

Da-Wei Li

List of Publications by Year in descending order

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133
papers

5,259
citations

126708

33
h-index

102304

66
g-index

139
all docs

139
docs citations

139
times ranked

4857
citing authors

#	ARTICLE	IF	CITATIONS
1	Plant virus infection disrupts vacuolar acidification and autophagic degradation for the effective infection. <i>Autophagy</i> , 2022, 18, 705-706.	4.3	5
2	SAMDC3 enhances resistance to <i>Barley stripe mosaic virus</i> by promoting the ubiquitination and proteasomal degradation of viral β protein. <i>New Phytologist</i> , 2022, 234, 618-633.	3.5	10
3	Coat proteins of necroviruses target 14-3-3a to subvert MAPKKK1-mediated antiviral immunity in plants. <i>Nature Communications</i> , 2022, 13, 716.	5.8	27
4	The Carboxyl Terminal Regions of P0 Protein Are Required for Systemic Infections of Poleroviruses. <i>International Journal of Molecular Sciences</i> , 2022, 23, 1945.	1.8	1
5	Incidence and prevalence levels of three aphid-transmitted viruses in crucifer crops in China. <i>Journal of Integrative Agriculture</i> , 2022, 21, 774-780.	1.7	10
6	A Powerful Method for Studying Protein-Protein Interactions in Plants: Coimmunoprecipitation (Co-IP) Assay. <i>Methods in Molecular Biology</i> , 2022, 2400, 87-92.	0.4	5
7	Host casein kinase 1-mediated phosphorylation modulates phase separation of a rhabdovirus phosphoprotein and virus infection. <i>ELife</i> , 2022, 11, .	2.8	21
8	Barley stripe mosaic virus β protein targets thioredoxin h-type 1 to dampen salicylic acid-mediated defenses. <i>Plant Physiology</i> , 2022, 189, 1715-1727.	2.3	7
9	A viral protein disrupts vacuolar acidification to facilitate virus infection in plants. <i>EMBO Journal</i> , 2022, 41, e108713.	3.5	15
10	Palmitoylation of β protein directs a dynamic switch between <i>Barley stripe mosaic virus</i> replication and movement. <i>EMBO Journal</i> , 2022, 41, .	3.5	3
11	Tobacco Necrosis Virus-A ^C Single Coat Protein Amino Acid Substitutions Determine Host-Specific Systemic Infections of <i>Nicotiana benthamiana</i> and Soybean. <i>Molecular Plant-Microbe Interactions</i> , 2021, 34, 49-61.	1.4	11
12	Proximity labeling: an emerging tool for probing in planta molecular interactions. <i>Plant Communications</i> , 2021, 2, 100137.	3.6	36
13	Three-dimensional reconstruction and comparison of vacuolar membranes in response to viral infection. <i>Journal of Integrative Plant Biology</i> , 2021, 63, 353-364.	4.1	14
14	Hordeiviruses (Virgaviridae). , 2021, , 420-429.		9
15	The serine/threonine/tyrosine kinase STY46 defends against hordeivirus infection by phosphorylating β protein. <i>Plant Physiology</i> , 2021, 186, 715-730.	2.3	19
16	Genome-Wide and Comprehensive Analysis of the Multiple Stress-Related CAF1 (CCR4-Associated Factor) Tj ETQq0,0,0 rgBT /Overlock 1,6 2		
17	<i>Barley stripe mosaic virus</i> β protein disrupts chloroplast antioxidant defenses to optimize viral replication. <i>EMBO Journal</i> , 2021, 40, e107660.	3.5	27
18	Highly efficient heritable genome editing in wheat using an RNA virus and bypassing tissue culture. <i>Molecular Plant</i> , 2021, 14, 1787-1798.	3.9	85

