Roger A C Jones

List of Publications by Year
in descending order

226 papers	8,171 citations	53794 45 h-index	82547 72 g-index
229 all docs	229 docs citations	229 times ranked	3162 citing authors

#	Article	IF	Citations
1	Phylogenetics and Evolution of Potato Virus V: Another Potyvirus that Originated in the Andes. Plant Disease, 2022, 106, 691-700.	1.4	8
2	Occurrence of cucumber green mottle mosaic virus in Western Australia. Australasian Plant Pathology, 2022, 51, 1-8.	1.0	7
3	Alteration of plant species mixtures by virus infection: Managed pastures the forgotten dimension. Plant Pathology, 2022, 71, 1255-1281.	2.4	5
4	Phylogenetics and evolution of wheat streak mosaic virus: ItsÂglobal origin and the source of the Australian epidemic. Plant Pathology, 2022, 71, 1660-1673.	2.4	5
5	Historical virus isolate collections: An invaluable resource connecting plant virologyâ€™s preâ€sequencing and postâ€sequencing eras. Plant Pathology, 2021, 70, 235-248.	2.4	19
6	Global Plant Virus Disease Pandemics and Epidemics. Plants, 2021, 10, 233.	3.5	158
7	The Phylogeography of Potato Virus X Shows the Fingerprints of Its Human Vector. Viruses, 2021, 13, 644.	3.3	12
8	Potato Virus Y Biological Strain Group Y^D: Hypersensitive Resistance Genes Elicited and Phylogenetic Placement. Plant Disease, 2021, 105, 3600-3609.	1.4	4
9	Potato Virus A Isolates from Three Continents: Their Biological Properties, Phylogenetics, and Prehistory. Phytopathology, 2021, 111, 217-226.	2.2	24
10	Virus Diseases of Cereal and Oilseed Crops in Australia: Current Position and Future Challenges. Viruses, 2021, 13, 2051.	3.3	19
11	Disease Pandemics and Major Epidemics Arising from New Encounters between Indigenous Viruses and Introduced Crops. Viruses, 2020, 12, 1388.	3.3	50
12	Host plant affiliations of aphid vector species found in a remote tropical environment. Virus Research, 2020, 281, 197934.	2.2	4
13	Genetic Diversity of Nine Non-Recombinant Potato virus Y Isolates From Three Biological Strain Groups: Historical and Geographical Insights. Plant Disease, 2020, 104, 2317-2323.	1.4	17
14	Epidemiology of Zucchini yellow mosaic virus in cucurbit crops in a remote tropical environment. Virus Research, 2020, 281, 197897.	2.2	7
15	The Potyviruses: An Evolutionary Synthesis Is Emerging. Viruses, 2020, 12, 132.	3.3	74
16	Global Dimensions of Plant Virus Diseases: Current Status and Future Perspectives. Annual Review of Virology, 2019, 6, 387-409.	6.7	173
17	Effects of a Potato Spindle Tuber Viroid Tomato Strain on the Symptoms, Biomass, and Yields of Classical Indicator and Currently Grown Potato and Tomato Cultivars. Plant Disease, 2019, 103, 3009-3017.	1.4	11
18	Genomic sequence and host range studies reveal considerable variation within the species Arracacha virus B. Archives of Virology, 2019, 164, 2849-2852.	2.1	2

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19	Potato virus Y; the Andean connection. Virus Evolution, 2019, 5, vez037.	4.9	34
20	Biological and Molecular Properties of <i>Wild potato mosaic virus</i> Isolates from Pepino (<i>Solanum muricatum</i>). Plant Disease, 2019, 103, 1746-1756.	1.4	12
21	<i>Zucchini yellow mosaic virus</i> Genomic Sequences from Papua New Guinea: Lack of Genetic Connectivity with Northern Australian or East Timorese Genomes, and New Recombination Findings. Plant Disease, 2019, 103, 1326-1336.	1.4	5
22	Genetic Connectivity Between Papaya Ringspot Virus Genomes from Papua New Guinea and Northern Australia, and New Recombination Insights. Plant Disease, 2019, 103, 737-747.	1.4	11
23	Complete Coding Sequence of <i>Andean Potato Mottle Virus</i> from a 40-Year-Old Sample from Peru. Microbiology Resource Announcements, 2019, 8, .	0.6	1
24	New Isolates of <i>Sweet potato feathery mottle virus</i> and <i>Sweet potato virus C</i>: Biological and Molecular Properties, and Recombination Analysis Based on Complete Genomes. Plant Disease, 2018, 102, 1899-1914.	1.4	11
