

Toufic Elbeaino

List of Publications by Year in descending order

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59
papers

1,919
citations

361296

20
h-index

276775

41
g-index

71
all docs

71
docs citations

71
times ranked

2047
citing authors

#	ARTICLE	IF	CITATIONS
1	First detection and molecular characterization of grapevine phytoplasmas in Kosovo. <i>Journal of Phytopathology</i> , 2022, 170, 34-40.	0.5	0
2	Biological, epidemiological and population structure analyses of vitiviruses in Iran. <i>European Journal of Plant Pathology</i> , 2021, 159, 117-129.	0.8	2
3	Fimoviruses (Fimoviridae). , 2021, , 396-404.		1
4	Maize (<i>Zea mays</i> L.): A New Host for <i>Ligustrum witchesâ€™ Broom Phytoplasma</i> . <i>Pathogens</i> , 2021, 10, 723.	1.2	2
5	Development of singleplex and multiplex real-time (Taqman ^Â) RT-PCR assays for the detection of viruses associated with fig mosaic disease. <i>Journal of Virological Methods</i> , 2021, 293, 114145.	1.0	8
6	<i>Pistacia vera</i> L. hosts pistachio ampelovirus A in Tunisia. <i>Journal of Plant Pathology</i> , 2021, 103, 1335.	0.6	2
7	2021 Taxonomic update of phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. <i>Archives of Virology</i> , 2021, 166, 3513-3566.	0.9	62
8	First detection of <i>Candidatus</i> <i>Phytoplasma asteris</i> and <i>Candidatus</i> <i>Phytoplasma solani</i> related strains in fig trees. <i>Journal of Phytopathology</i> , 2020, 168, 63-71.	0.5	5
9	Generation and molecular docking analysis of specific single-chain variable fragments selected by phage display against the recombinant nucleocapsid protein of fig mosaic virus. <i>Journal of Virological Methods</i> , 2020, 276, 113796.	1.0	6
10	Development of an FTP-LAMP assay based on TaqMan real-time PCR and LAMP for the specific detection of <i>Xylella fastidiosa</i> De Donno and mulberry strains in both plants and insect vectors. <i>Journal of Microbiological Methods</i> , 2020, 175, 105992.	0.7	5
11	2020 taxonomic update for phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. <i>Archives of Virology</i> , 2020, 165, 3023-3072.	0.9	184
12	Complete Sequence, Genome Organization and Molecular Detection of Grapevine Line Pattern Virus, a New Putative Anulavirus Infecting Grapevine. <i>Viruses</i> , 2020, 12, 602.	1.5	4
13	Detection and Phylogenetic Analysis of Viruses Linked with Fig Mosaic Disease in Seventeen Fig Cultivars in Palestine. <i>Plant Pathology Journal</i> , 2020, 36, 267-279.	0.7	5
14	The Detection of viruses in olive cultivars in Greece, using a rapid and effective RNA extraction method, for certification of virus-tested propagation material. <i>Phytopathologia Mediterranea</i> , 2020, 59, 203-211.	0.6	15
15	Development of RT-PCR assays for the detection and the resultant phylogenetic analysis of four grapevine vitiviruses based on the coat protein sequences. <i>Journal of Virological Methods</i> , 2019, 273, 113712.	1.0	1
16	Taxonomy of the order Bunyavirales: second update 2018. <i>Archives of Virology</i> , 2019, 164, 927-941.	0.9	115
17	Taxonomy of the order Bunyavirales: update 2019. <i>Archives of Virology</i> , 2019, 164, 1949-1965.	0.9	285
18	Detection and seed transmission of Bermudagrass phytoplasma in maize in Turkey. <i>Journal of Phytopathology</i> , 2019, 167, 248-255.	0.5	15

