Zhi-Ming Yuan

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3944468/publications.pdf

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138 papers 5,382 citations

32 h-index 65 g-index

147 all docs

 $\begin{array}{c} 147 \\ \text{docs citations} \end{array}$

147 times ranked

9474 citing authors

#	Article	IF	CITATIONS
1	A human neutralizing antibody targets the receptor-binding site of SARS-CoV-2. Nature, 2020, 584, 120-124.	13.7	1,237
2	Effect of an Inactivated Vaccine Against SARS-CoV-2 on Safety and Immunogenicity Outcomes. JAMA - Journal of the American Medical Association, 2020, 324, 951.	3.8	671
3	An adenovirus-vectored COVID-19 vaccine confers protection from SARS-COV-2 challenge in rhesus macaques. Nature Communications, 2020, 11, 4207.	5 . 8	194
4	Infection with novel coronavirus (SARS-CoV-2) causes pneumonia in Rhesus macaques. Cell Research, 2020, 30, 670-677.	5.7	194
5	Cyclosporine Inhibits Flavivirus Replication through Blocking the Interaction between Host Cyclophilins and Viral NS5 Protein. Antimicrobial Agents and Chemotherapy, 2009, 53, 3226-3235.	1.4	116
6	Characterization of Dengue Virus NS4A and NS4B Protein Interaction. Journal of Virology, 2015, 89, 3455-3470.	1.5	116
7	High-Level Field Resistance to Bacillus sphaericus C3-41 in Culex quinquefasciatus from Southern China. Biocontrol Science and Technology, 2000, 10, 41-49.	0.5	100
8	Gemcitabine, lycorine and oxysophoridine inhibit novel coronavirus (SARS-CoV-2) in cell culture. Emerging Microbes and Infections, 2020, 9, 1170-1173.	3.0	100
9	Two Distinct Sets of NS2A Molecules Are Responsible for Dengue Virus RNA Synthesis and Virion Assembly. Journal of Virology, 2015, 89, 1298-1313.	1.5	90
10	Mapping the Interactions between the NS4B and NS3 Proteins of Dengue Virus. Journal of Virology, 2015, 89, 3471-3483.	1.5	83
11	Complete Genome Sequence of the Mosquitocidal Bacterium <i>Bacillus sphaericus</i> C3-41 and Comparison with Those of Closely Related Bacillus Species. Journal of Bacteriology, 2008, 190, 2892-2902.	1.0	80
12	The occurrence of Bacillus cereus, B. thuringiensis and B. mycoides in Chinese pasteurized full fat milk. International Journal of Food Microbiology, 2008, 121, 195-200.	2.1	74
13	Metagenomic Virome Analysis of Culex Mosquitoes from Kenya and China. Viruses, 2018, 10, 30.	1.5	74
14	A Single Amino Acid in Nonstructural Protein NS4B Confers Virulence to Dengue Virus in AG129 Mice through Enhancement of Viral RNA Synthesis. Journal of Virology, 2011, 85, 7775-7787.	1.5	73
15	A new Cry toxin with a unique twoâ€component dependency from <i>Bacillus sphaericus</i> . FASEB Journal, 2007, 21, 4112-4120.	0.2	69
16	Identification of palmatine as an inhibitor of West Nile virus. Archives of Virology, 2010, 155, 1325-1329.	0.9	68
17	A SARS-CoV-2 neutralizing antibody with extensive Spike binding coverage and modified for optimal therapeutic outcomes. Nature Communications, 2021, 12, 2623.	5 . 8	64
18	Dengue virus subgenomic RNA induces apoptosis through the Bcl-2-mediated PI3k/Akt signaling pathway. Virology, 2014, 448, 15-25.	1.1	63