25	First Complete Genome Sequence of Cucurbit aphid-borne yellows virus from Papua New Guinea. Genome Announcements, 2018, 6, .	0.8	7
26	<i>Sweet potato feathery mottle virus</i> and <i>Sweet potato virus C</i> from East Timorese and Australian Sweetpotato: Biological and Molecular Properties, and Biosecurity Implications. Plant Disease, 2018, 102, 589-599.	1.4	17
27	Strain-Specific Hypersensitive and Extreme Resistance Phenotypes Elicited by <i>Potato virus Y</i> Among 39 Potato Cultivars Released in Three World Regions Over a 117-Year Period. Plant Disease, 2018, 102, 185-196.	1.4	32
28	The Biology and Phylogenetics of <i>Potato virus S</i> Isolates from the Andean Region of South America. Plant Disease, 2018, 102, 869-885.	1.4	35
29	Complete Genomic Sequence of the Potyvirus <i>Mashua Virus Y</i> , Obtained from a 33-Year-Old Mashua (<i>Tropaeaolum tuberosum</i>) Sample. Microbiology Resource Announcements, 2018, 7, .	0.6	5
30	A 33-Year-Old Plant Sample Contributes the First Complete Genomic Sequence of <i>Potato Virus U</i>. Microbiology Resource Announcements, 2018, 7, .	0.6	4
31	Full-Genome Sequencing of a Virus from a 33-Year-Old Sample Demonstrates that <i>Arracacha Mottle Virus</i> Is Synonymous with <i>Arracacha Virus Y</i>. Microbiology Resource Announcements, 2018, 7, .	0.6	4
32	Complete Genome Sequence of Potato Virus T from Bolivia, Obtained from a 33-Year-Old Sample. Microbiology Resource Announcements, 2018, 7, .	0.6	4
33	Plant and Insect Viruses in Managed and Natural Environments: Novel and Neglected Transmission Pathways. Advances in Virus Research, 2018, 101, 149-187.	2.1	45
34	<i>Papaya ringspot virus</i> Populations From East Timorese and Northern Australian Cucurbit Crops: Biological and Molecular Properties, and Absence of Genetic Connectivity. Plant Disease, 2017, 101, 985-993.	1.4	30
35	Biological and Molecular Properties of a<i>Turnip mosaic virus</i>(TuMV) Strain that Breaks TuMV Resistances in<i>Brassica napus</i>. Plant Disease, 2017, 101, 674-683.	1.4	29
36	<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.	1.4	24

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37	Establishing alighting preferences and species transmission differences for Pea seed-borne mosaic virus aphid vectors. Virus Research, 2017, 241, 145-155.	2.2	9
38	Analysis of an RNA-seq Strand-Specific Library from an East Timorese Cucumber Sample Reveals a Complete <i>Cucurbit aphid-borne yellows virus</i> Genome. Genome Announcements, 2017, 5, .	0.8	12
39	A Risk Assessment Framework for Seed Degeneration: Informing an Integrated Seed Health Strategy for Vegetatively Propagated Crops. Phytopathology, 2017, 107, 1123-1135.	2.2	53
40	<i>Cucumber green mottle mosaic virus</i>: Rapidly Increasing Global Distribution, Etiology, Epidemiology, and Management. Annual Review of Phytopathology, 2017, 55, 231-256.	7.8	140
41	Forecasting model for Pea seed-borne mosaic virus epidemics in field pea crops in a Mediterranean-type environment. Virus Research, 2017, 241, 163-171.	2.2	5
42	Seed fractionation as a phytosanitary control measure for Pea seed-borne mosaic virus infection of field pea seed-stocks. European Journal of Plant Pathology, 2017, 148, 733-737.	1.7	4
43	First Complete Genome Sequence of <i>Cucumber green mottle mosaic virus</i> Isolated from Australia. Genome Announcements, 2017, 5, .	0.8	9
44	<i>Pea seed-borne mosaic virus</i>Pathosystem Drivers under Mediterranean-Type Climatic Conditions: Deductions from 23 Epidemic Scenarios. Plant Disease, 2017, 101, 929-940.	1.4	3
45	First Complete Genome Sequence of <i>Arracacha virus A</i> Isolated from a 38-Year-Old Sample from Peru. Genome Announcements, 2017, 5, .	0.8	8
46	Two Complete Genome Sequences of Squash mosaic virus from 20-Year-Old Cucurbit Leaf Samples from Australia. Genome Announcements, 2017, 5, .	0.8	8
