

Sabrina Palmano

List of Publications by Year in descending order

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34
papers

908
citations

393982

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#	ARTICLE	IF	CITATIONS
1	Silencing of ATP Synthase \hat{I}^2 Impairs Egg Development in the Leafhopper <i>Scaphoideus titanus</i> , Vector of the Phytoplasma Associated with Grapevine Flavescence DorÃ©e. <i>International Journal of Molecular Sciences</i> , 2022, 23, 765.	1.8	4
2	Silencing of ATP synthase \hat{I}^2 reduces phytoplasma multiplication in a leafhopper vector. <i>Journal of Insect Physiology</i> , 2021, 128, 104176.	0.9	7
3	Recovery from Grapevine Flavescence DorÃ©e in Areas of High Infection Pressure. <i>Agronomy</i> , 2020, 10, 1479.	1.3	4
4	Molecular memory of Flavescence dorÃ©e phytoplasma in recovering grapevines. <i>Horticulture Research</i> , 2020, 7, 126.	2.9	17
5	Biological characterization of <i>Euscelidius variegatus</i> iflavirus 1. <i>Journal of Invertebrate Pathology</i> , 2020, 173, 107370.	1.5	5
6	Differential gene expression in two grapevine cultivars recovered from â€œflavescence dorÃ©eâ€. <i>Microbiological Research</i> , 2019, 220, 72-82.	2.5	7
7	Transcriptomic Analyses of Phytoplasmas. <i>Methods in Molecular Biology</i> , 2019, 1875, 239-251.	0.4	1
8	Towards the identification of genes involved in resistance/tolerance to â€œflavescence dorÃ©eâ€. <i>Phytopathogenic Mollicutes</i> , 2019, 9, 223.	0.1	0
9	RNAi silencing to validate the role of insect genes in phytoplasma transmission. <i>Phytopathogenic Mollicutes</i> , 2019, 9, 135.	0.1	0
10	miRVIT: A Novel miRNA Database and Its Application to Uncover <i>Vitis</i> Responses to Flavescence dorÃ©e Infection. <i>Frontiers in Plant Science</i> , 2018, 9, 1034.	1.7	26
11	Dissecting interplays between <i>Vitis vinifera</i> L. and grapevine virus B (GVB) under field conditions. <i>Molecular Plant Pathology</i> , 2018, 19, 2651-2666.	2.0	26
12	Identification of putative effector genes and their transcripts in three strains related to â€ Candidatus <i>Phytoplasma aurantifolia</i> â€. <i>Microbiological Research</i> , 2017, 199, 57-66.	2.5	18
13	Transmission of <i>Penicillium aurantiogriseum</i> partitiâ€like virus 1 to a new fungal host (<i>Cryphonectria parasitica</i>) confers higher resistance to salinity and reveals adaptive genomic changes. <i>Environmental Microbiology</i> , 2017, 19, 4480-4492.	1.8	56
14	Structural modification of cuminaldehyde thiosemicarbazone increases inhibition specificity toward aflatoxin biosynthesis and sclerotia development in <i>Aspergillus flavus</i> . <i>Applied Microbiology and Biotechnology</i> , 2017, 101, 6683-6696.	1.7	17
15	Space-Time Point Pattern Analysis of Flavescence DorÃ©e Epidemic in a Grapevine Field: Disease Progression and Recovery. <i>Frontiers in Plant Science</i> , 2016, 7, 1987.	1.7	34
16	Decreasing Global Transcript Levels over Time Suggest that Phytoplasma Cells Enter Stationary Phase during Plant and Insect Colonization. <i>Applied and Environmental Microbiology</i> , 2015, 81, 2591-2602.	1.4	33
17	Diagnosis of Phytoplasmas by Real-Time PCR Using Locked Nucleic Acid (LNA) Probes. <i>Methods in Molecular Biology</i> , 2015, 1302, 113-122.	0.4	4
18	RNA-Seq profile of flavescence dorÃ©e phytoplasma in grapevine. <i>BMC Genomics</i> , 2014, 15, 1088.	1.2	34

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19	A DNA Origami Nanorobot Controlled by Nucleic Acid Hybridization. <i>Small</i> , 2014, 10, 2918-2926.	5.2	47
20	Metabolic and transcript analysis of the flavonoid pathway in diseased and recovered Nebbiolo and Barbera grapevines (<i>Vitis</i> spp.) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50,702 Td (v Cell and Environment, 2014, 37, 2183-2200.	2.8	57
21	Genome wide sequence analysis grants unbiased definition of species boundaries in <i>Candidatus Phytoplasma</i> . <i>Systematic and Applied Microbiology</i> , 2013, 36, 539-548.	1.2	28
22	Hydrogen Peroxide Accumulation and Transcriptional Changes in Grapevines Recovered from Flavescence dorée Disease. <i>Phytopathology</i> , 2013, 103, 776-784.	1.1	48
23	Novel aspects of grapevine response to phytoplasma infection investigated by a proteomic and phospho-proteomic approach with data integration into functional networks. <i>BMC Genomics</i> , 2013, 14, 38.	1.2	94
24	Genome drafts of four phytoplasma strains of the ribosomal group 16SrIII. <i>Microbiology (United)</i> Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 59	0.7	59
25	Quantitation of Grapevine leafroll associated virus-1 and -3, Grapevine virus A, Grapevine fanleaf virus and Grapevine fleck virus in field-collected <i>Vitis vinifera</i> L. "Nebbiolo"™ by real-time reverse transcription-PCR. <i>Journal of Virological Methods</i> , 2011, 172, 1-7.	1.0	38
26	On the alleged origin of geminiviruses from extrachromosomal DNAs of phytoplasmas. <i>BMC Evolutionary Biology</i> , 2011, 11, 185.	3.2	19
27	Response of the <i>Vitis vinifera</i> L. cv. "Nebbiolo"™ proteome to Flavescence dorée phytoplasma infection. <i>Proteomics</i> , 2011, 11, 212-224.	1.3	67
28	Cloning of the Glycerinaldehyde 3-phosphate Dehydrogenase Gene of Flavescence dorée Phytoplasma and Development of Serological and Molecular Tools for Studying its Expression. <i>Journal of Phytopathology</i> , 2010, 158, 382-386.	0.5	1
29	Detection of Flavescence dorée and Bois noir phytoplasmas, Grapevine leafroll associated virus-1 and -3 and Grapevine virus A from the same crude extract by reverse transcription-RealTime Taqman assays. <i>Plant Pathology</i> , 2009, 58, 838-845.	1.2	37
30	Detection of Flavescence Dorée Phytoplasma in Grapevine by Reverse-Transcription PCR. <i>Plant Disease</i> , 2007, 91, 1496-1501.	0.7	26
31	Characterization of Four Viral Species Belonging to the Family Potyviridae Isolated from <i>Ranunculus asiaticus</i> . <i>Phytopathology</i> , 2006, 96, 560-566.	1.1	37
32	Title is missing!. <i>European Journal of Plant Pathology</i> , 2003, 109, 817-825.	0.8	17
33	Development of a PCR test for the detection of <i>Curtobacterium flaccumfaciens</i> pv. <i>flaccumfaciens</i> . <i>Antonie Van Leeuwenhoek</i> , 2001, 80, 1-10.	0.7	29
34	Diversity of phytoplasmas isolated from insects, determined by a DNA heteroduplex mobility assay and a length polymorphism of the 16S-23S rDNA spacer region analysis. <i>Journal of Applied Microbiology</i> , 2000, 89, 744-750F.	1.4	11