

Hideki KondÅ•

List of Publications by Year in descending order

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56
papers

3,548
citations

117625

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161849

54
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docs citations

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times ranked

3339
citing authors

#	ARTICLE	IF	CITATIONS
1	Plant viruses and viroids in Japan. <i>Journal of General Plant Pathology</i> , 2022, 88, 105-127.	1.0	16
2	Coat protein of Chinese wheat mosaic virus upregulates and interacts with cytosolic glyceraldehyde-3-phosphate dehydrogenase, a negative regulator of plant autophagy, to promote virus infection. <i>Journal of Integrative Plant Biology</i> , 2022, 64, 1631-1645.	8.5	9
3	Pathogenetic roles of beet necrotic yellow vein virus RNA5 in the exacerbation of symptoms and yield reduction, development of scab-like symptoms, and resistance breaking in sugar beet. <i>Plant Pathology</i> , 2021, 70, 219-232.	2.4	9
4	Identification of a Novel Quinvirus in the Family Betaflexiviridae That Infects Winter Wheat. <i>Frontiers in Microbiology</i> , 2021, 12, 715545.	3.5	7
5	2021 Taxonomic update of phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. <i>Archives of Virology</i> , 2021, 166, 3513-3566.	2.1	62
6	2020 taxonomic update for phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. <i>Archives of Virology</i> , 2020, 165, 3023-3072.	2.1	184
7	Diverse Partitiviruses From the Phytopathogenic Fungus, <i>Rosellinia necatrix</i> . <i>Frontiers in Microbiology</i> , 2020, 11, 1064.	3.5	22
8	Diversity and epidemiology of plant rhabdoviruses. <i>Virus Research</i> , 2020, 281, 197942.	2.2	56
9	Virome Analysis of Aphid Populations That Infest the Barley Field: The Discovery of Two Novel Groups of Nege/Kita-Like Viruses and Other Novel RNA Viruses. <i>Frontiers in Microbiology</i> , 2020, 11, 509.	3.5	46
10	Novel Victorivirus from a Pakistani Isolate of <i>Alternaria alternata</i> Lacking a Typical Translational Stop/Restart Sequence Signature. <i>Viruses</i> , 2019, 11, 577.	3.3	35
11	Taxonomy of the order Mononegavirales: second update 2018. <i>Archives of Virology</i> , 2019, 164, 1233-1244.	2.1	70
12	Dicer functions transcriptionally and posttranscriptionally in a multilayer antiviral defense. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 2274-2281.	7.1	33
13	Two novel fungal negative-strand RNA viruses related to mymonaviruses and phenuiviruses in the shiitake mushroom (<i>Lentinula edodes</i>). <i>Virology</i> , 2019, 533, 125-136.	2.4	72
14	Identification of a Novel Hypovirulence-Inducing Hypovirus From <i>Alternaria alternata</i> . <i>Frontiers in Microbiology</i> , 2019, 10, 1076.	3.5	60
15	Taxonomy of the order Mononegavirales: update 2019. <i>Archives of Virology</i> , 2019, 164, 1967-1980.	2.1	224
16	A novel insect-infecting virga/nege-like virus group and its pervasive endogenization into insect genomes. <i>Virus Research</i> , 2019, 262, 37-47.	2.2	49
17	Taxonomy of the order Mononegavirales: update 2018. <i>Archives of Virology</i> , 2018, 163, 2283-2294.	2.1	153
18	Novel, diverse RNA viruses from Mediterranean isolates of the phytopathogenic fungus, <i>Rosellinia necatrix</i> : insights into evolutionary biology of fungal viruses. <i>Environmental Microbiology</i> , 2018, 20, 1464-1483.	3.8	92

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19	Evidence for a novel negative-stranded RNA mycovirus isolated from the plant pathogenic fungus <i>Fusarium graminearum</i> . <i>Virology</i> , 2018, 518, 232-240.	2.4	41
20	Plant rhabdoviruses—their origins and vector interactions. <i>Current Opinion in Virology</i> , 2018, 33, 198-207.	5.4	70
21	Dichorhavirus in their Host Plants and Mite Vectors. <i>Advances in Virus Research</i> , 2018, 102, 119-148.	2.1	51
22	Taxonomy of the order Mononegavirales: update 2017. <i>Archives of Virology</i> , 2017, 162, 2493-2504.	2.1	173
23	A possible occurrence of genome reassortment among bipartite rhabdoviruses. <i>Virology</i> , 2017, 508, 18-25.	2.4	39
24	The family Rhabdoviridae: mono- and bipartite negative-sense RNA viruses with diverse genome organization and common evolutionary origins. <i>Virus Research</i> , 2017, 227, 158-170.	2.2	200
25	Taxonomy of the order Mononegavirales: update 2016. <i>Archives of Virology</i> , 2016, 161, 2351-2360.	2.1	407
26	Genetic Diversity of Beet Necrotic Yellow Vein Virus. , 2016, , 109-131.		5
27	Reprint of “Sequence and phylogenetic analyses of novel totivirus-like double-stranded RNAs from field-collected powdery mildew fungi”. <i>Virus Research</i> , 2016, 219, 39-50.	2.2	1
28	A novel betapartitivirus RnPV6 from <i>Rosellinia necatrix</i> tolerates host RNA silencing but is interfered by its defective RNAs. <i>Virus Research</i> , 2016, 219, 62-72.	2.2	47
29	Sequence and phylogenetic analyses of novel totivirus-like double-stranded RNAs from field-collected powdery mildew fungi. <i>Virus Research</i> , 2016, 213, 353-364.	2.2	35
30	Detection and Analysis of Non-retroviral RNA Virus-Like Elements in Plant, Fungal, and Insect Genomes. <i>Methods in Molecular Biology</i> , 2015, 1236, 73-88.	0.9	25
31	<i>Cymbidium chlorotic mosaic virus</i> , a new sobemovirus isolated from a spring orchid (<i>Cymbidium</i>) Tj ETQq1 1 0.784314 rgBT /Overlook	2.1	6
32	A novel single-stranded RNA virus isolated from a phytopathogenic filamentous fungus, <i>Rosellinia necatrix</i> , with similarity to hypo-like viruses. <i>Frontiers in Microbiology</i> , 2014, 5, 360.	3.5	75
33	Complete genome sequence of <i>Habenaria mosaic virus</i> , a new potyvirus infecting a terrestrial orchid (<i>Habenaria radiata</i>) in Japan. <i>Archives of Virology</i> , 2014, 159, 163-166.	2.1	7
34	Transcriptional mapping of the messenger and leader RNAs of orchid fleck virus, a bisegmented negative-strand RNA virus. <i>Virology</i> , 2014, 452-453, 166-174.	2.4	20
35	Dichorhavirus: a proposed new genus for <i>Brevipalpus</i> mite-transmitted, nuclear, bacilliform, bipartite, negative-strand RNA plant viruses. <i>Archives of Virology</i> , 2014, 159, 607-619.	2.1	61
36	Nyamiviridae: Proposal for a new family in the order Mononegavirales. <i>Archives of Virology</i> , 2013, 158, 2209-2226.	2.1	29

