

# Annalisa Giampetruzzi

## List of Publications by Year in descending order

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Version: 2024-02-01

33  
papers

1,438  
citations

361413

20  
h-index

414414

32  
g-index

33  
all docs

33  
docs citations

33  
times ranked

1413  
citing authors

| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 1  | <i>Xylella fastidiosa</i> ™s relationships: the bacterium, the host plants, and the plant microbiome. <i>New Phytologist</i> , 2022, 234, 1598-1605.  | 7.3 | 17        |
| 2  | Introduction and adaptation of an emerging pathogen to olive trees in Italy. <i>Microbial Genomics</i> , 2021, 7, .   | 2.0 | 14        |
| 3  | Grapevine Pinot gris virus variants in vines with chlorotic mottling and leaf deformation. <i>Journal of Plant Pathology</i> , 2020, 102, 531-531.  | 1.2 | 8         |
| 4  | Emergence of a Plant Pathogen in Europe Associated with Multiple Intercontinental Introductions. <i>Applied and Environmental Microbiology</i> , 2020, 86, .  | 3.1 | 57        |
| 5  | Coding-Complete Genome Sequence of a <i>Black Queen Cell Virus</i> Isolate from Honey Bees ( <i>Apis</i> ) Tj ETQq1 1 0,784314,3gBT /Over   | 0,6 |           |
| 6  | Differences in the Endophytic Microbiome of Olive Cultivars Infected by <i>Xylella fastidiosa</i> across Seasons. <i>Pathogens</i> , 2020, 9, 723.  | 2.8 | 39        |
| 7  | Draft Genome Sequence Resources of Three Strains (TOS4, TOS5, and TOS14) of <i>Xylella fastidiosa</i> Infecting Different Host Plants in the Newly Discovered Outbreak in Tuscany, Italy. <i>Phytopathology</i> , 2019, 109, 1516-1518. | 2.2 | 11        |
| 8  | Infections of the <i>Xylella fastidiosa</i> subsp. <i>pauca</i> Strain "De Donno" in Alfalfa ( <i>Medicago sativa</i> ) Elicits an Overactive Immune Response. <i>Plants</i> , 2019, 8, 335.  | 3.5 | 12        |
| 9  | Draft Genome Resources of Two Strains ("ESVL" and "VIA5901") of <i>Xylella fastidiosa</i> Associated with Almond Leaf Scorch Disease in Alicante, Spain. <i>Phytopathology</i> , 2019, 109, 219-221.                                    | 2.2 | 24        |
| 10 | Identification and Characterization of Resistance-Breaking (RB) Isolates of Citrus tristeza virus. <i>Methods in Molecular Biology</i> , 2019, 2015, 105-126.   | 0.9 | 1         |
| 11 | Ionic Differences between Susceptible and Resistant Olive Cultivars Infected by <i>Xylella fastidiosa</i> in the Outbreak Area of Salento, Italy. <i>Pathogens</i> , 2019, 8, 272.  | 2.8 | 37        |
| 12 | Small RNA Isolation from Tissues of Grapevine and Woody Plants. <i>Methods in Molecular Biology</i> , 2018, 1746, 27-36.  | 0.9 | 2         |
| 13 | Molecular and biological characterization of a novel mild strain of citrus tristeza virus in California. <i>Archives of Virology</i> , 2018, 163, 1795-1804.  | 2.1 | 31        |
| 14 | Draft Genome Sequence of <i>Xylella fastidiosa</i> subsp. <i>fastidiosa</i> Strain VIA5235, Isolated from <i>Prunus avium</i> in Mallorca Island, Spain. <i>Microbiology Resource Announcements</i> , 2018, 7, .                        | 0.6 | 13        |
| 15 | Identification and Characterization of <i>Citrus tristeza virus</i> Isolates Breaking Resistance in Trifoliate Orange in California. <i>Phytopathology</i> , 2017, 107, 901-908.  | 2.2 | 33        |
| 16 | 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        |
| 17 | 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        |
| 18 | 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        |

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|----|---|-----|-----------|
| 19 | Identification and characterization of an isolate of apple green crinkle associated virus involved in a severe disease of quince ( <i>Cydonia oblonga</i> , Mill.). <i>Archives of Virology</i> , 2017, 162, 299-306. | 2.1 | 25        |
| 20 | 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         |
| 21 | 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       |
| 22 | Transmission of grapevine Pinot gris virus by <i>Colomerus vitis</i> (Acari: Eriophyidae) to grapevine. <i>Archives of Virology</i> , 2016, 161, 2595-2599.   | 2.1 | 60        |
| 23 | 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         |
| 24 | Draft Genome Sequence of CO33, a Coffee-Infecting Isolate of <i>Xylella fastidiosa</i> . <i>Genome Announcements</i> , 2015, 3, .   | 0.8 | 10        |
| 25 | Draft Genome Sequence of the <i>Xylella fastidiosa</i> CoDiRO Strain. <i>Genome Announcements</i> , 2015, 3, .  | 0.8 | 51        |
| 26 | Genetic Variability of <i>Grapevine Pinot gris virus</i> and Its Association with Grapevine Leaf Mottling and Deformation. <i>Phytopathology</i> , 2015, 105, 555-563.  | 2.2 | 79        |
| 27 | Deep sequencing and analysis of small RNAs in sweet orange grafted on sour orange infected with two citrus tristeza virus isolates prevalent in Sicily. <i>Archives of Virology</i> , 2015, 160, 2583-2589.           | 2.1 | 28        |
| 28 | Roles and Programming of Arabidopsis ARGONAUTE Proteins during Turnip Mosaic Virus Infection. <i>PLoS Pathogens</i> , 2015, 11, e1004755.   | 4.7 | 175       |
| 29 | Deep-sequencing analysis of an apricot tree with vein clearing symptoms reveals the presence of a novel betaflexivirus. <i>Virus Research</i> , 2014, 181, 1-5.   | 2.2 | 27        |
| 30 | Identification and Characterization of <i>Citrus yellow vein clearing virus</i> , A Putative New Member of the Genus <i>Mandarivirus</i> . <i>Phytopathology</i> , 2012, 102, 1168-1175.                              | 2.2 | 90        |
| 31 | A new grapevine virus discovered by deep sequencing of virus- and viroid-derived small RNAs in Cv Pinot gris. <i>Virus Research</i> , 2012, 163, 262-268.   | 2.2 | 227       |
| 32 | Deep sequencing analysis of viral short RNAs from an infected Pinot Noir grapevine. <i>Virology</i> , 2010, 408, 49-56.   | 2.4 | 109       |
| 33 | An assay for the detection of grapevine leafroll-associated virus 3 using a single-chain fragment variable antibody. <i>Archives of Virology</i> , 2009, 154, 19-26.  | 2.1 | 11        |