

# Ruhui Li

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6831794/publications.pdf>

Version: 2024-02-01

58  
papers

1,077  
citations

471061

17  
h-index

454577

30  
g-index

59  
all docs

59  
docs citations

59  
times ranked

1013  
citing authors

#	ARTICLE	IF	CITATIONS
1	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.	1.0	215
2	Detection of Geminiviruses in Sweetpotato by Polymerase Chain Reaction. <i>Plant Disease</i> , 2004, 88, 1347-1351.	0.7	70
3	Triticum mosaic poacevirus enlists P1 rather than HC-Pro to suppress RNA silencing-mediated host defense. <i>Virology</i> , 2012, 433, 104-115.	1.1	68
4	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, 432-439.	0.7	45
5	Simultaneous detection and identification of four sugarcane viruses by one-step RT-PCR. <i>Journal of Virological Methods</i> , 2009, 162, 64-68.	1.0	41
6	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, 161-166.	1.0	30
7	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, 118-125.	0.7	30
8	Development of a polyprobe to detect six viroids of pome and stone fruit trees. <i>Journal of Virological Methods</i> , 2011, 171, 91-97.	1.0	29
9	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, 162-169.	1.0	28
10	Molecular analysis of the complete genomic sequences of four isolates of Gooseberry vein banding associated virus. <i>Virus Genes</i> , 2011, 43, 130-137.	0.7	28
11	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.	1.2	28
12	Virome of <i>Camellia japonica</i> : Discovery of and Molecular Characterization of New Viruses of Different Taxa in Camellias. <i>Frontiers in Microbiology</i> , 2020, 11, 945.	1.5	28
13	Detection of tobamoviruses by RT-PCR using a novel pair of degenerate primers. <i>Journal of Virological Methods</i> , 2018, 259, 122-128.	1.0	26
14	Molecular characterization of a novel luteovirus from peach identified by high-throughput sequencing. <i>Archives of Virology</i> , 2017, 162, 2903-2905.	0.9	23
15	Characterization of a new apple luteovirus identified by high-throughput sequencing. <i>Virology Journal</i> , 2018, 15, 85.	1.4	23
16	Molecular characterization of a novel rhabdovirus infecting blackcurrant identified by high-throughput sequencing. <i>Archives of Virology</i> , 2018, 163, 1363-1366.	0.9	22
17	Genome characterization of sweet potato symptomless virus 1: a mastrevirus with an unusual nonanucleotide sequence. <i>Archives of Virology</i> , 2017, 162, 2881-2884.	0.9	20
18	Molecular characterization of a novel luteovirus infecting apple by next-generation sequencing. <i>Archives of Virology</i> , 2018, 163, 761-765.	0.9	17

#	ARTICLE	IF	CITATIONS
19	Characterization of a flowering cherry strain of Cherry necrotic rusty mottle virus. Archives of Virology, 2008, 153, 973-978.	0.9	16
20	Loquat Is a New Natural Host of Apple Stem Grooving Virus and Apple Chlorotic Leaf Spot Virus in China. Plant Disease, 2019, 103, 3290-3290.	0.7	16
21	Simultaneous detection of four causal agents of tobacco bushy top disease by a multiplex one-step RT-PCR. Journal of Virological Methods, 2014, 205, 99-103.	1.0	15
22	Molecular characterization of a novel citrivirus from citrus using next-generation sequencing. Archives of Virology, 2018, 163, 3479-3482.	0.9	15
23	Characterization of three new viruses of the family Betaflexiviridae associated with camellia ringspot disease. Virus Research, 2019, 272, 197668.	1.1	15
24	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. European Journal of Plant Pathology, 2013, 137, 797-804.	0.8	14
25	Complete genome sequence of Paris mosaic necrosis virus, a distinct member of the genus Potyvirus. Archives of Virology, 2018, 163, 787-790.	0.9	13
26	Complete genome sequence of a divergent strain of Japanese yam mosaic virus from China. Archives of Virology, 2015, 160, 573-576.	0.9	12
27	Characterization of the partial RNA1 and RNA2 3' untranslated region of Tomato ringspot virus isolates from North America. Canadian Journal of Plant Pathology, 2011, 33, 94-99.	0.8	11
28	One-step multiplex RT-PCR for simultaneous detection of four pome tree viroids. European Journal of Plant Pathology, 2012, 133, 765-772.	0.8	11
29	Complete genome sequence of the original Taiwanese isolate of sweet potato latent virus and its relationship to other potyviruses infecting sweet potato. Archives of Virology, 2013, 158, 2189-2192.	0.9	11
30	Biological characterization and complete genomic sequence of Apium virus Y infecting celery. Virus Research, 2011, 155, 76-82.	1.1	10
31	Complete genome sequence of yam chlorotic necrosis virus, a novel macluravirus infecting yam. Archives of Virology, 2018, 163, 2275-2278.	0.9	10
32	Complete genomic sequence of tea-oil camellia deltapartivirus 1, a novel virus from Camellia oleifera. Archives of Virology, 2020, 165, 227-231.	0.9	10
33	Complete genome sequence of Celery mosaic virus and its relationship to other members of the genus Potyvirus. Archives of Virology, 2011, 156, 917-920.	0.9	9
34	Identification and molecular characterization of tea-oil camellia-associated totivirus 1. Archives of Virology, 2021, 166, 2347-2351.	0.9	9
35	First Report of Beet western yellows virus Infecting Epiphyllum spp. Plant Disease, 2018, 102, 464-464.	0.7	9
36	Complete genome sequence of two isolates of pokeweed mosaic virus and its relationship to other members of the genus Potyvirus. Archives of Virology, 2012, 157, 2023-2026.	0.9	8