47	First Complete Squash leaf curl China virus Genomic Segment DNA-A Sequence from East Timor. Genome Announcements, 2017, 5, .	0.8	15
48	The phylogenetics of the global population of potato virus Y and its necrogenic recombinants. Virus Evolution, 2017, 3, vex002.	4.9	57
49	Metagenomic Analysis of Cucumber RNA from East Timor Reveals an <i>Aphid lethal paralysis virus</i> Genome. Genome Announcements, 2017, 5, .	0.8	23
50	Analysis of an RNA-seq Strand-Specific Library Sample Reveals a Complete Genome of <i>Hardenbergia mosaic virus</i> from Native Wisteria, an Indigenous Virus from Southwest Australia. Genome Announcements, 2017, 5, .	0.8	4
51	First Complete Genome Sequence of Suakwa aphid-borne yellows virus from East Timor. Genome Announcements, 2016, 4, .	0.8	13
52	Programmed cell death pathways induced by early plantâ€virus infection are determined by isolate virulence and stage of infection. Plant Pathology, 2016, 65, 1518-1528.	2.4	12
53	Effects of Temperature on Disease Severity in Plants of Subterranean Clover Infected Singly or in Mixed Infection with <i>Bean yellow mosaic virus</i> and <i>Kabatiella caulivora</i>. Journal of Phytopathology, 2016, 164, 608-619.	1.0	14
54	Complete Genome Sequences of the Potyvirus Sweet potato virus 2 from East Timor and Australia. Genome Announcements, 2016, 4, .	0.8	14

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55	<i>Pea seed-borne mosaic virus</i>in Field Pea: Widespread Infection, Genetic Diversity, and Resistance Gene Effectiveness. Plant Disease, 2016, 100, 2475-2482.	1.4	15
56	First Complete Genome Sequence of <i>Bean common mosaic necrosis virus</i> from East Timor. Genome Announcements, 2016, 4, .	0.8	13
57	Strain specificity of Turnip mosaic virus resistance gene TuRBJU 01 in Brassica juncea. European Journal of Plant Pathology, 2016, 145, 209-213.	1.7	16
58	A proposal to rationalize within-species plant virus nomenclature: benefits and implications of inaction. Archives of Virology, 2016, 161, 2051-2057.	2.1	26
59	Future Scenarios for Plant Virus Pathogens as Climate Change Progresses. Advances in Virus Research, 2016, 95, 87-147.	2.1	107
60	First Complete Genome Sequence of Pepper vein yellows virus from Australia. Genome Announcements, 2016, 4, .	0.8	18
61	Systemic Hypersensitive Resistance to <i>Turnip mosaic virus</i> in <i>Brassica juncea</i> is Associated With Multiple Defense Responses, Especially Phloem Necrosis and Xylem Occlusion. Plant Disease, 2016, 100, 1261-1270.	1.4	17
62	<i>Pea seed-borne mosaic virus</i>: Stability and Wind-Mediated Contact Transmission in Field Pea. Plant Disease, 2016, 100, 953-958.	1.4	17
63	Deep Sequencing Reveals the Complete Genome Sequence of Sweet potato virus G from East Timor. Genome Announcements, 2016, 4, .	0.8	15
64	Improving <i>Potato virus Y</i> strain nomenclature: lessons from comparing isolates obtained over a 73â€year period. Plant Pathology, 2016, 65, 322-333.	2.4	57
65	Complete Genome Sequences of the Carlavirus Sweet potato chlorotic fleck virus from East Timor and Australia. Genome Announcements, 2016, 4, .	0.8	16
66	Potato spindle tuber viroid: alternative host reservoirs and strain found in a remote subtropical irrigation area. European Journal of Plant Pathology, 2016, 145, 433-446.	1.7	27
67	Spread of introduced viruses to new plants in natural ecosystems and the threat this poses to plant biodiversity. Molecular Plant Pathology, 2015, 16, 541-545.	4.2	29
68	Biological and molecular variation amongst Australian <i>Turnip mosaic virus</i> isolates. Plant Pathology, 2015, 64, 1215-1223.	2.4	19
69	Studies on resistance phenotypes to Turnip mosaic virus in five species of Brassicaceae, and identification of a virus resistance gene in Brassica juncea. European Journal of Plant Pathology, 2015, 141, 647-666.	1.7	22
70	<i>Potato spindle tuber viroid</i>: Stability on Common Surfaces and Inactivation With Disinfectants. Plant Disease, 2015, 99, 770-775.	1.4	14
