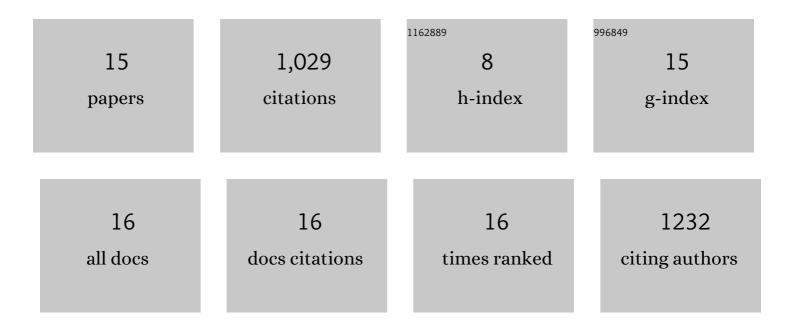
## Jiawei Wang

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5075767/publications.pdf Version: 2024-02-01



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#	Article	IF	CITATIONS
1	MFCIS: an automatic leaf-based identification pipeline for plant cultivars using deep learning and persistent homology. Horticulture Research, 2021, 8, 172.	2.9	7
2	Chromosome-scale genome assembly of sweet cherry (Prunus avium L.) cv. Tieton obtained using long-read and Hi-C sequencing. Horticulture Research, 2020, 7, 122.	2.9	44
3	A de novo assembly of the sweet cherry ( <i>Prunus avium</i> cv. Tieton) genome using linked-read sequencing technology. Peerl, 2020, 8, e9114.	0.9	8
4	Transcriptomic analysis reveals insights into the response to <i>Hop stunt viroid</i> (HSVd) in sweet cherry ( <i>Prunus avium</i> L.) fruits. PeerJ, 2020, 8, e10005.	0.9	7
5	Complete Genome Sequence of a Little Cherry Virus-2 Isolate from Sweet Cherry in China. Microbiology Resource Announcements, 2019, 8, .	0.3	1
6	Multilocus genotyping identifies a highly homogeneous phytoplasma lineage associated with sweet cherry virescence disease in China and its carriage by an erythroneurine leafhopper. Crop Protection, 2018, 106, 13-22.	1.0	13
7	â€~ Candidatus Phytoplasma brasiliense'-related strains associated with papaya bunchy top disease in northern Peru represent a distinct geographic lineage. Crop Protection, 2017, 92, 99-106.	1.0	7
8	Novel phytoplasma strains of Xâ€disease group unveil genetic markers that distinguish North American and South American geographic lineages within subgroups 16SrIllâ€J and 16SrIllâ€U. Annals of Applied Biology, 2017, 171, 405-416.	1.3	8
9	Complete nucleotide sequence of little cherry virus 1 (LChV-1) infecting sweet cherry in China. Archives of Virology, 2016, 161, 749-753.	0.9	11
10	Phytoplasmas change the source–sink relationship of field-grown sweet cherry by disturbing leaf function. Physiological and Molecular Plant Pathology, 2015, 92, 22-27.	1.3	17
11	Genomewide identification and expression analysis of the ARF gene family in apple. Journal of Genetics, 2014, 93, 785-797.	0.4	40
12	A multiplex RT-PCR assay for simultaneous detection of four viruses from sweet cherry. Scientia Horticulturae, 2014, 180, 118-122.	1.7	8
13	Rapid detection of Prunus necrotic ringspot virus using magnetic nanoparticle-assisted reverse transcription loop-mediated isothermal amplification. Journal of Virological Methods, 2014, 208, 85-89.	1.0	19
14	The genome of the pear ( <i>Pyrus bretschneideri</i> Rehd.). Genome Research, 2013, 23, 396-408.	2.4	832
15	Improvement for Agronomic Traits of Partial Waxy Wheat by Combination of Backcrossing with a PCR-based DNA Marker. Journal of Genetics and Genomics, 2007, 34, 836-841.	1.7	7