

Ruhui Li

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

56
papers

743
citations

14
h-index

25
g-index

59
ext. papers

949
ext. citations

2.5
avg, IF

3.95
L-index

#	Paper	IF	Citations
56	A reliable and inexpensive method of nucleic acid extraction for the PCR-based detection of diverse plant pathogens. <i>Journal of Virological Methods</i> , 2008 , 154, 48-55	2.6	162
55	Detection of Geminiviruses in Sweetpotato by Polymerase Chain Reaction. <i>Plant Disease</i> , 2004 , 88, 1347-1351	1.3	53
54	Triticum mosaic poacevirus enlists P1 rather than HC-Pro to suppress RNA silencing-mediated host defense. <i>Virology</i> , 2012 , 433, 104-115	3.6	50
53	Complete nucleotide sequence and taxonomy of Sugarcane streak mosaic virus, member of a novel genus in the family Potyviridae. <i>Virus Genes</i> , 2010 , 40, 432-9	2.3	39
52	Simultaneous detection and identification of four sugarcane viruses by one-step RT-PCR. <i>Journal of Virological Methods</i> , 2009 , 162, 64-8	2.6	34
51	Molecular analysis of the complete genomic sequences of four isolates of Gooseberry vein banding associated virus. <i>Virus Genes</i> , 2011 , 43, 130-7	2.3	24
50	Development of a polyprobe to detect six viroids of pome and stone fruit trees. <i>Journal of Virological Methods</i> , 2011 , 171, 91-7	2.6	23
49	An improved reverse transcription-polymerase chain reaction (RT-PCR) assay for the detection of two cherry flexiviruses in <i>Prunus</i> spp. <i>Journal of Virological Methods</i> , 2005 , 129, 162-9	2.6	22
48	Phylogenetic relationships of closely related potyviruses infecting sweet potato determined by genomic characterization of Sweet potato virus G and Sweet potato virus 2. <i>Virus Genes</i> , 2012 , 45, 118-25	2.3	21
47	Elimination of five viruses from sugarcane using in vitro culture of axillary buds and apical meristems. <i>Plant Cell, Tissue and Organ Culture</i> , 2012 , 109, 439-445	2.7	20
46	Simultaneous detection and differentiation of four closely related sweet potato potyviruses by a multiplex one-step RT-PCR. <i>Journal of Virological Methods</i> , 2012 , 186, 161-6	2.6	19
45	Molecular characterization of a novel luteovirus from peach identified by high-throughput sequencing. <i>Archives of Virology</i> , 2017 , 162, 2903-2905	2.6	15
44	Molecular characterization of a novel rhabdovirus infecting blackcurrant identified by high-throughput sequencing. <i>Archives of Virology</i> , 2018 , 163, 1363-1366	2.6	15
43	Molecular characterization of a novel luteovirus infecting apple by next-generation sequencing. <i>Archives of Virology</i> , 2018 , 163, 761-765	2.6	14
42	Simultaneous detection of four causal agents of tobacco bushy top disease by a multiplex one-step RT-PCR. <i>Journal of Virological Methods</i> , 2014 , 205, 99-103	2.6	14
41	Characterization of a flowering cherry strain of Cherry necrotic rusty mottle virus. <i>Archives of Virology</i> , 2008 , 153, 973-8	2.6	14
40	Detection of tobamoviruses by RT-PCR using a novel pair of degenerate primers. <i>Journal of Virological Methods</i> , 2018 , 259, 122-128	2.6	12

39	Molecular characterization of a novel citrivirus from citrus using next-generation sequencing. <i>Archives of Virology</i> , 2018 , 163, 3479-3482	2.6	11
38	Virome of : Discovery of and Molecular Characterization of New Viruses of Different Taxa in Camellias. <i>Frontiers in Microbiology</i> , 2020 , 11, 945	5.7	10
37	Development of a multiplex TaqMan real-time RT-PCR assay for simultaneous detection of Asian prunus viruses, plum bark necrosis stem pitting associated virus, and peach latent mosaic viroid. <i>European Journal of Plant Pathology</i> , 2013 , 137, 797-804	2.1	10
36	One-step multiplex RT-PCR for simultaneous detection of four pome tree viroids. <i>European Journal of Plant Pathology</i> , 2012 , 133, 765-772	2.1	10
35	Characterization of three new viruses of the family Betaflexiviridae associated with camellia ringspot disease. <i>Virus Research</i> , 2019 , 272, 197668	6.4	9
34	Characterization of a new apple luteovirus identified by high-throughput sequencing. <i>Virology Journal</i> , 2018 , 15, 85	6.1	9
33	Complete genome sequence of the original Taiwanese isolate of sweet potato latent virus and its relationship to other potyviruses infecting sweet potato. <i>Archives of Virology</i> , 2013 , 158, 2189-92	2.6	9
32	Genome characterization of sweet potato symptomless virus 1: a mastrevirus with an unusual nonanucleotide sequence. <i>Archives of Virology</i> , 2017 , 162, 2881-2884	2.6	8
31	Biological characterization and complete genomic sequence of Apium virus Y infecting celery. <i>Virus Research</i> , 2011 , 155, 76-82	6.4	8
30	Simultaneous detection of Cherry necrotic rusty mottle virus and Cherry green ring mottle virus using real-time PCR and high resolution melting analysis. <i>Molecular and Cellular Probes</i> , 2014 , 28, 186-91 ³⁻³	3.3	7
29	Complete genome sequence of two isolates of pokeweed mosaic virus and its relationship to other members of the genus Potyvirus. <i>Archives of Virology</i> , 2012 , 157, 2023-6	2.6	7
28	Complete genome sequence of Paris mosaic necrosis virus, a distinct member of the genus Potyvirus. <i>Archives of Virology</i> , 2018 , 163, 787-790	2.6	7
27	Characterization and detection of a new badnavirus infecting Epiphyllum spp. <i>Archives of Virology</i> , 2019 , 164, 1837-1841	2.6	6
26	Loquat Is a New Natural Host of Apple Stem Grooving Virus and Apple Chlorotic Leaf Spot Virus in China. <i>Plant Disease</i> , 2019 , 103, 3290-3290	1.5	6
25	Characterizations of Carrot thin leaf virus based on host reactions and complete genomic sequences. <i>European Journal of Plant Pathology</i> , 2014 , 138, 15-22	2.1	6
24	Complete genome sequence of a divergent strain of Japanese yam mosaic virus from China. <i>Archives of Virology</i> , 2015 , 160, 573-6	2.6	6
23	Characterization of the partial RNA1 and RNA2 3' untranslated region of Tomato ringspot virus isolates from North America. <i>Canadian Journal of Plant Pathology</i> , 2011 , 33, 94-99	1.6	6
22	Complete genomic sequence of tea-oil camellia deltapartivirus 1, a novel virus from Camellia oleifera. <i>Archives of Virology</i> , 2020 , 165, 227-231	2.6	6