71	<i>Potato virus Y</i>: Contact Transmission, Stability, Inactivation, and Infection Sources. Plant Disease, 2015, 99, 387-394.	1.4	26
72	Effects of Introduced and Indigenous Viruses on Native Plants: Exploring Their Disease Causing Potential at the Agro-Ecological Interface. PLoS ONE, 2014, 9, e91224.	2.5	45

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73	<i>Polymyxa graminis</i> Isolates from Australia: Identification in Wheat Roots and Soil, Molecular Characterization, and Wide Genetic Diversity. Plant Disease, 2014, 98, 1567-1575.	1.4	13
74	Black Pod Syndrome of Lupinus angustifolius Is Caused by Late Infection with Bean yellow mosaic virus. Plant Disease, 2014, 98, 739-745.	1.4	8
75	First Report of Wheat mosaic virus Infecting Wheat in Western Australia. Plant Disease, 2014, 98, 285-285.	1.4	27
76	Epidemiology of Wheat streak mosaic virus in wheat in a Mediterranean-type environment. European Journal of Plant Pathology, 2014, 140, 797-813.	1.7	16
77	Control of Plant Virus Diseases in Cool-Season Grain Legume Crops. Advances in Virus Research, 2014, 90, 207-253.	2.1	31
78	Hardenbergia mosaic virus: Crossing the barrier between native and introduced plant species. Virus Research, 2014, 184, 87-92.	2.2	16
79	Trends in plant virus epidemiology: Opportunities from new or improved technologies. Virus Research, 2014, 186, 3-19.	2.2	50
80	Plant virus ecology and epidemiology: historical perspectives, recent progress and future prospects. Annals of Applied Biology, 2014, 164, 320-347.	2.5	67
81	Preliminary studies on resistance phenotypes to Turnip mosaic virus in Brassica napus and B. carinata from different continents and effects of temperature on their expression. European Journal of Plant Pathology, 2014, 139, 687-706.	1.7	19
82	Genetic improvement of subterranean clover (Trifolium subterraneum L.). 2. Breeding for disease and pest resistance. Crop and Pasture Science, 2014, 65, 1207.	1.5	33
83	Plant Virology and Next Generation Sequencing: Experiences with a Potyvirus. PLoS ONE, 2014, 9, e104580.	2.5	72
84	Split Personality of a Potyvirus: To Specialize or Not to Specialize?. PLoS ONE, 2014, 9, e105770.	2.5	50
85	<i>Zucchini yellow mosaic virus</i>: Contact Transmission, Stability on Surfaces, and Inactivation with Disinfectants. Plant Disease, 2013, 97, 765-771.	1.4	30
86	Virus diseases of perennial pasture legumes in Australia: incidences, losses, epidemiology, and management. Crop and Pasture Science, 2013, 64, 199.	1.5	17
87	Virus diseases of pasture grasses in Australia: incidences, losses, epidemiology, and management. Crop and Pasture Science, 2013, 64, 216.	1.5	18
88	Biological Properties of <i>Potato virus X</i> in Potato: Effects of Mixed Infection with <i>Potato virus S</i> and Resistance Phenotypes in Cultivars from Three Continents. Plant Disease, 2012, 96, 43-54.	1.4	29
89	Virus diseases of annual pasture legumes: incidences, losses, epidemiology, and management. Crop and Pasture Science, 2012, 63, 399.	1.5	33
90	Detection of Viruses in Sweetpotato from Honduras and Guatemala Augmented by Deep-Sequencing of Small-RNAs. Plant Disease, 2012, 96, 1430-1437.	1.4	68

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91	Effects of tissue sampling position, primary and secondary infection, cultivar, and storage temperature and duration on the detection, concentration and distribution of three viruses within infected potato tubers. Australasian Plant Pathology, 2012, 41, 197-210.	1.0	4
92	First Report of <i>Alfalfa mosaic virus</i> Infecting Tedera (<i>Bituminaria bituminosa</i> (L.) C.H.) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 1384-1384.	1.4	8
93	Minimising losses caused by Zucchini yellow mosaic virus in vegetable cucurbit crops in tropical, sub-tropical and Mediterranean environments through cultural methods and host resistance. Virus Research, 2011, 159, 141-160.	2.2	41
