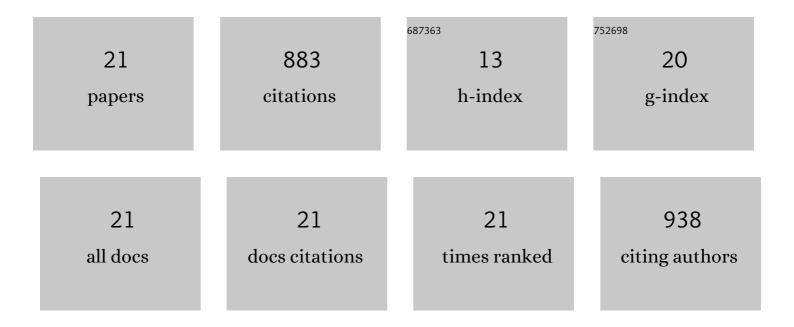
Giuliana Loconsole

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4863483/publications.pdf Version: 2024-02-01



| # | Article | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Infectivity and Transmission of <i>Xylella fastidiosa</i> by <i>Philaenus spumarius</i> (Hemiptera: Aphrophoridae) in Apulia, Italy. Journal of Economic Entomology, 2014, 107, 1316-1319. | 1.8 | 152 |
| 2 | Spittlebugs as vectors of Xylella fastidiosa in olive orchards in Italy. Journal of Pest Science, 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. Virology, 2012, 432, 162-172. | 2.4 | 130 |
| 4 | Transcriptome profiling of two olive cultivars in response to infection by the CoDiRO strain of Xylella fastidiosa subsp. pauca. BMC Genomics, 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. Phytopathology, 2017, 107, 816-827. | 2.2 | 61 |
| 6 | Draft Genome Sequence of the Xylella fastidiosa CoDiRO Strain. Genome Announcements, 2015, 3, . | 0.8 | 51 |
| 7 | Complete Genome Sequence of the Olive-Infecting Strain Xylella fastidiosa subsp. <i>pauca</i> De Donno. Genome Announcements, 2017, 5, . | 0.8 | 34 |
| 8 | A new variant of Xylella fastidiosa subspecies multiplex detected in different host plants in the recently emerged outbreak in the region of Tuscany, Italy. European Journal of Plant Pathology, 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. Journal of Virological Methods, 2013, 193, 478-486. | 2.1 | 28 |
| 10 | Development of real-time PCR based assays for simultaneous and improved detection of citrus viruses. European Journal of Plant Pathology, 2010, 128, 251-259. | 1.7 | 27 |
| 11 | Pilot project on Xylella fastidiosa to reduce risk assessment uncertainties. EFSA Supporting Publications, 2016, 13, 1013E. | 0.7 | 23 |
| 12 | Identification and characterization of privet leaf blotchâ€associated virus, a novel <i>idaeovirus</i> . Molecular Plant Pathology, 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. Molecular and Cellular Probes, 2013, 27, 221-229. | 2.1 | 18 |
| 14 | Olea Europaea Geminivirus: A Novel Bipartite Geminivirid Infecting Olive Trees. Viruses, 2021, 13, 481. | 3.3 | 16 |
| 15 | Introduction and adaptation of an emerging pathogen to olive trees in Italy. Microbial Genomics, 2021, 7, . | 2.0 | 14 |
| 16 | Draft Genome Sequence of CO33, a Coffee-Infecting Isolate of Xylella fastidiosa. Genome Announcements, 2015, 3, . | 0.8 | 10 |
| 17 | Isolation and Partial Characterization of a Novel Cytorhabdovirus from Citrus Trees Showing Foliar Symptoms in Iran. Plant Disease, 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 Jasminum officinale in Italy. Plants, 2022, 11, 309. | 3.5 | 5 |

| # | Article | IF | CITATIONS |
|----|--|-----|-----------|
| 19 | Detection of Citrus tristeza virus and Coinfecting Viroids. Methods in Molecular Biology, 2019, 2015, 67-78. | 0.9 | 2 |
| 20 | High throughput sequencing from Angolan citrus accessions discloses the presence of emerging CTV strains. Virology Journal, 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. Acta Horticulturae, 2015, , 817-824. | 0.2 | 0 |