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19	A Simple Method for the Acquisition and Transmission of Brassica Yellows Virus from Transgenic Plants and Frozen Infected Leaves by Aphids. <i>Plants</i> , 2021, 10, 1944.	1.6	4
20	Comparative Analysis of Biological Characteristics among PO Proteins from Different Brassica Yellows Virus Genotypes. <i>Biology</i> , 2021, 10, 1076.	1.3	2
21	First Report of Cucurbit Aphid-Borne Yellows Virus in Passion Fruit Plants Exhibiting Mosaic and Mottling in China. <i>Plant Disease</i> , 2020, 104, 601-601.	0.7	4
22	Construction of an Infectious <i>Poa semilatifolia</i> virus cDNA Clone and Comparisons of Hordeivirus Cytopathology and Pathogenicity. <i>Phytopathology</i> , 2020, 110, 215-227.	1.1	9
23	Functional Characterization of RNA Silencing Suppressor PO from Pea Mild Chlorosis Virus. <i>International Journal of Molecular Sciences</i> , 2020, 21, 7136.	1.8	6
24	The Barley stripe mosaic virus Ω protein promotes viral cell-to-cell movement by enhancing ATPase-mediated assembly of ribonucleoprotein movement complexes. <i>PLoS Pathogens</i> , 2020, 16, e1008709.	2.1	22
25	Development of polyclonal antisera against movement proteins from three poleroviruses infecting cucurbits. <i>Phytopathology Research</i> , 2020, 2, .	0.9	1
26	Genome-Wide microRNA Profiling Using Oligonucleotide Microarray Reveals Regulatory Networks of microRNAs in <i>Nicotiana benthamiana</i> During Beet Necrotic Yellow Vein Virus Infection. <i>Viruses</i> , 2020, 12, 310.	1.5	18
27	CCR4, a RNA decay factor, is hijacked by a plant cytorhabdovirus phosphoprotein to facilitate virus replication. <i>ELife</i> , 2020, 9, .	2.8	20
28	A barley stripe mosaic virus-based guide RNA delivery system for targeted mutagenesis in wheat and maize. <i>Molecular Plant Pathology</i> , 2019, 20, 1463-1474.	2.0	91
29	Interaction between Brassica yellows virus silencing suppressor PO and plant SKP1 facilitates stability of PO <i>in vivo</i> against degradation by proteasome and autophagy pathways. <i>New Phytologist</i> , 2019, 222, 1458-1473.	3.5	41
30	Development of polyclonal antiserum against movement protein from Potato leafroll virus and its application for the virus detection. <i>Phytopathology Research</i> , 2019, 1, .	0.9	5
31	The Three Essential Motifs in PO for Suppression of RNA Silencing Activity of Potato leafroll virus Are Required for Virus Systemic Infection. <i>Viruses</i> , 2019, 11, 170.	1.5	12
32	Brassica yellows virus PO protein impairs the antiviral activity of NbRAF2 in <i>Nicotiana benthamiana</i> . <i>Journal of Experimental Botany</i> , 2018, 69, 3127-3139.	2.4	22
33	<i>Barley stripe mosaic virus</i> infection requires PKA-mediated phosphorylation of Ω for suppression of both RNA silencing and the host cell death response. <i>New Phytologist</i> , 2018, 218, 1570-1585.	3.5	40
34	Hsc70-2 is required for Beet black scorch virus infection through interaction with replication and capsid proteins. <i>Scientific Reports</i> , 2018, 8, 4526.	1.6	18
35	Barley Stripe Mosaic Virus Ω Interacts with Glycolate Oxidase and Inhibits Peroxisomal ROS Production to Facilitate Virus Infection. <i>Molecular Plant</i> , 2018, 11, 338-341.	3.9	46
36	Construction of infectious clones of lychnis ringspot virus and evaluation of its relationship with barley stripe mosaic virus by reassortment of genomic RNA segments. <i>Virus Research</i> , 2018, 243, 106-109.	1.1	10

