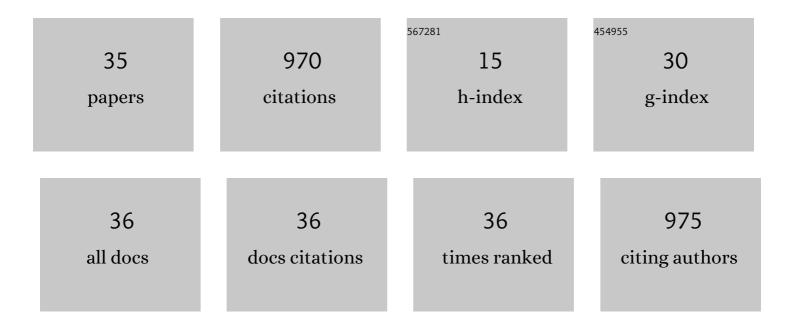
Guoping Wang

List of Publications by Year in descending order

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CHODING WANG

#	Article	IF	CITATIONS
1	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
2	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
3	A dsRNA virus with filamentous viral particles. Nature Communications, 2017, 8, 168.	12.8	84
4	Characterization of a novel double-stranded RNA mycovirus conferring hypovirulence from the phytopathogenic fungus Botryosphaeria dothidea. Virology, 2016, 493, 75-85.	2.4	83
5	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
6	A mycovirus modulates the endophytic and pathogenic traits of a plant associated fungus. ISME Journal, 2021, 15, 1893-1906.	9.8	49
7	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
8	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
9	Molecular characterization of a novel emaravrius infecting Actinidia spp. in China. Virus Research, 2020, 275, 197736.	2.2	34
10	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
11	Characterization of a novel botybirnavirus isolated from a phytopathogenic Alternaria fungus. Archives of Virology, 2017, 162, 3907-3911.	2.1	33
12	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
13	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
14	The detection of ACLSV and ASPV in pear plants by RT-LAMP assays. Journal of Virological Methods, 2018, 252, 80-85.	2.1	31
15	The 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	18
16	Characterization of a novel victorivirus isolated from the phytopathogenic fungus Botryosphaeria dothidea. Archives of Virology, 2019, 164, 1609-1617.	2.1	17
17	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
18	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

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19	Functional analysis of apple stem pitting virus coat protein variants. Virology Journal, 2019, 16, 20.	3.4	12
20	Characterization of Diaporthe species associated with peach constriction canker, with two novel species from China. MycoKeys, 2021, 80, 77-90.	1.9	11
21	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
22	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
23	Molecular Characteristics of Jujube Yellow Mottle-Associated Virus Infecting Jujube (Ziziphus jujuba) Tj ETQq1 1).784314 3.3	rg&T /Overlo
24	A novel <i>Actinidia</i> cytorhabdovirus characterized using genomic and viral protein interaction features. Molecular Plant Pathology, 2021, 22, 1271-1287.	4.2	8
25	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
26	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
27	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
28	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
29	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
30	Seed Transmission of Three Viruses in Two Pear Rootstock Species Pyrus betulifolia and P. calleryana. Viruses, 2022, 14, 599.	3.3	5
31	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
32	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
33	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
34	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
35	Surface Remeshing on Triangular Domain for CAD Applications. , 2007, , .		0