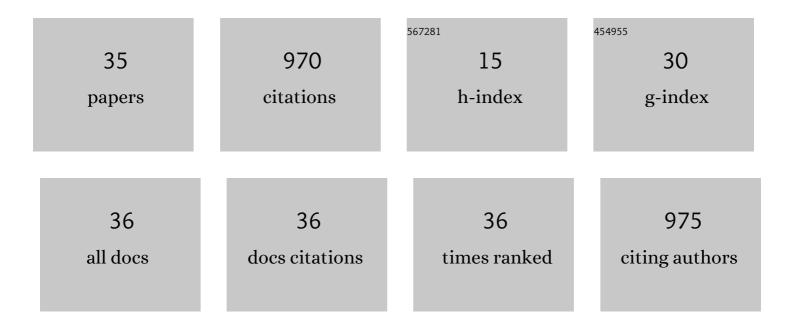
Guoping Wang

List of Publications by Year in descending order

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18

#	Article	IF	CITATIONS
1	Novel insight into the distribution and dissemination of <i>Candidatus</i> Liberibacter asiaticus, the causal agent of citrus Huanglongbing. Plant Biotechnology Journal, 2022, 20, 247-249.	8.3	6
2	Transcriptome Analysis of the Molecular Patterns of Pear Plants Infected by Two Colletotrichum fructicola Pathogenic Strains Causing Contrasting Sets of Leaf Symptoms. Frontiers in Plant Science, 2022, 13, 761133.	3.6	7
3	Molecular Characteristics and Incidence of Apple Rubbery Wood Virus 2 and Citrus Virus A Infecting Pear Trees in China. Viruses, 2022, 14, 576.	3.3	4
4	Seed Transmission of Three Viruses in Two Pear Rootstock Species Pyrus betulifolia and P. calleryana. Viruses, 2022, 14, 599.	3.3	5
5	A Novel Heptasegmented Positive-Sense Single-Stranded RNA Virus from the Phytopathogenic Fungus <i>Colletotrichum fructicola</i> . Journal of Virology, 2022, 96, e0031822.	3.4	10
6	Timeâ€resolved fluorescent microsphere lateral flow biosensors for rapid detection of <i>Candidatus</i> Liberibacter asiaticus. Plant Biotechnology Journal, 2022, 20, 1235-1237.	8.3	4
7	A mycovirus modulates the endophytic and pathogenic traits of a plant associated fungus. ISME Journal, 2021, 15, 1893-1906.	9.8	49
8	The p23 of Citrus Tristeza Virus Interacts with Host FKBP-Type Peptidyl-Prolylcis-Trans Isomerase 17-2 and Is Involved in the Intracellular Movement of the Viral Coat Protein. Cells, 2021, 10, 934.	4.1	6
9	Characterization of Diaporthe species associated with peach constriction canker, with two novel species from China. MycoKeys, 2021, 80, 77-90.	1.9	11
10	A novel <i>Actinidia</i> cytorhabdovirus characterized using genomic and viral protein interaction features. Molecular Plant Pathology, 2021, 22, 1271-1287.	4.2	8
11	2021 Taxonomic update of phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. Archives of Virology, 2021, 166, 3513-3566.	2.1	62
12	Molecular Characteristics of Jujube Yellow Mottle-Associated Virus Infecting Jujube (Ziziphus jujuba) Tj ETQq0 0	Ο rgBT /Ον	erlgck 10 Tf
13	Molecular characterization of a novel emaravrius infecting Actinidia spp. in China. Virus Research, 2020, 275, 197736.	2.2	34
14	Next-Generation Sequencing Combined With Conventional Sanger Sequencing Reveals High Molecular Diversity in Actinidia Virus 1 Populations From Kiwifruit Grown in China. Frontiers in Microbiology, 2020, 11, 602039.	3.5	6
15	Identification and characterization of water chestnut Soymovirus-1 (WCSV-1), a novel Soymovirus in water chestnuts (Eleocharis dulcis). BMC Plant Biology, 2019, 19, 159.	3.6	2

16	Functional analysis of apple stem pitting virus coat protein variants. Virology Journal, 2019, 16, 20.	3.4	12
17	Characterization of a novel victorivirus isolated from the phytopathogenic fungus Botryosphaeria dothidea. Archives of Virology, 2019, 164, 1609-1617.	2.1	17

18The Coat Protein of Citrus Yellow Vein Clearing Virus Interacts with Viral Movement Proteins and
Serves as an RNA Silencing Suppressor. Viruses, 2019, 11, 329.3.3

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#	Article	IF	CITATIONS
19	A sensitive nested multiplex RT-PCR assay for the simultaneous detection of three common viruses infecting pear plants. Journal of Virological Methods, 2019, 263, 105-110.	2.1	4
20	The detection of ACLSV and ASPV in pear plants by RT-LAMP assays. Journal of Virological Methods, 2018, 252, 80-85.	2.1	31
21	Molecular characterization of an <i>Apple stem grooving virus</i> isolate from kiwifruit (<i>Actinidia chinensis</i>) in China. Canadian Journal of Plant Pathology, 2018, 40, 76-83.	1.4	13
22	Identification and Characterization of a Novel Hepta-Segmented dsRNA Virus From the Phytopathogenic Fungus Colletotrichum fructicola. Frontiers in Microbiology, 2018, 9, 754.	3.5	35
23	A rapid silica spin column-based method of RNA extraction from fruit trees for RT-PCR detection of viruses. Journal of Virological Methods, 2017, 247, 61-67.	2.1	32
24	Characterization of a novel botybirnavirus isolated from a phytopathogenic Alternaria fungus. Archives of Virology, 2017, 162, 3907-3911.	2.1	33
25	A dsRNA virus with filamentous viral particles. Nature Communications, 2017, 8, 168.	12.8	84
26	Characterization of a novel double-stranded RNA mycovirus conferring hypovirulence from the phytopathogenic fungus Botryosphaeria dothidea. Virology, 2016, 493, 75-85.	2.4	83
27	Genetic diversity and evolution of <i>Apple stem pitting virus</i> isolates from pear in China. Canadian Journal of Plant Pathology, 2016, 38, 218-230.	1.4	16
28	Characterization of Colletotrichum fructicola, a new causal agent of leaf black spot disease of sandy pear (Pyrus pyrifolia). European Journal of Plant Pathology, 2015, 143, 651-662.	1.7	32
29	Biological and Molecular Characterization of Five <i>Phomopsis</i> Species Associated with Pear Shoot Canker in China. Plant Disease, 2015, 99, 1704-1712.	1.4	40
30	Different roles for RNA silencing and RNA processing components in virus recovery and virus-induced gene silencing in plants. Journal of Experimental Botany, 2015, 66, 919-932.	4.8	125
31	The genome sequences of three isolates of <i>Apple chlorotic leaf spot virus</i> from pear (<i>Pyrus</i> sp.) in China. Canadian Journal of Plant Pathology, 2014, 36, 396-402.	1.4	6
32	Hypovirulence of the Phytopathogenic Fungus Botryosphaeria dothidea: Association with a Coinfecting Chrysovirus and a Partitivirus. Journal of Virology, 2014, 88, 7517-7527.	3.4	115
33	Simultaneous detection and differentiation of three viruses in pear plants by a multiplex RT-PCR. Journal of Virological Methods, 2014, 196, 113-119.	2.1	33
34	Genetic variability and population structure ofGrapevine virus Ain China based on the analysis of its coat protein gene. Canadian Journal of Plant Pathology, 2011, 33, 227-233.	1.4	9
35	Surface Remeshing on Triangular Domain for CAD Applications. , 2007, , .		0