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37	Hijacking of the nucleolar protein fibrillarin by TGB1 is required for cell-to-cell movement of <i>Barley stripe mosaic virus</i> . <i>Molecular Plant Pathology</i> , 2018, 19, 1222-1237.	2.0	41
38	Three-Dimensional Analysis of Chloroplast Structures Associated with Virus Infection. <i>Plant Physiology</i> , 2018, 176, 282-294.	2.3	62
39	Brassica yellows virus™ movement protein upregulates anthocyanin accumulation, leading to the development of purple leaf symptoms on <i>Arabidopsis thaliana</i> . <i>Scientific Reports</i> , 2018, 8, 16273.	1.6	19
40	Identification of the Potential Virulence Factors and RNA Silencing Suppressors of Mulberry Mosaic Dwarf-Associated Geminivirus. <i>Viruses</i> , 2018, 10, 472.	1.5	41
41	<i>Barley stripe mosaic virus</i> β Protein Subverts Autophagy to Promote Viral Infection by Disrupting the ATG7-ATG8 Interaction. <i>Plant Cell</i> , 2018, 30, 1582-1595.	3.1	114
42	The Conserved Proline18 in the P3a Is Important for Brassica Yellows Virus Systemic Infection. <i>Frontiers in Microbiology</i> , 2018, 9, 613.	1.5	16
43	Synergistic infection of BrYV and PEMV 2 increases the accumulations of both BrYV and BrYV-derived siRNAs in <i>Nicotiana benthamiana</i> . <i>Scientific Reports</i> , 2017, 7, 45132.	1.6	36
44	RNA-seq analysis of <i>Brachypodium distachyon</i> responses to Barley stripe mosaic virus infection. <i>Crop Journal</i> , 2017, 5, 1-10.	2.3	4
45	ICTV Virus Taxonomy Profile: Virgaviridae. <i>Journal of General Virology</i> , 2017, 98, 1999-2000.	1.3	134
46	Characterization of microRNAs of <i>Beta macrocarpa</i> and their responses to Beet necrotic yellow vein virus infection. <i>PLoS ONE</i> , 2017, 12, e0186500.	1.1	7
47	The Barley stripe mosaic virus β protein promotes chloroplast-targeted replication by enhancing unwinding of RNA duplexes. <i>PLoS Pathogens</i> , 2017, 13, e1006319.	2.1	65
48	Rice black streaked dwarf virus P7-2 forms a SCF complex through binding to <i>Oryza sativa</i> SKP1-like proteins, and interacts with GID2 involved in the gibberellin pathway. <i>PLoS ONE</i> , 2017, 12, e0177518.	1.1	28
49	A self-perpetuating repressive state of a viral replication protein blocks superinfection by the same virus. <i>PLoS Pathogens</i> , 2017, 13, e1006253.	2.1	42
50	Improved Pathogenicity of a Beet Black Scorch Virus Variant by Low Temperature and Co-infection with Its Satellite RNA. <i>Frontiers in Microbiology</i> , 2016, 7, 1771.	1.5	13
51	Simultaneous detection and differentiation of three genotypes of Brassica yellows virus by multiplex reverse transcription-polymerase chain reaction. <i>Virology Journal</i> , 2016, 13, 189.	1.4	17
52	Incomplete DRB4-dependence of the DCL4-mediated antiviral defense. <i>Scientific Reports</i> , 2016, 6, 39244.	1.6	4
53	Overexpression of hepatocyte nuclear factor 4 α in human mesenchymal stem cells suppresses hepatocellular carcinoma development through Wnt/ β -catenin signaling pathway downregulation. <i>Cancer Biology and Therapy</i> , 2016, 17, 558-565.	1.5	43
54	Induction of heat shock protein 27 by bicyclol attenuates d-galactosamine/lipopolysaccharide-induced liver injury. <i>European Journal of Pharmacology</i> , 2016, 791, 482-490.	1.7	11