94	A virus of an isolated indigenous flora spreads naturally to an introduced crop species. Annals of Applied Biology, 2011, 159, 339-347.	2.5	12
95	Genetic variability of the coat protein sequence of pea seed-borne mosaic virus isolates and the current relationship between phylogenetic placement and resistance groups. Archives of Virology, 2011, 156, 1287-1290.	2.1	13
96	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.	2.1	38
97	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.	2.1	18
98	Zucchini yellow mosaic virus: biological properties, detection procedures and comparison of coat protein gene sequences. Archives of Virology, 2011, 156, 2119-2131.	2.1	51
99	Advances in winter pulse pathology research in Australia. Australasian Plant Pathology, 2011, 40, 549-567.	1.0	23
100	Virus symptomatology in accessions of the Medicago truncatula core collection and identification of virus resistance phenotypes. Crop and Pasture Science, 2011, 62, 686.	1.5	3
101	Molecular Genetic Characterization of Olpidium virulentus Isolates Associated with Big-Vein Diseased Lettuce Plants. Plant Disease, 2010, 94, 563-569.	1.4	29
102	Control of Beet western yellows virus in Brassica napus crops: infection resistance in Australian genotypes and effectiveness of imidacloprid seed dressing. Crop and Pasture Science, 2010, 61, 321.	1.5	17
103	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.	1.4	16
104	Genome sequences and phylogenetic placement of two isolates of<i>Bean common mosaic virus</i>from<i>Macroptilium atropurpureum</i>in north-west Australia. Australasian Plant Pathology, 2010, 39, 184.	1.0	14
105	Comparison of the coat protein genes of Lettuce big-vein associated virus isolates from Australia with those of isolates from other continents. Archives of Virology, 2010, 155, 765-770.	2.1	10
106	Genetic variability in the coat protein gene of Potato virus S isolates and distinguishing its biologically distinct strains. Archives of Virology, 2010, 155, 1163-1169.	2.1	33
107	Genetic variability in the coat protein gene of Potato virus X and the current relationship between phylogenetic placement and resistance groupings. Archives of Virology, 2010, 155, 1349-1356.	2.1	27
108	Comparison of the coat protein genes of Mirafiori lettuce big-vein virus isolates from Australia with those of isolates from other continents. Archives of Virology, 2010, 155, 1519-1522.	2.1	10

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109	An epidemiological model for externally acquired vector-borne viruses applied to Beet western yellows virus in Brassica napus crops in a Mediterranean-type environment. Crop and Pasture Science, 2010, 61, 132.	1.5	14
110	Principles of Predicting Plant Virus Disease Epidemics. Annual Review of Phytopathology, 2010, 48, 179-203.	7.8	61
111	Quantifying Effects of Seedborne Inoculum on Virus Spread, Yield Losses, and Seed Infection in the <i>Pea seed-borne mosaic virus</i>â€“Field Pea Pathosystem. Phytopathology, 2009, 99, 1156-1167.	2.2	52
112	A single tube, quantitative real-time RT-PCR assay that detects four potato viruses simultaneously. Journal of Virological Methods, 2009, 161, 289-296.	2.1	55
113	BYDV PREDICTOR: a simulation model to predict aphid arrival, epidemics of <i>Barley yellow dwarf virus</i> and yield losses in wheat crops in a Mediterraneanâ€type environment. Plant Pathology, 2009, 58, 186-202.	2.4	71
114	Plant virus emergence and evolution: Origins, new encounter scenarios, factors driving emergence, effects of changing world conditions, and prospects for control. Virus Research, 2009, 141, 113-130.	2.2	320
115	Global status of tospovirus epidemics in diverse cropping systems: Successes achieved and challenges ahead. Virus Research, 2009, 141, 219-236.	2.2	491
116	Resistance to Subterranean clover mottle virus in Medicago truncatula and genetic mapping of a resistance locus. Crop and Pasture Science, 2009, 60, 480.	1.5	6
