Hideki KondÅ•

List of Publications by Year
in descending order

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56 papers	3,548 citations	117625 34 h-index	161849 54 g-index
57 all docs	57 docs citations	57 times ranked	3339 citing authors

#	Article	IF	Citations
1	Taxonomy of the order Mononegavirales: update 2016. Archives of Virology, 2016, 161, 2351-2360.	2.1	407
2	Taxonomy of the order Mononegavirales: update 2019. Archives of Virology, 2019, 164, 1967-1980.	2.1	224
3	The family Rhabdoviridae: mono- and bipartite negative-sense RNA viruses with diverse genome organization and common evolutionary origins. Virus Research, 2017, 227, 158-170.	2.2	200
4	2020 taxonomic update for phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. Archives of Virology, 2020, 165, 3023-3072.	2.1	184
5	Widespread Endogenization of Genome Sequences of Non-Retroviral RNA Viruses into Plant Genomes. PLoS Pathogens, 2011, 7, e1002146.	4.7	173
6	Taxonomy of the order Mononegavirales: update 2017. Archives of Virology, 2017, 162, 2493-2504.	2.1	173
7	Taxonomy of the order Mononegavirales: update 2018. Archives of Virology, 2018, 163, 2283-2294.	2.1	153
8	Orchid fleck virus is a rhabdovirus with an unusual bipartite genome. Journal of General Virology, 2006, 87, 2413-2421.	2.9	92
9	Novel, diverse RNA viruses from Mediterranean isolates of the phytopathogenic fungus, Rosellinia necatrix: insights into evolutionary biology of fungal viruses. Environmental Microbiology, 2018, 20, 1464-1483.	3.8	92
10	Orchid Fleck Virus: Brevipalpus californicus Mite Transmission, Biological Properties and Genome Structure. Experimental and Applied Acarology, 2003, 30, 215-223.	1.6	79
11	Viruses of the White Root Rot Fungus, Rosellinia necatrix. Advances in Virus Research, 2013, 86, 177-214.	2.1	79
12	A novel single-stranded RNA virus isolated from a phytopathogenic filamentous fungus, Rosellinia necatrix, with similarity to hypo-like viruses. Frontiers in Microbiology, 2014, 5, 360.	3.5	75
13	Two novel fungal negative-strand RNA viruses related to mymonaviruses and phenuiviruses in the shiitake mushroom (Lentinula edodes). Virology, 2019, 533, 125-136.	2.4	72
14	RNA4-encoded p31 of beet necrotic yellow vein virus is involved in efficient vector transmission, symptom severity and silencing suppression in roots. Journal of General Virology, 2007, 88, 1611-1619.	2.9	70
15	Evidence for negative-strand RNA virus infection in fungi. Virology, 2013, 435, 201-209.	2.4	70
16	Plant rhabdovirusesâ€"their origins and vector interactions. Current Opinion in Virology, 2018, 33, 198-207.	5.4	70
17	Taxonomy of the order Mononegavirales: second update 2018. Archives of Virology, 2019, 164, 1233-1244.	2.1	70
18	The Evolutionary History of Beet necrotic yellow vein virus Deduced from Genetic Variation, Geographical Origin and Spread, and the Breaking of Host Resistance. Molecular Plant-Microbe Interactions, 2011, 24, 207-218.	2.6	64

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19	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
20	Dichorhavirus: a proposed new genus for Brevipalpus mite-transmitted, nuclear, bacilliform, bipartite, negative-strand RNA plant viruses. Archives of Virology, 2014, 159, 607-619.	2.1	61
21	Identification of a Novel Hypovirulence-Inducing Hypovirus From Alternaria alternata. Frontiers in Microbiology, 2019, 10, 1076.	3.5	60
22	Biological and genetic diversity of plasmodiophorid-transmitted viruses and their vectors. Journal of General Plant Pathology, 2013, 79, 307-320.	1.0	58
23	Identification of amino acids of the beet necrotic yellow vein virus p25 protein required for induction of the resistance response in leaves of Beta vulgaris plants. Journal of General Virology, 2008, 89, 1314-1323.	2.9	57
24	Diversity and epidemiology of plant rhabdoviruses. Virus Research, 2020, 281, 197942.	2.2	56
25	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. Virology, 2013, 435, 493-503.	2.4	52
26	Identification of the amino acid residues and domains in the cysteineâ€rich protein of <scp>C</scp>hinese wheat mosaic virus that are important for <scp>RNA</scp> silencing suppression and subcellular localization. Molecular Plant Pathology, 2013, 14, 265-278.	4.2	51
27	Dichorhaviruses in their Host Plants and Mite Vectors. Advances in Virus Research, 2018, 102, 119-148.	2.1	51
28	A novel insect-infecting virga/nege-like virus group and its pervasive endogenization into insect genomes. Virus Research, 2019, 262, 37-47.	2.2	49
29	A novel betapartitivirus RnPV6 from Rosellinia necatrix tolerates host RNA silencing but is interfered by its defective RNAs. Virus Research, 2016, 219, 62-72.	2.2	47
30	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. Frontiers in Microbiology, 2020, 11, 509.	3.5	46
31	Evidence for a novel negative-stranded RNA mycovirus isolated from the plant pathogenic fungus Fusarium graminearum. Virology, 2018, 518, 232-240.	2.4	41
32	A possible occurrence of genome reassortment among bipartite rhabdoviruses. Virology, 2017, 508, 18-25.	2.4	39
33	The cysteine-rich proteins of beet necrotic yellow vein virus and tobacco rattle virus contribute to efficient suppression of silencing in roots. Journal of General Virology, 2012, 93, 1841-1850.	2.9	37
34	Sequence and phylogenetic analyses of novel totivirus-like double-stranded RNAs from field-collected powdery mildew fungi. Virus Research, 2016, 213, 353-364.	2.2	35
35	Novel Victorivirus from a Pakistani Isolate of Alternaria alternata Lacking a Typical Translational Stop/Restart Sequence Signature. Viruses, 2019, 11, 577.	3.3	35
36	Dicer functions transcriptionally and posttranscriptionally in a multilayer antiviral defense. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 2274-2281.	7.1	33