21	Molecular characterization and detection of a new closterovirus identified from blackcurrant by high-throughput sequencing. <i>Virus Genes</i> , 2018 , 54, 828-832	2.3	6
20	First identification and molecular characterization of a novel cherry robigovirus. <i>Archives of Virology</i> , 2019 , 164, 3103-3106	2.6	5
19	Complete genome sequence of yam chlorotic necrosis virus, a novel macluravirus infecting yam. <i>Archives of Virology</i> , 2018 , 163, 2275-2278	2.6	5
18	Complete genome sequence of Celery mosaic virus and its relationship to other members of the genus Potyvirus. <i>Archives of Virology</i> , 2011 , 156, 917-20	2.6	5
17	Camellia ringspot-associated virus 4, a proposed new foveavirus from Camellia japonica. <i>Archives of Virology</i> , 2020 , 165, 1707-1710	2.6	4
16	Molecular characterization and detection of two carlaviruses infecting cactus. <i>Archives of Virology</i> , 2019 , 164, 1873-1876	2.6	3
15	Molecular characterization of a new badnavirus infecting green Sichuan pepper (<i>Zanthoxylum schinifolium</i>). <i>Archives of Virology</i> , 2019 , 164, 2613-2616	2.6	3
14	First Report of Beet western yellows virus Infecting Epiphyllum spp. <i>Plant Disease</i> , 2018 , 102, 464-464	1.5	3
13	Mulberry (<i>Morus alba</i>) is a new natural host of Citrus leaf blotch virus in China. <i>Plant Disease</i> , 2020 ,	1.5	3
12	First identification and molecular characterization of a new badnavirus infecting camellia. <i>Archives of Virology</i> , 2020 , 165, 2115-2118	2.6	2
11	First identification and molecular characterization of a novel cavemovirus infecting Epiphyllum spp. <i>Archives of Virology</i> , 2020 , 165, 2083-2086	2.6	2
10	Discovery and molecular characterization of a novel trichovirus infecting sweet cherry. <i>Virus Genes</i> , 2020 , 56, 380-385	2.3	2
9	Simultaneous detection and differentiation of three Potyviridae viruses in sweet potato by a multiplex TaqMan real time RT-PCR assay. <i>Journal of Virological Methods</i> , 2018 , 252, 24-31	2.6	2
8	Molecular Detection and Characterization of Chinese Yam Mild Mosaic Virus Isolates. <i>Journal of Phytopathology</i> , 2015 , 163, 1036-1040	1.8	1
7	<i>Senna bicapsularis</i> : A New Natural Host of Bean Yellow Mosaic Virus in China. <i>Plant Disease</i> , 2019 , 103, 2144-2144	1.5	1
6	First Report of Alternanthera Mosaic Virus Infecting Epiphyllum spp.. <i>Plant Disease</i> , 2019 , 103, 780-780	1.5	1
5	Identification and molecular characterization of tea-oil camellia-associated totivirus 1. <i>Archives of Virology</i> , 2021 , 166, 2347-2351	2.6	1
4	First Report of Blackcurrant Reversion Virus in <i>Ribes nigrum</i> Germplasm in the United States. <i>Plant Disease</i> , 2019 , 103, 1051	1.5	1

3	A Multiyear Survey and Identification of Pepper- and Tomato-Infecting Viruses in Yunnan Province, China. <i>Frontiers in Microbiology</i> , 2021 , 12, 623875	5.7	1
2	First Report of Citrus Vein Enation Virus from Citrus Cultivar Huangguogan in Sichuan Province, China. <i>Plant Disease</i> , 2019 , 103, 2701-2701	1.5	0
1	Genomic characterization of a new enamovirus infecting common bean.. <i>Archives of Virology</i> , 2022 , 167, 999	2.6	