117	Molecular Characterization of <i>Sweet potato feathery mottle virus</i> (SPFMV) Isolates from Easter Island, French Polynesia, New Zealand, and Southern Africa. Plant Disease, 2009, 93, 933-939.	1.4	26
118	Role of Recombination in the Evolution of Host Specialization Within <i>Bean yellow mosaic virus</i>. Phytopathology, 2009, 99, 512-518.	2.2	43
119	Discussion paper: The naming of Potato virus Y strains infecting potato. Archives of Virology, 2008, 153, 1-13.	2.1	262
120	Natural resistance to<i>Alfalfa mosaic virus</i>in different lupin species. Australasian Plant Pathology, 2008, 37, 112.	1.0	10
121	<i>Cucumber mosaic virus</i> infection of chickpea stands: temporal and spatial patterns of spread and yieldâ€limiting potential. Plant Pathology, 2008, 57, 842-853.	2.4	23
122	Molecular Genetic Characterization of <i>Sweet potato virus G</i> (SPVG) Isolates from Areas of the Pacific Ocean and Southern Africa. Plant Disease, 2008, 92, 1313-1320.	1.4	24
123	Phylogenetic Analysis of Bean yellow mosaic virus Isolates from Four Continents: Relationship Between the Seven Groups Found and Their Hosts and Origins. Plant Disease, 2008, 92, 1596-1603.	1.4	43
124	An Epidemiological Model for Externally Sourced Vector-Borne Viruses Applied to <i>Bean yellow mosaic virus</i> in Lupin Crops in a Mediterranean-Type Environment. Phytopathology, 2008, 98, 1280-1290.	2.2	19
125	Further studies on Pea seed-borne mosaic virus in cool-season crop legumes: responses to infection and seed quality defects. Australian Journal of Agricultural Research, 2008, 59, 1130.	1.5	33
126	Finding Wheat streak mosaic virus in south-west Australia. Australian Journal of Agricultural Research, 2008, 59, 836.	1.5	20

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127	The epidemiology of Wheat streak mosaic virus in Australia: case histories, gradients, mite vectors, and alternative hosts. Australian Journal of Agricultural Research, 2008, 59, 844.	1.5	41
128	Evaluation of resistance to Turnip mosaic virus in Australian Brassica napus genotypes. Australian Journal of Agricultural Research, 2007, 58, 67.	1.5	15
129	Yield-limiting potential of Beet western yellows virus in Brassica napus. Australian Journal of Agricultural Research, 2007, 58, 788.	1.5	44
130	Wheat streak mosaic virus in Australia: Relationship to Isolates from the Pacific Northwest of the USA and Its Dispersion Via Seed Transmission. Plant Disease, 2007, 91, 164-170.	1.4	60
131	Lack of Seed Coat Contamination with Cucumber mosaic virus in Lupin Permits Reliable, Large-Scale Detection of Seed Transmission in Seed Samples. Plant Disease, 2007, 91, 504-508.	1.4	18
132	Virus impact at the interface of an ancient ecosystem and a recent agroecosystem: studies on three legumeâ€infecting potyviruses in the southwest Australian floristic region. Plant Pathology, 2007, 56, 729-742.	2.4	77
133	Detection of Sweet potato chlorotic fleck virus and<i>Sweet potato feathery mottle virus</i>â€“ strain O in Australia. Australasian Plant Pathology, 2007, 36, 591.	1.0	20
134	Control of Plant Virus Diseases. Advances in Virus Research, 2006, 67, 205-244.	2.1	89
135	Wild Plants and Viruses: Underâ€Investigated Ecosystems. Advances in Virus Research, 2006, 67, 1-47.	2.1	94
136	Occurrence of Beet western yellows virus and its aphid vectors in over-summering broad-leafed weeds and volunteer crop plants in the grainbelt region of south-western Australia. Australian Journal of Agricultural Research, 2006, 57, 975.	1.5	43
137	Localized Distribution of Iris yellow spot virus Within Leeks and Its Reliable Large-Scale Detection. Plant Disease, 2006, 90, 729-733.	1.4	24
138	Potyvirus Complexes in Sweetpotato: Occurrence in Australia, Serological and Molecular Resolution, and Analysis of the Sweet potato virus 2 (SPV2) Component. Plant Disease, 2006, 90, 1120-1128.	1.4	43
139	Phytoplasma from little leaf disease affected sweetpotato in Western Australia: detection and phylogeny. Annals of Applied Biology, 2006, 149, 9-14.	2.5	11