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37	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. Virus Research, 2013, 177, 75-86.	2.2	31
38	Comparative Cytopathology and Immunocytochemistry of Japanese, Australian and Brazilian Isolates of Orchid fleck virus. Journal of General Plant Pathology, 2001, 67, 231-237.	1.0	30
39	The enigmatic genome of Chara australis virus. Journal of General Virology, 2011, 92, 2679-2690.	2.9	30
40	Nyamiviridae: Proposal for a new family in the order Mononegavirales. Archives of Virology, 2013, 158, 2209-2226.	2.1	29
41	Orchid Fleck Virus Structural Proteins N and P Form Intranuclear Viroplasm-Like Structures in the Absence of Viral Infection. Journal of Virology, 2013, 87, 7423-7434.	3.4	29
42	Detection and Analysis of Non-retroviral RNA Virus-Like Elements in Plant, Fungal, and Insect Genomes. Methods in Molecular Biology, 2015, 1236, 73-88.	0.9	25
43	Identification and characterization of structural proteins of orchid fleck virus. Archives of Virology, 2009, 154, 37-45.	2.1	22
44	Diverse Partitiviruses From the Phytopathogenic Fungus, Rosellinia necatrix. Frontiers in Microbiology, 2020, 11, 1064.	3.5	22
45	Transcriptional mapping of the messenger and leader RNAs of orchid fleck virus, a bisegmented negative-strand RNA virus. Virology, 2014, 452-453, 166-174.	2.4	20
46	Lower Levels of Transgene Silencing in Roots is Associated with Reduced DNA Methylation Levels at Non-Symmetrical Sites but not at Symmetrical Sites. Plant Molecular Biology, 2006, 60, 423-435.	3.9	19
47	Plant viruses and viroids in Japan. Journal of General Plant Pathology, 2022, 88, 105-127.	1.0	16
48	Pathogenetic roles of beet necrotic yellow vein virus RNA5 in the exacerbation of symptoms and yield reduction, development of scabâ€like symptoms, and Rz1â€resistance breaking in sugar beet. Plant Pathology, 2021, 70, 219-232.	2.4	9
49	Coat protein of Chinese wheat mosaic virus upregulates and interacts with cytosolic glyceraldehydeâ€3â€phosphate dehydrogenase, a negative regulator of plant autophagy, to promote virus infection. Journal of Integrative Plant Biology, 2022, 64, 1631-1645.	8.5	9
50	Complete genome sequence of Habenaria mosaic virus, a new potyvirus infecting a terrestrial orchid (Habenaria radiata) in Japan. Archives of Virology, 2014, 159, 163-166.	2.1	7
51	Identification of a Novel Quinvirus in the Family Betaflexiviridae That Infects Winter Wheat. Frontiers in Microbiology, 2021, 12, 715545.	3.5	7
52	Cymbidium chlorotic mosaic virus, a new sobemovirus isolated from a spring orchid (Cymbidium) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 1	2.1	6
53	Genetic Diversity of Beet Necrotic Yellow Vein Virus. , 2016, , 109-131.		5
54	Epidemic progress of beet necrotic yellow vein virus: evidence from an investigation in Japan spanning half a century. Plant Pathology, 0, , .	2.4	5

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55	Reprint of â€œSequence and phylogenetic analyses of novel totivirus-like double-stranded RNAs from field-collected powdery mildew fungiâ€• Virus Research, 2016, 219, 39-50.	2.2	1
56	Distinctive in vitro ATP Hydrolysis Activity of AtVIPP1, a Chloroplastic ESCRT-III Superfamily Protein in Arabidopsis. Frontiers in Plant Science, 0, 13, .	3.6	0