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55	Two amino acids near the N-terminus of Cucumber mosaic virus 2b play critical roles in the suppression of RNA silencing and viral infectivity. <i>Molecular Plant Pathology</i> , 2016, 17, 173-183.	2.0	33
56	Random Plant Viral Variants Attain Temporal Advantages During Systemic Infections and in Turn Resist other Variants of the Same Virus. <i>Scientific Reports</i> , 2015, 5, 15346.	1.6	24
57	Phosphorylation of Beet black scorch virus coat protein by PKA is required for assembly and stability of virus particles. <i>Scientific Reports</i> , 2015, 5, 11585.	1.6	26
58	Transcriptome Analysis of Beta macrocarpa and Identification of Differentially Expressed Transcripts in Response to Beet Necrotic Yellow Vein Virus Infection. <i>PLoS ONE</i> , 2015, 10, e0132277.	1.1	11
59	Risk factors for new onset diabetes mellitus after liver transplantation: A meta-analysis. <i>World Journal of Gastroenterology</i> , 2015, 21, 6329.	1.4	56
60	Phosphorylation of TGB1 by protein kinase CK2 promotes barley stripe mosaic virus movement in monocots and dicots. <i>Journal of Experimental Botany</i> , 2015, 66, 4733-4747.	2.4	44
61	Morphogenesis of Endoplasmic Reticulum Membrane-Invaginated Vesicles during Beet Black Scorch Virus Infection: Role of Auxiliary Replication Protein and New Implications of Three-Dimensional Architecture. <i>Journal of Virology</i> , 2015, 89, 6184-6195.	1.5	56
62	Development of three full-length infectious cDNA clones of distinct brassica yellows virus genotypes for agrobacterium-mediated inoculation. <i>Virus Research</i> , 2015, 197, 13-16.	1.1	25
63	Architecture of viral replication factories. <i>Oncotarget</i> , 2015, 6, 30439-30440.	0.8	4
64	Contrast-enhanced micro-computed tomography using ExiTron nano6000 for assessment of liver injury. <i>World Journal of Gastroenterology</i> , 2015, 21, 8043.	1.4	4
65	Deep Sequencing-Based Transcriptome Profiling Reveals Comprehensive Insights into the Responses of <i>Nicotiana benthamiana</i> to Beet necrotic yellow vein virus Infections Containing or Lacking RNA4. <i>PLoS ONE</i> , 2014, 9, e85284.	1.1	26
66	Infection of Beet necrotic yellow vein virus with RNA4-encoded P31 specifically up-regulates pathogenesis-related protein 10 in <i>Nicotiana benthamiana</i> . <i>Virology Journal</i> , 2014, 11, 118.	1.4	19
67	Complete genome sequence analysis identifies a new genotype of brassica yellows virus that infects cabbage and radish in China. <i>Archives of Virology</i> , 2014, 159, 2177-2180.	0.9	29
68	Amino Acid Sequence Motifs Essential for PO-Mediated Suppression of RNA Silencing in an Isolate of Potato leafroll virus from Inner Mongolia. <i>Molecular Plant-Microbe Interactions</i> , 2014, 27, 515-527.	1.4	47
69	First Report of Potato Virus H on <i>Solanum muricatum</i> in China. <i>Plant Disease</i> , 2014, 98, 1016-1016.	0.7	11
70	First Report of Barley yellow striate mosaic virus on Wheat in China. <i>Plant Disease</i> , 2014, 98, 1450-1450.	0.7	26
71	N-terminal basic amino acid residues of Beet black scorch virus capsid protein play a critical role in virion assembly and systemic movement. <i>Virology Journal</i> , 2013, 10, 200.	1.4	20
72	Efficient and fine mapping of RMES1 conferring resistance to sorghum aphid <i>Melanaphis sacchari</i> . <i>Molecular Breeding</i> , 2013, 31, 777-784.	1.0	28

