

# Tetsuo Tamada

## List of Publications by Year in descending order

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33  
papers

1,655  
citations

331670

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docs citations

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times ranked

710  
citing authors

#	ARTICLE	IF	CITATIONS
1	Production and Pathogenicity of Isolates of Beet Necrotic Yellow Vein Virus with Different Numbers of RNA Components. <i>Journal of General Virology</i> , 1989, 70, 3399-3409.	2.9	116
2	Association of beet necrotic yellow vein virus with isolates of <i>Polymyxa betae</i> Keskin.. <i>Nihon Shokubutsu Byori Gakkaiho = Annals of the Phytopathological Society of Japan</i> , 1986, 52, 235-247.	0.1	104
3	Evidence that Beet Necrotic Yellow Vein Virus RNA-4 Is Essential for Efficient Transmission by the Fungus <i>Polymyxa betae</i> . <i>Journal of General Virology</i> , 1989, 70, 3391-3398.	2.9	103
4	Evidence That RNA Silencing-Mediated Resistance to Beet necrotic yellow vein virus Is Less Effective in Roots Than in Leaves. <i>Molecular Plant-Microbe Interactions</i> , 2005, 18, 194-204.	2.6	98
5	Beet Necrotic Yellow Vein Virus from Rizomania-Affected Sugar Beet in Japan. <i>Nihon Shokubutsu Byori Gakkaiho = Annals of the Phytopathological Society of Japan</i> , 1973, 39, 325-332_1.	0.1	96
6	Mapping Functions on the Multipartite Genome of Beet Necrotic Yellow Vein Virus. <i>Annual Review of Phytopathology</i> , 1992, 30, 291-313.	7.8	96
7	High resolution analysis of the readthrough domain of beet necrotic yellow vein virus readthrough protein: a KTER motif is important for efficient transmission of the virus by <i>Polymyxa betae</i> . <i>Journal of General Virology</i> , 1996, 77, 1359-1367.	2.9	95
8	Orchid fleck virus is a rhabdovirus with an unusual bipartite genome. <i>Journal of General Virology</i> , 2006, 87, 2413-2421.	2.9	92
9	RNA 3 Deletion Mutants of Beet Necrotic Yellow Vein Virus Do Not Cause Rhizomania Disease in Sugar Beets. <i>Phytopathology</i> , 1999, 89, 1000-1006.	2.2	83
10	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
11	The Evolutionary History of <i>Beet necrotic yellow vein virus</i> 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
12	Evidence for three groups of sequence variants of beet necrotic yellow vein virus RNA 5. <i>Archives of Virology</i> , 1999, 144, 879-892.	2.1	59
13	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
14	Vascular movement of beet necrotic yellow vein virus in <i>Beta macrocarpa</i> is probably dependent on an RNA 3 sequence domain rather than a gene product.. <i>Journal of General Virology</i> , 1998, 79, 385-393.	2.9	58
15	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
16	Differential contributions of plant Dicer-like proteins to antiviral defences against potato virus X in leaves and roots. <i>Plant Journal</i> , 2015, 81, 781-793.	5.7	51
17	Complete nucleotide sequence of the Japanese isolate S of beet necrotic yellow vein virus RNA and comparison with European isolates. <i>Archives of Virology</i> , 1996, 141, 2163-2175.	2.1	39
18	Nucleotide sequence analysis of RNA-5 of five isolates of beet necrotic yellow vein virus and the identity of a deletion mutant. <i>Journal of General Virology</i> , 1996, 77, 575-580.	2.9	38

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19	BENYVIRUSES. , 1999, , 154-160.		37
20	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
21	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
22	The enigmatic genome of <i>Chara australis</i> virus. <i>Journal of General Virology</i> , 2011, 92, 2679-2690.	2.9	30
23	Identification and characterization of structural proteins of orchid fleck virus. <i>Archives of Virology</i> , 2009, 154, 37-45.	2.1	22
24	Host range and molecular analysis of <i>Beet leaf yellowing virus</i> and <i>Beet western yellows virus</i> and <i>Brassica yellows virus</i> in Japan. <i>Plant Pathology</i> , 2019, 68, 1045-1058.	2.4	22
25	Identification of a New Wheat Yellow Mosaic Virus Strain with Specific Pathogenicity towards Major Wheat Cultivars Grown in Hokkaido.. <i>Nihon Shokubutsu Byori Gakkaiho = Annals of the Phytopathological Society of Japan</i> , 1997, 63, 107-109.	0.1	19
26	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
27	Susceptibility and resistance of <i>Beta vulgaris</i> subsp. <i>maritima</i> to foliar rub-inoculation with Beet necrotic yellow vein virus. <i>Journal of General Plant Pathology</i> , 2007, 73, 76-80.	1.0	14
28	General Features of Beet Necrotic Yellow Vein Virus. , 2016, , 55-83.		12
29	Pathogenetic roles of beet necrotic yellow vein virus RNA5 in the exacerbation of symptoms and yield reduction, development of scab-like symptoms, and <i>Rz1</i> resistance breaking in sugar beet. <i>Plant Pathology</i> , 2021, 70, 219-232.	2.4	9
30	The Plasmodiophorid Protist <i>Polymyxa betae</i> . , 2016, , 135-153.		9
31	Identification of a Novel Quinvirus in the Family Betaflexiviridae That Infects Winter Wheat. <i>Frontiers in Microbiology</i> , 2021, 12, 715545.	3.5	7
32	Genetic Diversity of Beet Necrotic Yellow Vein Virus. , 2016, , 109-131.		5
33	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