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37	Endoplasmic reticulum export and vesicle formation of the movement protein of Chinese wheat mosaic virus are regulated by two transmembrane domains and depend on the secretory pathway. <i>Virology</i> , 2013, 435, 493-503.	2.4	52
38	Characterization of burdock mottle virus, a novel member of the genus Benyvirus, and the identification of benyvirus-related sequences in the plant and insect genomes. <i>Virus Research</i> , 2013, 177, 75-86.	2.2	31
39	Identification of the amino acid residues and domains in the cysteine-rich protein of Chinese wheat mosaic virus that are important for RNA silencing suppression and subcellular localization. <i>Molecular Plant Pathology</i> , 2013, 14, 265-278.	4.2	51
40	Viruses of the White Root Rot Fungus, <i>Rosellinia necatrix</i> . <i>Advances in Virus Research</i> , 2013, 86, 177-214.	2.1	79
41	Biological and genetic diversity of plasmodiophorid-transmitted viruses and their vectors. <i>Journal of General Plant Pathology</i> , 2013, 79, 307-320.	1.0	58
42	Evidence for negative-strand RNA virus infection in fungi. <i>Virology</i> , 2013, 435, 201-209.	2.4	70
43	Orchid Fleck Virus Structural Proteins N and P Form Intranuclear Viroplasm-Like Structures in the Absence of Viral Infection. <i>Journal of Virology</i> , 2013, 87, 7423-7434.	3.4	29
44	The cysteine-rich proteins of beet necrotic yellow vein virus and tobacco rattle virus contribute to efficient suppression of silencing in roots. <i>Journal of General Virology</i> , 2012, 93, 1841-1850.	2.9	37
45	The Evolutionary History of Beet necrotic yellow vein virus Deduced from Genetic Variation, Geographical Origin and Spread, and the Breaking of Host Resistance. <i>Molecular Plant-Microbe Interactions</i> , 2011, 24, 207-218.	2.6	64
46	The enigmatic genome of <i>Chara australis</i> virus. <i>Journal of General Virology</i> , 2011, 92, 2679-2690.	2.9	30
47	Widespread Endogenization of Genome Sequences of Non-Retroviral RNA Viruses into Plant Genomes. <i>PLoS Pathogens</i> , 2011, 7, e1002146.	4.7	173
48	Identification and characterization of structural proteins of orchid fleck virus. <i>Archives of Virology</i> , 2009, 154, 37-45.	2.1	22
49	Identification of amino acids of the beet necrotic yellow vein virus p25 protein required for induction of the resistance response in leaves of <i>Beta vulgaris</i> plants. <i>Journal of General Virology</i> , 2008, 89, 1314-1323.	2.9	57
50	RNA4-encoded p31 of beet necrotic yellow vein virus is involved in efficient vector transmission, symptom severity and silencing suppression in roots. <i>Journal of General Virology</i> , 2007, 88, 1611-1619.	2.9	70
51	Lower Levels of Transgene Silencing in Roots is Associated with Reduced DNA Methylation Levels at Non-Symmetrical Sites but not at Symmetrical Sites. <i>Plant Molecular Biology</i> , 2006, 60, 423-435.	3.9	19
52	Orchid fleck virus is a rhabdovirus with an unusual bipartite genome. <i>Journal of General Virology</i> , 2006, 87, 2413-2421.	2.9	92
53	Orchid Fleck Virus: <i>Brevipalpus californicus</i> Mite Transmission, Biological Properties and Genome Structure. <i>Experimental and Applied Acarology</i> , 2003, 30, 215-223.	1.6	79
54	Comparative Cytopathology and Immunocytochemistry of Japanese, Australian and Brazilian Isolates of Orchid fleck virus. <i>Journal of General Plant Pathology</i> , 2001, 67, 231-237.	1.0	30

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55	Epidemic progress of beet necrotic yellow vein virus: evidence from an investigation in Japan spanning half a century. <i>Plant Pathology</i> , 0, , .	2.4	5
56	Distinctive in vitro ATP Hydrolysis Activity of AtVIPP1, a Chloroplastic ESCRT-III Superfamily Protein in <i>Arabidopsis</i> . <i>Frontiers in Plant Science</i> , 0, 13, .	3.6	0