

Laura Miozzi

List of Publications by Year in descending order

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Version: 2024-02-01

45
papers

1,985
citations

257450

24
h-index

243625

44
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46
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46
docs citations

46
times ranked

2429
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of grapevine microRNAs and their targets using high throughput sequencing and degradome analysis. <i>Plant Journal</i> , 2010, 62, 960-76.	5.7	335
2	Global and cell-type gene expression profiles in tomato plants colonized by an arbuscular mycorrhizal fungus. <i>New Phytologist</i> , 2009, 184, 975-987.	7.3	187
3	From root to fruit: RNA-Seq analysis shows that arbuscular mycorrhizal symbiosis may affect tomato fruit metabolism. <i>BMC Genomics</i> , 2014, 15, 221.	2.8	149
4	Deep sequencing analysis of viral short RNAs from an infected Pinot Noir grapevine. <i>Virology</i> , 2010, 408, 49-56.	2.4	109
5	Novel functional microRNAs from virus-free and infected <i>Vitis vinifera</i> plants under water stress. <i>Scientific Reports</i> , 2016, 6, 20167.	3.3	81
6	Transcriptomics of the Interaction between the Monopartite Phloem-Limited Geminivirus Tomato Yellow Leaf Curl Sardinia Virus and <i>Solanum lycopersicum</i> Highlights a Role for Plant Hormones, Autophagy and Plant Immune System Fine Tuning during Infection. <i>PLoS ONE</i> , 2014, 9, e89951.	2.5	77
7	Genome-wide identification of viral and host transcripts targeted by viral siRNAs in <i>Vitis vinifera</i> . <i>Molecular Plant Pathology</i> , 2013, 14, 30-43.	4.2	69
8	Comparative Analysis of Expression Profiles in Shoots and Roots of Tomato Systemically Infected by Tomato spotted wilt virus Reveals Organ-Specific Transcriptional Responses. <i>Molecular Plant-Microbe Interactions</i> , 2009, 22, 1504-1513.	2.6	64
9	The arbuscular mycorrhizal symbiosis attenuates symptom severity and reduces virus concentration in tomato infected by Tomato yellow leaf curl Sardinia virus (TYLCSV). <i>Mycorrhiza</i> , 2014, 24, 179-186.	2.8	61
10	In silico analysis of fungal small RNA accumulation reveals putative plant mRNA targets in the symbiosis between an arbuscular mycorrhizal fungus and its host plant. <i>BMC Genomics</i> , 2019, 20, 169.	2.8	60
11	Nondestructive Raman Spectroscopy as a Tool for Early Detection and Discrimination of the Infection of Tomato Plants by Two Economically Important Viruses. <i>Analytical Chemistry</i> , 2019, 91, 9025-9031.	6.5	57
12	Two new natural begomovirus recombinants associated with the tomato yellow leaf curl disease co-exist with parental viruses in tomato epidemics in Italy. <i>Virus Research</i> , 2009, 143, 15-23.	2.2	56
13	Identification of grapevine microRNAs and their targets using high-throughput sequencing and degradome analysis. <i>Plant Journal</i> , 2010, 62, no-no.	5.7	53
14	Arbuscular Mycorrhizal Symbiosis: Plant Friend or Foe in the Fight Against Viruses?. <i>Frontiers in Microbiology</i> , 2019, 10, 1238.	3.5	52
15	Bacterial and fungal communities associated with <i>Tuber magnatum</i> -productive niches. <i>Plant Biosystems</i> , 2010, 144, 323-332.	1.6	45
16	Bioinformatics approaches for viral metagenomics in plants using short RNAs: model case of study and application to a <i>Cicer arietinum</i> population. <i>Frontiers in Microbiology</i> , 2014, 5, 790.	3.5	42
17	Analysis of small RNAs derived from tomato yellow leaf curl Sardinia virus reveals a cross reaction between the major viral hotspot and the plant host genome. <i>Virus Research</i> , 2013, 178, 287-296.	2.2	39
18	Phospholipase A2 up-regulation during mycorrhiza formation in <i>Tuber borchii</i> . <i>New Phytologist</i> , 2005, 167, 229-238.	7.3	38