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19	Production of a polyclonal antiserum against recombinant nucleocapsid protein and its application for the detection of fig mosaic virus. <i>Journal of Virological Methods</i> , 2019, 265, 22-25.	1.0	9
20	First detection of grapevine virus F in Iran. <i>Journal of Plant Pathology</i> , 2018, 100, 111-112.	0.6	3
21	Taxonomy of the family Arenaviridae and the order Bunyavirales: update 2018. <i>Archives of Virology</i> , 2018, 163, 2295-2310.	0.9	157
22	Effect of <i>Cucumber mosaic virus</i> (<i>CMV</i>) infection on antineoplastic alkaloids from periwinkle (<i>Catharanthus roseus</i> L.) cultured in the Mecca region and resistance induction by plant-growth-promoting rhizobacteria (PGPR). <i>Biotechnology and Biotechnological Equipment</i> , 2018, 32, 49-57.	0.5	6
23	High-Throughput Sequencing Reveals <i>Cyclamen persicum</i> Mill. as a Natural Host for Fig Mosaic Virus. <i>Viruses</i> , 2018, 10, 684.	1.5	16
24	First report on the occurrence of grapevine leafroll-associated virus 7 in Morocco. <i>Journal of Plant Pathology</i> , 2018, 100, 355-355.	0.6	1
25	ICTV Virus Taxonomy Profile: Fimoviridae. <i>Journal of General Virology</i> , 2018, 99, 1478-1479.	1.3	90
26	Inhibitory activity of different medicinal extracts from Thuja leaves, ginger roots, Harmal seeds and turmeric rhizomes against Fig leaf mottle-associated virus 1 (FLMaV-1) infecting figs in Mecca region. <i>Saudi Journal of Biological Sciences</i> , 2017, 24, 936-944.	1.8	11
27	Next generation sequencing and molecular analysis of artichoke Italian latent virus. <i>Archives of Virology</i> , 2017, 162, 1805-1809.	0.9	3
28	Viruses infecting different Mediterranean genotypes of <i>Ficus carica</i> and their distribution in different plant organs. <i>Acta Horticulturae</i> , 2017, , 273-278.	0.1	1
29	First Report on the Occurrence of <i>Grapevine leafroll-associated virus-4</i> Strain 6 in Lebanon. <i>Plant Disease</i> , 2017, 101, 1066-1066.	0.7	3
30	Pittosporum cryptic virus 1: genome sequence completion using next-generation sequencing. <i>Archives of Virology</i> , 2016, 161, 2039-2042.	0.9	7
31	Occurrence of Deformed wing virus, Chronic bee paralysis virus and mtDNA variants in haplotype K of <i>Varroa destructor</i> mites in Syrian apiaries. <i>Experimental and Applied Acarology</i> , 2016, 69, 11-19.	0.7	6
32	Unusual genomic features of a badnavirus infecting mulberry. <i>Journal of General Virology</i> , 2016, 97, 3073-3087.	1.3	19
33	Deep sequencing of dsRNAs recovered from mosaic-diseased pigeonpea reveals the presence of a novel emaravirus: pigeonpea sterility mosaic virus 2. <i>Archives of Virology</i> , 2015, 160, 2019-2029.	0.9	62
34	The sequencing of the complete genome of a Tomato black ring virus (TBRV) and of the RNA2 of three Grapevine chrome mosaic virus (GCMV) isolates from grapevine reveals the possible recombinant origin of GCMV. <i>Virus Genes</i> , 2015, 50, 165-171.	0.7	21
35	Specific Detection and Identification of American Mulberry-Infecting and Italian Olive-Associated Strains of <i>Xylella fastidiosa</i> by Polymerase Chain Reaction. <i>PLoS ONE</i> , 2015, 10, e0129330.	1.1	12
36	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.	1.1	27