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73	Nonstructural protein P7-2 encoded by Rice black-streaked dwarf virus interacts with SKP1, a core subunit of SCF ubiquitin ligase. <i>Virology Journal</i> , 2013, 10, 325.	1.4	30
74	Selection of reference genes for gene expression studies in virus-infected monocots using quantitative real-time PCR. <i>Journal of Biotechnology</i> , 2013, 168, 7-14.	1.9	33
75	Enhanced Virus Resistance in Transgenic Maize Expressing a dsRNA-Specific Endoribonuclease Gene from <i>E. coli</i> . <i>PLoS ONE</i> , 2013, 8, e60829.	1.1	17
76	Discovery and Characterization of a Novel Carlavirus Infecting Potatoes in China. <i>PLoS ONE</i> , 2013, 8, e69255.	1.1	28
77	<i>Trichosanthes kirilowii</i> : A New Host of Cucurbit mild mosaic virus in China. <i>Plant Disease</i> , 2013, 97, 1388-1388.	0.7	5
78	Identification of an Internal RNA Element Essential for Replication and Translational Enhancement of Tobacco Necrosis Virus AC. <i>PLoS ONE</i> , 2013, 8, e57938.	1.1	5
79	Temperature-Dependent Survival of Turnip Crinkle Virus-Infected Arabidopsis Plants Relies on an RNA Silencing-Based Defense That Requires DCL2, AGO2, and HEN1. <i>Journal of Virology</i> , 2012, 86, 6847-6854.	1.5	168
80	Two distinct sites are essential for virulent infection and support of variant satellite RNA replication in spontaneous beet black scorch virus variants. <i>Journal of General Virology</i> , 2012, 93, 2718-2728.	1.3	10
81	The R-rich motif of Beet black scorch virus P7a movement protein is important for the nuclear localization, nucleolar targeting and viral infectivity. <i>Virus Research</i> , 2012, 167, 207-218.	1.1	15
82	Brachypodium distachyon line Bd3-1 resistance is elicited by the barley stripe mosaic virus triple gene block 1 movement protein. <i>Journal of General Virology</i> , 2012, 93, 2729-2739.	1.3	33
83	Validation of Reference Genes for Gene Expression Studies in Virus-Infected <i>Nicotiana benthamiana</i> Using Quantitative Real-Time PCR. <i>PLoS ONE</i> , 2012, 7, e46451.	1.1	337
84	Nucleotide sequence of a chickpea chlorotic stunt virus relative that infects pea and faba bean in China. <i>Archives of Virology</i> , 2012, 157, 1393-1396.	0.9	6
85	Complete genomic sequence analysis reveals a novel fabavirus infecting cucurbits in China. <i>Archives of Virology</i> , 2012, 157, 597-600.	0.9	16
86	Fine Mapping of the Bsr1 Barley Stripe Mosaic Virus Resistance Gene in the Model Grass <i>Brachypodium distachyon</i> . <i>PLoS ONE</i> , 2012, 7, e38333.	1.1	67
87	The Complete Genome Sequence of Southern rice black-streaked dwarf virus Isolated from Vietnam. <i>Plant Pathology Journal</i> , 2012, 28, 428-432.	0.7	5
88	Rapid Detection and Differentiation of Three Cucurbit-infecting Poloroviruses by Multiplex RT-PCR. <i>Journal of Agricultural Science</i> , 2012, 4, .	0.1	1
89	Nuclear localization of Beet black scorch virus capsid protein and its interaction with importin β . <i>Virus Research</i> , 2011, 155, 307-315.	1.1	26
90	Studies on interaction of cucurbit aphid-borne yellow virus proteins using yeast two-hybrid system and bimolecular fluorescence complementation. <i>Acta Virologica</i> , 2011, 55, 235-241.	0.3	0

