

Monica A Kehoe

List of Publications by Year in descending order

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Version: 2024-02-01

30
papers

647
citations

686830

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642321

23
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33
all docs

33
docs citations

33
times ranked

549
citing authors

#	ARTICLE	IF	CITATIONS
1	Plant Virology and Next Generation Sequencing: Experiences with a Potyvirus. PLoS ONE, 2014, 9, e104580.	1.1	72
2	Analyses of Twelve New Whole Genome Sequences of Cassava Brown Streak Viruses and Ugandan Cassava Brown Streak Viruses from East Africa: Diversity, Supercomputing and Evidence for Further Speciation. PLoS ONE, 2015, 10, e0139321.	1.1	70
3	Cassava brown streak virus has a rapidly evolving genome: implications for virus speciation, variability, diagnosis and host resistance. Scientific Reports, 2016, 6, 36164.	1.6	66
4	Split Personality of a Potyvirus: To Specialize or Not to Specialize?. PLoS ONE, 2014, 9, e105770.	1.1	50
5	Indigenous and introduced potyviruses of legumes and Passiflora spp. from Australia: biological properties and comparison of coat protein nucleotide sequences. Archives of Virology, 2011, 156, 1757-1774.	0.9	38
6	The Biology and Phylogenetics of <i>Potato virus S</i> Isolates from the Andean Region of South America. Plant Disease, 2018, 102, 869-885.	0.7	35
7	Real time portable genome sequencing for global food security. F1000Research, 0, 7, 1101.	0.8	32
8	Unusual occurrence of a DAG motif in the Ipomovirus Cassava brown streak virus and implications for its vector transmission. PLoS ONE, 2017, 12, e0187883.	1.1	29
9	A proposal to rationalize within-species plant virus nomenclature: benefits and implications of inaction. Archives of Virology, 2016, 161, 2051-2057.	0.9	26
10	<i>Zucchini yellow mosaic virus</i> Populations from East Timorese and Northern Australian Cucurbit Crops: Molecular Properties, Genetic Connectivity, and Biosecurity Implications. Plant Disease, 2017, 101, 1236-1245.	0.7	24
11	Genetic diversity and recombination between turnip yellows virus strains in Australia. Archives of Virology, 2021, 166, 813-829.	0.9	23
12	A proposal to help resolve the disagreement between naming of potato virus Y strain groups defined by resistance phenotypes and those defined by sequencing. Archives of Virology, 2011, 156, 2273-2278.	0.9	18
13	Resistance Phenotypes in Diverse Accessions, Breeding Lines, and Cultivars of Three Mustard Species Inoculated with <i>Turnip mosaic virus</i> . Plant Disease, 2010, 94, 1290-1298.	0.7	16
14	Two Complete Genome Sequences of Phasey Bean Mild Yellows Virus, a Novel Member of the <i>Luteoviridae</i> from Australia. Genome Announcements, 2016, 4, .	0.8	14
15	Application of Loop-Mediated Isothermal Amplification in an Early Warning System for Epidemics of an Externally Sourced Plant Virus. Plants, 2019, 8, 139.	1.6	14
16	First report of Cassava brown streak viruses on wild plant species in Mozambique. Physiological and Molecular Plant Pathology, 2019, 105, 88-95.	1.3	14
17	<i>Pratylenchus quasitereoides</i> n. sp. from cereals in Western Australia. Zootaxa, 2014, 3866, 277-88.	0.2	12
18	Evolutionary insights of <i>Bean common mosaic necrosis virus</i> and <i>Cowpea aphid-borne mosaic virus</i> . PeerJ, 2019, 7, e6297.	0.9	12

#	ARTICLE	IF	CITATIONS
19	First Report of Grapevine Rupestris Vein Feathering Virus in Grapevine in Australia. <i>Plant Disease</i> , 2021, 105, 515.	0.7	10
20	First Complete Genome Sequence of <i>Cucumber green mottle mosaic virus</i> Isolated from Australia. <i>Genome Announcements</i> , 2017, 5, .	0.8	9
21	Biology and genetic diversity of phasey bean mild yellows virus, a common virus in legumes in Australia. <i>Archives of Virology</i> , 2021, 166, 1575-1589.	0.9	9
22	Epidemiology of Zucchini yellow mosaic virus in cucurbit crops in a remote tropical environment. <i>Virus Research</i> , 2020, 281, 197897.	1.1	7
23	Phylogenomic relationship and evolutionary insights of sweet potato viruses from the western highlands of Kenya. <i>PeerJ</i> , 2018, 6, e5254.	0.9	7
24	A metagenomic study of DNA viruses from samples of local varieties of common bean in Kenya. <i>PeerJ</i> , 2019, 7, e6465.	0.9	7
25	Occurrence of cucumber green mottle mosaic virus in Western Australia. <i>Australasian Plant Pathology</i> , 2022, 51, 1-8.	0.5	7
26	Turnip yellows virus and Soybean dwarf virus in Western Australia. <i>Australasian Plant Pathology</i> , 2019, 48, 323-329.	0.5	6
27	Host plant affiliations of aphid vector species found in a remote tropical environment. <i>Virus Research</i> , 2020, 281, 197934.	1.1	4
28	Genomic characterisation and evolutionary relationships of groundnut rosette virus from the western highlands of Kenya. <i>Tropical Plant Pathology</i> , 2018, 43, 583-585.	0.8	3
29	Genetic diversity and SNPs from the chloroplast coding regions of virus-infected cassava. <i>PeerJ</i> , 2020, 8, e8632.	0.9	1
30	Using Genomics to Design a Pathovar-Specific Loop-Mediated Isothermal Amplification (LAMP) Assay, for the Improved Detection of <i>Xanthomonas citri</i> pv. <i>citri</i> . <i>Microorganisms</i> , 2022, 10, 1153.	1.6	0