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37	Phylogenetic and recombination analysis of the homing protein domain of grapevine fanleaf virus (GFLV) isolates associated with "yellow mosaic"™ and "infectious malformation"™ syndromes in grapevine. <i>Archives of Virology</i> , 2014, 159, 2757-2764.		16
38	Deep sequencing of pigeonpea sterility mosaic virus discloses five RNA segments related to emaraviruses. <i>Virus Research</i> , 2014, 188, 27-31.	1.1	67
39	A novel phytoplasma associated with witches'™ broom disease of <i>Ligustrum ovalifolium</i> in Turkey. <i>European Journal of Plant Pathology</i> , 2013, 137, 113-117.	0.8	3
40	Emaravirus-specific degenerate PCR primers allowed the identification of partial RNA-dependent RNA polymerase sequences of Maize red stripe virus and Pigeonpea sterility mosaic virus. <i>Journal of Virological Methods</i> , 2013, 188, 37-40.	1.0	33
41	Detection and Molecular Characterization of <i>Pepper Mild Mottle Virus</i> from Turkey. <i>Journal of Phytopathology</i> , 2013, 161, 434-438.	0.5	20
42	Complete sequence of RNA1 of grapevine Anatolian ringspot virus. <i>Archives of Virology</i> , 2012, 157, 2013-2016.	0.9	5
43	Grapevine deformation virus: Completion of the sequence and evidence on its origin from recombination events between Grapevine fanleaf virus and Arabis mosaic virus. <i>Virus Research</i> , 2012, 166, 136-140.	1.1	20
44	Occurrence of <i>Hop Stunt Viroid</i> in Mulberry (<i>Morus alba</i>) in Lebanon and Italy. <i>Journal of Phytopathology</i> , 2012, 160, 48-51.	0.5	16
45	Current Status of Fig Mosaic Disease in Iran. <i>Journal of Phytopathology</i> , 2012, 160, 324-330.	0.5	10
46	Complete sequence of Fig fleck-associated virus, a novel member of the family Tymoviridae. <i>Virus Research</i> , 2011, 161, 198-202.	1.1	33
47	The Prevalence of Three Viruses Infecting Fig in Southern Turkey. <i>Journal of Phytopathology</i> , 2011, 159, 181-183.	0.5	11
48	Detection of Olive-infecting Viruses in Tunisia. <i>Journal of Phytopathology</i> , 2011, 159, 283-286.	0.5	22
49	The complete nucleotide sequence and genome organization of Fig cryptic virus, a novel bipartite dsRNA virus infecting fig, widely distributed in the Mediterranean basin. <i>Virus Genes</i> , 2011, 42, 415-421.	0.7	40
50	Complete nucleotide sequence and genome organisation of grapevine Bulgarian latent virus. <i>Archives of Virology</i> , 2011, 156, 875-879.	0.9	11
51	Inter-laboratory validation of PCR-based protocol for detection of olive viruses. <i>EPPO Bulletin</i> , 2010, 40, 423-428.	0.6	20
52	Complete nucleotide sequence and genome organization of Olive latent virus 3, a new putative member of the family Tymoviridae. <i>Virus Research</i> , 2010, 152, 10-18.	1.1	31
53	Complete nucleotide sequence of four RNA segments of fig mosaic virus. <i>Archives of Virology</i> , 2009, 154, 1719-1727.	0.9	87
54	A multipartite single-stranded negative-sense RNA virus is the putative agent of fig mosaic disease. <i>Journal of General Virology</i> , 2009, 90, 1281-1288.	1.3	108

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55	Incidence of Viruses and Nematode Vectors in Lebanese Vineyards. <i>Journal of Phytopathology</i> , 2008, 156, 304-310.	0.5	12
56	Development of degenerate and species-specific primers for the differential and simultaneous RT-PCR detection of grapevine-infecting nepoviruses of subgroups A, B and C. <i>Journal of Virological Methods</i> , 2007, 141, 34-40.	1.0	51
57	Viruses of grapevine in Syria. <i>EPPO Bulletin</i> , 2006, 36, 523-528.	0.6	9
58	On the presence and distribution of olive viruses in Lebanon. <i>EPPO Bulletin</i> , 2005, 35, 33-36.	0.6	27
59	Preliminary evaluation of the status of olive-infecting viruses in Syria. <i>EPPO Bulletin</i> , 2005, 35, 249-252.	0.6	20