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19	A Primer on the Analysis of High-Throughput Sequencing Data for Detection of Plant Viruses. <i>Microorganisms</i> , 2021, 9, 841.	3.6	36
20	Recombination profiles between Tomato yellow leaf curl virus and Tomato yellow leaf curl Sardinia virus in laboratory and field conditions: evolutionary and taxonomic implications. <i>Journal of General Virology</i> , 2012, 93, 2712-2717.	2.9	34
21	Small RNA profiles of wild-type and silencing suppressor-deficient tomato spotted wilt virus infected <i>Nicotiana benthamiana</i> . <i>Virus Research</i> , 2015, 208, 30-38.	2.2	34
22	Arbuscular Mycorrhizal Symbiosis Limits Foliar Transcriptional Responses to Viral Infection and Favors Long-Term Virus Accumulation. <i>Molecular Plant-Microbe Interactions</i> , 2011, 24, 1562-1572.	2.6	33
23	Pyramiding <i>Ty1</i> and <i>Ty3</i> and <i>Ty2</i> in tomato hybrids dramatically inhibits symptom expression and accumulation of tomato yellow leaf curl disease inducing viruses. <i>Archives of Phytopathology and Plant Protection</i> , 2017, 50, 213-227.	1.3	33
24	The Induction of an Effective dsRNA-Mediated Resistance Against Tomato Spotted Wilt Virus by Exogenous Application of Double-Stranded RNA Largely Depends on the Selection of the Viral RNA Target Region. <i>Frontiers in Plant Science</i> , 2020, 11, 533338.	3.6	28
25	Arbuscular Mycorrhizal Symbiosis Primes Tolerance to Cucumber Mosaic Virus in Tomato. <i>Viruses</i> , 2020, 12, 675.	3.3	23
26	Different Genetic Sources Contribute to the Small RNA Population in the Arbuscular Mycorrhizal Fungus <i>Gigaspora margarita</i> . <i>Frontiers in Microbiology</i> , 2020, 11, 395.	3.5	23
27	Comparison of small RNA profiles in <i>Nicotiana benthamiana</i> and <i>Solanum lycopersicum</i> infected by <i>Polygonum ringspot tospovirus</i> reveals host-specific responses to viral infection. <i>Virus Research</i> , 2016, 211, 38-45.	2.2	21
28	Functional Annotation and Identification of Candidate Disease Genes by Computational Analysis of Normal Tissue Gene Expression Data. <i>PLoS ONE</i> , 2008, 3, e2439.	2.5	20
29	Deep Sequencing Data and Infectivity Assays Indicate that Chickpea Chlorotic Dwarf Virus is the Etiological Agent of the "Hard Fruit Syndrome" of Watermelon. <i>Viruses</i> , 2017, 9, 311.	3.3	18
30	A Short Indel-Lacking-Resistance Gene Triggers Silencing of the Photosynthetic Machinery Components Through TYLCSV-Associated Endogenous siRNAs in Tomato. <i>Frontiers in Plant Science</i> , 2018, 9, 1470.	3.6	15
31	Real-Time PCR Protocols for the Quantification of the Begomovirus Tomato Yellow Leaf Curl Sardinia Virus in Tomato Plants and in Its Insect Vector. <i>Methods in Molecular Biology</i> , 2015, 1236, 61-72.	0.9	13
32	The first complete genome sequences of two distinct European tomato spotted wilt virus isolates. <i>Archives of Virology</i> , 2015, 160, 591-595.	2.1	13
33	Impact of high or low levels of phosphorus and high sodium in soils on productivity and stress tolerance of <i>Arundo donax</i> plants. <i>Plant Science</i> , 2019, 289, 110260.	3.6	13
34	The complete genome sequence of <i>Polygonum ringspot virus</i> . <i>Archives of Virology</i> , 2014, 159, 3149-3152.	2.1	9
35	In silico prediction of miRNAs targeting ToLCV and their regulation in susceptible and resistant tomato plants. <i>Australasian Plant Pathology</i> , 2017, 46, 379-386.	1.0	9
36	The interaction of <i>Lolium latent virus</i> major coat protein with ankyrin repeat protein NbANKr redirects it to chloroplasts and modulates virus infection. <i>Journal of General Virology</i> , 2018, 99, 730-742.	2.9	9

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37	No Evidence for Seed Transmission of Tomato Yellow Leaf Curl Sardinia Virus in Tomato. <i>Cells</i> , 2021, 10, 1673.	4.1	8
38	Drawing siRNAs of Viral Origin Out from Plant siRNAs Libraries. <i>Methods in Molecular Biology</i> , 2015, 1236, 111-123.	0.9	5
39	ORTom: a multi-species approach based on conserved co-expression to identify putative functional relationships among genes in tomato. <i>Plant Molecular Biology</i> , 2010, 73, 519-532.	3.9	4
40	First Report of Grapevine Latent Viroid Infecting Grapevine (<i>Vitis vinifera</i>) in Italy. <i>Plant Disease</i> , 2018, 102, 1672.	1.4	3
41	Modulation of class III peroxidase pathways and phenylpropanoids in <i>Arundo donax</i> under salt and phosphorus stress. <i>Plant Physiology and Biochemistry</i> , 2022, 183, 151-159.	5.8	3
42	The complete nucleotide sequence of an isolate of Tomato yellow leaf curl Sardinia virus found in Sicily. <i>Archives of Virology</i> , 2010, 155, 1539-1542.	2.1	1
43	Evidence of new viruses infecting freesia hybrids showing necrotic disease. <i>Acta Horticulturae</i> , 2018, , 21-28.	0.2	1
44	Identification of Known and Novel <i>Arundo donax</i> L. MicroRNAs and Their Targets Using High-Throughput Sequencing and Degradome Analysis. <i>Life</i> , 2022, 12, 651.	2.4	1
45	Women in the European Virus Bioinformatics Center. <i>Viruses</i> , 2022, 14, 1522.	3.3	1