140	Relative abilities of different aphid species to act as vectors ofCarrot virus Y. Australasian Plant Pathology, 2006, 35, 23.	1.0	3
141	'CandidatusPhytoplasma australiense' is associated with diseases of red clover and paddy melon in south-west Australia. Australasian Plant Pathology, 2006, 35, 283.	1.0	6
142	Genetic diversity of the nucleocapsid gene ofIris yellow spot virus. Australasian Plant Pathology, 2006, 35, 359.	1.0	11
143	Incidence and distribution of Barley yellow dwarf virus and Cereal yellow dwarf virus in over-summering grasses in a Mediterranean-type environment. Australian Journal of Agricultural Research, 2005, 56, 257.	1.5	52
144	Incidence and distribution of viruses infecting cucurbit crops in the Northern Territory and Western Australia. Australian Journal of Agricultural Research, 2005, 56, 847.	1.5	35

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145	Unravelling the genetic diversity of the three main viruses involved in Sweet Potato Virus Disease (SPVD), and its practical implications. Molecular Plant Pathology, 2005, 6, 199-211.	4.2	107
146	First report of Bean common mosaic virus in Western Australia. Plant Pathology, 2005, 54, 563-563.	2.4	11
147	Patterns of spread of Carrot virus Y in carrot plantings and validation of control measures. Annals of Applied Biology, 2005, 147, 57-67.	2.5	10
148	Patterns of spread of two non-persistently aphid-borne viruses in lupin stands under four different infection scenarios. Annals of Applied Biology, 2005, 146, 337-350.	2.5	48
149	Inheritance of hypersensitive resistance to Bean yellow mosaic virus in narrow-leafed lupin (Lupinus) Tj ETQq1 1 0.784314 rgBT /Overlo	2.5	21
150	Genetic variability of Tomato spotted wilt virus in Australia and validation of real time RT-PCR for its detection in single and bulked leaf samples. Annals of Applied Biology, 2005, 146, 517-530.	2.5	21
151	Suppressing spread of Tomato spotted wilt virus by drenching infected source or healthy recipient plants with neonicotinoid insecticides to control thrips vectors. Annals of Applied Biology, 2005, 146, 95-103.	2.5	30
152	Capsicum chlorosis virus infectingCapsicum annuumin the East Kimberley region of Western Australia. Australasian Plant Pathology, 2005, 34, 397.	1.0	12
153	Role of winter-active aphids spreading Barley yellow dwarf virus in decreasing wheat yields in a Mediterranean-type environment. Australian Journal of Agricultural Research, 2005, 56, 1089.	1.5	27
154	Seed Transmission of Wheat streak mosaic virus Shown Unequivocally in Wheat. Plant Disease, 2005, 89, 1048-1050.	1.4	63
155	Further studies on Carrot virus Y: hosts, symptomatology, search for resistance, and tests for seed transmissibility. Australian Journal of Agricultural Research, 2005, 56, 859.	1.5	13
156	Deploying partially resistant genotypes and plastic mulch on the soil surface to suppress spread of lettuce big-vein disease in lettuce. Australian Journal of Agricultural Research, 2004, 55, 131.	1.5	13
157	Yield losses caused by virus infection in four combinations of non-persistently aphid-transmitted virus and cool-season crop legume. Australian Journal of Experimental Agriculture, 2004, 44, 57.	1.0	24
158	Patterns of spread of Tomato spotted wilt virus in field crops of lettuce and pepper: spatial dynamics and validation of control measures. Annals of Applied Biology, 2004, 145, 231-245.	2.5	66
159	Analysing spatial patterns of spread of Lettuce necrotic yellows virus and lettuce big-vein disease in lettuce field plantings. Annals of Applied Biology, 2004, 145, 339-343.	2.5	15
160	Distribution and incidence of Carrot virus Y in Australia. Australasian Plant Pathology, 2004, 33, 83.	1.0	12
161	Resistance toCucumber mosaic virusinLupinus mutabilis(pearl lupin). Australasian Plant Pathology, 2004, 33, 591.	1.0	9
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