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91	Detection and characterization of spontaneous internal deletion mutants of Beet Necrotic yellow vein virus RNA3 from systemic host <i>Nicotiana benthamiana</i> . <i>Virology Journal</i> , 2011, 8, 335.	1.4	10
92	Rapid detection of wheat yellow mosaic virus by reverse transcription loop-mediated isothermal amplification. <i>Virology Journal</i> , 2011, 8, 550.	1.4	34
93	A novel strain of Beet western yellows virus infecting sugar beet with two distinct genotypes differing in the 5' terminal half of genome. <i>Virus Genes</i> , 2011, 42, 141-149.	0.7	10
94	Molecular characterization of two genotypes of a new polerovirus infecting brassicas in China. <i>Archives of Virology</i> , 2011, 156, 2251-2255.	0.9	47
95	Rice black-streaked dwarf virus P6 self-interacts to form punctate, viroplasm-like structures in the cytoplasm and recruits viroplasm-associated protein P9-1. <i>Virology Journal</i> , 2011, 8, 24.	1.4	37
96	A High Throughput Barley Stripe Mosaic Virus Vector for Virus Induced Gene Silencing in Monocots and Dicots. <i>PLoS ONE</i> , 2011, 6, e26468.	1.1	253
97	Development and Optimization of Tobacco necrosis virus A Induced Gene Silencing in <i>Nicotiana benthamiana</i> . <i>Progress in Biochemistry and Biophysics</i> , 2011, 38, 919-928.	0.3	2
98	Molecular characterization of two Chinese isolates of Beet western yellows virus infecting sugar beet. <i>Virus Genes</i> , 2010, 41, 105-110.	0.7	6
99	Ring structure amino acids affect the suppressor activity of melon aphid-borne yellows virus PO protein. <i>Virology</i> , 2010, 406, 21-27.	1.1	31
100	Development of Tobacco necrosis virus A as a vector for efficient and stable expression of FMDV VP1 peptides. <i>Plant Biotechnology Journal</i> , 2010, 8, 506-523.	4.1	28
101	Oral administration of plant-based rotavirus VP6 induces antigen-specific IgAs, IgGs and passive protection in mice. <i>Vaccine</i> , 2010, 28, 6021-6027.	1.7	29
102	Production of Antiserum to Recombinant Coat Protein for Detecting Lily mottle virus in Yunnan, China. <i>Journal of Phytopathology</i> , 2009, 157, 362-369.	0.5	5
103	Using porphyritic andesite as a new additive for improving hydrolysis and acidogenesis of solid organic wastes. <i>Bioresource Technology</i> , 2009, 100, 5594-5599.	4.8	16
104	Distribution and molecular diversity of three cucurbit-infecting poleroviruses in China. <i>Virus Research</i> , 2009, 145, 341-346.	1.1	39
105	Complete nucleotide sequence of a new strain of Tobacco necrosis virus A infecting soybean in China and infectivity of its full-length cDNA clone. <i>Virus Genes</i> , 2008, 36, 259-266.	0.7	19
106	Phylogenetic analysis of Beet necrotic yellow vein virus isolates from China. <i>Virus Genes</i> , 2008, 36, 429-432.	0.7	21
107	Effects on the local symptoms of subgenomic RNAs expressions and their translational products of Tobacco necrosis virus A Chinese isolate. <i>Science Bulletin</i> , 2008, 53, 1682-1690.	4.3	7
108	Complete sequence analysis reveals two distinct poleroviruses infecting cucurbits in China. <i>Archives of Virology</i> , 2008, 153, 1155-1160.	0.9	53

