</i> subsp. <i>pauca</i> De Donno. <i>Genome Announcements</i> , 2017, 5, .	0.8	34
8	A new variant of <i>Xylella fastidiosa</i> subspecies <i>multiplex</i> detected in different host plants in the recently emerged outbreak in the region of Tuscany, Italy. <i>European Journal of Plant Pathology</i> , 2019, 154, 1195-1200.	1.7	32
9	Validation of high-throughput real time polymerase chain reaction assays for simultaneous detection of invasive citrus pathogens. <i>Journal of Virological Methods</i> , 2013, 193, 478-486.	2.1	28
10	Development of real-time PCR based assays for simultaneous and improved detection of citrus viruses. <i>European Journal of Plant Pathology</i> , 2010, 128, 251-259.	1.7	27
11	Pilot project on <i>Xylella fastidiosa</i> to reduce risk assessment uncertainties. <i>EFSA Supporting Publications</i> , 2016, 13, 1013E.	0.7	23
12	Identification and characterization of privet leaf blotch-associated virus, a novel <i>idaeovirus</i> . <i>Molecular Plant Pathology</i> , 2017, 18, 925-936.	4.2	22
13	Rapid differentiation of citrus Hop stunt viroid variants by real-time RT-PCR and high resolution melting analysis. <i>Molecular and Cellular Probes</i> , 2013, 27, 221-229.	2.1	18
14	<i>Olea Europaea</i> Geminivirus: A Novel Bipartite Geminivirid Infecting Olive Trees. <i>Viruses</i> , 2021, 13, 481.	3.3	16
15	Introduction and adaptation of an emerging pathogen to olive trees in Italy. <i>Microbial Genomics</i> , 2021, 7, .	2.0	14
16	Draft Genome Sequence of CO33, a Coffee-Infecting Isolate of <i>Xylella fastidiosa</i> . <i>Genome Announcements</i> , 2015, 3, .	0.8	10
17	Isolation and Partial Characterization of a Novel Cytorhabdovirus from Citrus Trees Showing Foliar Symptoms in Iran. <i>Plant Disease</i> , 2016, 100, 66-71.	1.4	8
18	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

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19	Detection of Citrus tristeza virus and Coinfecting Viroids. <i>Methods in Molecular Biology</i> , 2019, 2015, 67-78.	0.9	2
20	High throughput sequencing from Angolan citrus accessions discloses the presence of emerging CTV strains. <i>Virology Journal</i> , 2021, 18, 62.	3.4	1
21	DEEP SEQUENCING OF SMALL RNAs FROM CITRUS AFFECTED BY GRAFT-TRANSMISSIBLE DISEASES OF UNKNOWN AETIOLOGY LEADS TO DISCOVERY OF TWO NOVEL VIRUSES. <i>Acta Horticulturae</i> , 2015, , 817-824.	0.2	0