#	ARTICLE	IF	CITATIONS
37	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, 186-191.	0.9	8
38	Molecular characterization and detection of a new closterovirus identified from blackcurrant by high-throughput sequencing. <i>Virus Genes</i> , 2018, 54, 828-832.	0.7	8
39	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.	0.8	7
40	Molecular Detection and Characterization of Chinese Yam Mild Mosaic Virus Isolates. <i>Journal of Phytopathology</i> , 2015, 163, 1036-1040.	0.5	7
41	Camellia ringspot-associated virus 4, a proposed new foveavirus from <i>Camellia japonica</i> . <i>Archives of Virology</i> , 2020, 165, 1707-1710.	0.9	7
42	First identification and molecular characterization of a novel cherry robigovirus. <i>Archives of Virology</i> , 2019, 164, 3103-3106.	0.9	6
43	Characterization and detection of a new badnavirus infecting <i>Epiphyllum</i> spp.. <i>Archives of Virology</i> , 2019, 164, 1837-1841.	0.9	6
44	Mulberry ( <i>Morus alba</i> ) Is a New Natural Host of Citrus Leaf Blotch Virus in China. <i>Plant Disease</i> , 2021, 105, 716-716.	0.7	6
45	Molecular characterization of a new badnavirus infecting green Sichuan pepper ( <i>Zanthoxylum</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 1	0.9	4
46	Discovery and molecular characterization of a novel trichovirus infecting sweet cherry. <i>Virus Genes</i> , 2020, 56, 380-385.	0.7	4
47	A Multiyear Survey and Identification of Pepper- and Tomato-Infecting Viruses in Yunnan Province, China. <i>Frontiers in Microbiology</i> , 2021, 12, 623875.	1.5	4
48	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.	1.0	3
49	Molecular characterization and detection of two carlaviruses infecting cactus. <i>Archives of Virology</i> , 2019, 164, 1873-1876.	0.9	3
50	First identification and molecular characterization of a new badnavirus infecting camellia. <i>Archives of Virology</i> , 2020, 165, 2115-2118.	0.9	3
51	First identification and molecular characterization of a novel cavemovirus infecting <i>Epiphyllum</i> spp.. <i>Archives of Virology</i> , 2020, 165, 2083-2086.	0.9	3
52	The Occurrence of Pea Enation Mosaic Virus 1 and Pea Enation Mosaic Virus 2 from Disease-Affected Pea Fields in China. <i>Plant Disease</i> , 2021, 105, 518.	0.7	3
53	<i>Senna bicapsularis</i> : A New Natural Host of Bean Yellow Mosaic Virus in China. <i>Plant Disease</i> , 2019, 103, 2144-2144.	0.7	3
54	First Report of Citrus Vein Enation Virus from Citrus Cultivar Huangguogan in Sichuan Province, China. <i>Plant Disease</i> , 2019, 103, 2701-2701.	0.7	2

#	ARTICLE	IF	CITATIONS
55	First Report of Blackcurrant Reversion Virus in <i>Ribes nigrum</i> Germplasm in the United States. <i>Plant Disease</i> , 2019, 103, 1051.	0.7	2
56	First Report of Alternanthera Mosaic Virus Infecting <i>Epiphyllum</i> spp.. <i>Plant Disease</i> , 2019, 103, 780-780.	0.7	2
57	Genomic characterization of a new enamovirus infecting common bean. <i>Archives of Virology</i> , 2022, 167, 999-1002.	0.9	1
58	Structure and Genome Organization of a Novel Fiji Strain of Sweet Potato Vein Clearing Virus Identified by High-Throughput Sequencing. <i>Genome Announcements</i> , 2018, 6, .	0.8	0