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109	First identification of Beet western yellows virus on sugarbeet and lettuce in China. <i>Plant Pathology</i> , 2008, 57, 390-390.	1.2	5
110	Complete Genome Sequences of Two Chinese Beet soil-borne virus Isolates Provide Evidence that the Genome is Highly Conserved. <i>Journal of Phytopathology</i> , 2008, 156, 487-488.	0.5	3
111	First report on the occurrence of Cucurbit aphid-borne yellows virus on nine cucurbitaceous species in China. <i>Plant Pathology</i> , 2008, 57, 390-390.	1.2	22
112	First report of Beet soil-borne virus on sugar beet in China. <i>Plant Pathology</i> , 2008, 57, 389-389.	1.2	5
113	Specificity of ARGONAUTE7-miR390 Interaction and Dual Functionality in TAS3 Trans-Acting siRNA Formation. <i>Cell</i> , 2008, 133, 128-141.	13.5	712
114	Molecular characterization of two Chinese isolates of Beet mosaic virus. <i>Virus Genes</i> , 2007, 35, 795-799.	0.7	6
115	Analysis of the subgenomic RNAs and the small open reading frames of Beet black scorch virus. <i>Journal of General Virology</i> , 2006, 87, 3077-3086.	1.3	30
116	Development of an ID-ELISA for the detection of Rice black-streaked dwarf virus in plants. <i>Journal of Virological Methods</i> , 2006, 134, 61-65.	1.0	32
117	Two virus-encoded RNA silencing suppressors, P14 of Beet necrotic yellow vein virus and S6 of Rice black streak dwarf virus. <i>Science Bulletin</i> , 2005, 50, 305-310.	1.7	21
118	Two virus-encoded RNA si-lencing suppressors, P14 of Beet necrotic yellow vein virus and S6 of Rice black streak dwarf virus. <i>Science Bulletin</i> , 2005, 50, 305.	1.7	3
119	Analysis of Nucleotide Sequences and Multimeric Forms of a Novel Satellite RNA Associated with Beet Black Scorch Virus. <i>Journal of Virology</i> , 2005, 79, 3664-3674.	1.5	26
120	Creation of trivalent transgenic watermelon resistant to virus infection. <i>Chinese Journal of Agricultural Biotechnology</i> , 2005, 2, 179-185.	0.1	3
121	Insights into the Oligomeric States, Conformational Changes, and Helicase Activities of SV40 Large Tumor Antigen. <i>Journal of Biological Chemistry</i> , 2004, 279, 38952-38959.	1.6	43
122	Mechanisms of Conformational Change for a Replicative Hexameric Helicase of SV40 Large Tumor Antigen. <i>Cell</i> , 2004, 119, 47-60.	13.5	291
123	Sequence analysis of the complete genome of rice black-streaked dwarf virus isolated from maize with rough dwarf disease. <i>Virus Genes</i> , 2003, 27, 163-168.	0.7	107
124	Structure of the replicative helicase of the oncoprotein SV40 large tumour antigen. <i>Nature</i> , 2003, 423, 512-518.	13.7	278
125	Effect of the RNA5 component on pathogenicity of beet ne-crotic yellow vein virus. <i>Science Bulletin</i> , 2003, 48, 796.	1.7	0
126	Functional analysis of beet necrotic yellow vein virus (BNYVV) RNA4 in fungal transmission. <i>Science Bulletin</i> , 2002, 47, 1281.	1.7	2

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127	The complete nucleotide sequence of Beet black scorch virus (BBSV), a new member of the genus Necrovirus. Archives of Virology, 2002, 147, 2431-2435.	0.9	28
128	Identification of rice black-streaked dwarf fijivirus in maize with rough dwarf disease in China. Archives of Virology, 2001, 146, 167-170.	0.9	78
129	Structure of Complement Receptor 2 in Complex with Its C3d Ligand. Science, 2001, 292, 1725-1728.	6.0	148
130	Wheat yellow mosaic virus Widely Occurring in Wheat (Triticum aestivum) in China. Plant Disease, 2000, 84, 627-630.	0.7	79
131	Over-expression of 72 ku protein of wheat yellow mosaic virus in E. coli and preparation of its antiserum. Science Bulletin, 2000, 45, 525-528.	1.7	3
132	Analysis of nucleotide sequence of wheat yellow mosaic virus genomic RNAs. Science in China Series C: Life Sciences, 1999, 42, 554-560.	1.3	11
133	The nucleotide sequence of a Chinese isolate of wheat yellow mosaic virus and its comparison with a Japanese isolate. Archives of Virology, 1999, 144, 2201-2206.	0.9	8