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19	Mosquito-Associated Viruses in China. Virologica Sinica, 2018, 33, 5-20.	1.2	59
20	Characterization of neutralizing antibody with prophylactic and therapeutic efficacy against SARS-CoV-2 in rhesus monkeys. Nature Communications, 2020, 11, 5752.	5.8	59
21	Comparison of Genotypes I and III in Japanese Encephalitis Virus Reveals Distinct Differences in Their Genetic and Host Diversity. Journal of Virology, 2014, 88, 11469-11479.	1.5	55
22	A Metagenomic Survey of Viral Abundance and Diversity in Mosquitoes from Hubei Province. PLoS ONE, 2015, 10, e0129845.	1.1	53
23	Rapid detection of filoviruses by real-time TaqMan polymerase chain reaction assays. Virologica Sinica, 2012, 27, 273-277.	1.2	52
24	Modulation of a thermoregulated type VI secretion system by AHL-dependent Quorum Sensing in Yersinia pseudotuberculosis. Archives of Microbiology, 2011, 193, 351-63.	1.0	50
25	Transmembrane Domains of NS2B Contribute to both Viral RNA Replication and Particle Formation in Japanese Encephalitis Virus. Journal of Virology, 2016, 90, 5735-5749.	1.5	48
26	The discovery and global distribution of novel mosquitoâ€essociated viruses in the last decade (2007â€2017). Reviews in Medical Virology, 2019, 29, e2079.	3.9	48
27	Comparative Metagenomic Profiling of Viromes Associated with Four Common Mosquito Species in China. Virologica Sinica, 2018, 33, 59-66.	1.2	46
28	Conjugative transfer, stability and expression of a plasmid encoding acry1Ac gene inBacillus cereusgroup strains. FEMS Microbiology Letters, 2004, 231, 45-52.	0.7	41
29	A cell-based large-scale screening of natural compounds for inhibitors of SARS-CoV-2. Signal Transduction and Targeted Therapy, 2020, 5, 218.	7.1	41
30	Recovery of a chemically synthesized Japanese encephalitis virus reveals two critical adaptive mutations in NS2B and NS4A. Journal of General Virology, 2014, 95, 806-815.	1.3	40
31	Stability of the Virome in Lab- and Field-Collected Aedes albopictus Mosquitoes across Different Developmental Stages and Possible Core Viruses in the Publicly Available Virome Data of <i>Aedes</i> Mosquitoes. MSystems, 2020, 5, .	1.7	40
32	The Interface between Methyltransferase and Polymerase of NS5 Is Essential for Flavivirus Replication. PLoS Neglected Tropical Diseases, 2014, 8, e2891.	1.3	38
33	Infectious Chikungunya Virus (CHIKV) with a Complete Capsid Deletion: a New Approach for a CHIKV Vaccine. Journal of Virology, 2019, 93, .	1.5	36
34	Single nucleotide deletion of cqm1 gene results in the development of resistance to Bacillus sphaericus in Culex quinquefasciatus. Journal of Insect Physiology, 2013, 59, 967-973.	0.9	35
35	The genetic diversity of cereulide biosynthesis gene cluster indicates a composite transposon Tnces in emetic Bacillus weihenstephanensis. BMC Microbiology, 2014, 14, 149.	1.3	33
36	Occurrence of psychrotolerant Bacillus cereus group strains in ice creams. International Journal of Food Microbiology, 2010, 137, 143-146.	2.1	29

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37	Critical role of Dengue Virus NS1 protein in viral replication. Virologica Sinica, 2014, 29, 162-169.	1.2	29
38	Transfer and expression of the mosquitocidal plasmid pBtoxis in Bacillus cereus group strains. FEMS Microbiology Letters, 2005, 245, 239-247.	0.7	28
39	Molecular Characterization of a Glucokinase with Broad Hexose Specificity from Bacillus sphaericus Strain C3-41. Applied and Environmental Microbiology, 2007, 73, 3581-3586.	1.4	28
40	Low toxicity and high immunogenicity of an inactivated vaccine candidate against COVID-19 in different animal models. Emerging Microbes and Infections, 2020, 9, 2606-2618.	3.0	28
41	Safety and immunogenicity of an inactivated SARS-CoV-2 vaccine in healthy adults aged 18 years or older: A randomized, double-blind, placebo-controlled, phase 1/2 trial. EClinicalMedicine, 2021, 38, 101010.	3.2	28
42	Genome comparison provides molecular insights into the phylogeny of the reassigned new genus Lysinibacillus. BMC Genomics, 2015, 16, 140.	1.2	25
43	Genetic interaction between NS4A and NS4B for replication of Japanese encephalitis virus. Journal of General Virology, 2015, 96, 1264-1275.	1.3	24
44	A new tubRZ operon involved in the maintenance of the Bacillus sphaericus mosquitocidal plasmid pBsph. Microbiology (United Kingdom), 2014, 160, 1112-1124.	0.7	23
45	Zika virus pathogenesis and current therapeutic advances. Pathogens and Global Health, 2021, 115, 21-39.	1.0	23
46	RBD-homodimer, a COVID-19 subunit vaccine candidate, elicits immunogenicity and protection in rodents and nonhuman primates. Cell Discovery, 2021, 7, 82.	3.1	22
47	Allelic Diversity and Population Structure of Bacillus sphaericus as Revealed by Multilocus Sequence Typing. Applied and Environmental Microbiology, 2011, 77, 5553-5556.	1.4	21
48	Development of a stable Gaussia luciferase enterovirus 71 reporter virus. Journal of Virological Methods, 2015, 219, 62-66.	1.0	21
49	Development of Neutralization Assay Using an eGFP Chikungunya Virus. Viruses, 2016, 8, 181.	1.5	21
50	Effects of exogenous aerobic bacteria on methane production and biodegradation of municipal solid waste in bioreactors. Waste Management, 2016, 55, 93-98.	3.7	21
51	A mouse model for SARS-CoV-2 infection by exogenous delivery of hACE2 using alphavirus replicon particles. Cell Research, 2020, 30, 1046-1048.	5.7	21
52	Identification and genomic comparison of temperate bacteriophages derived from emetic Bacillus cereus. PLoS ONE, 2017, 12, e0184572.	1.1	20
53	Identifying the pattern of molecular evolution for Zaire ebolavirus in the 2014 outbreak in West Africa. Infection, Genetics and Evolution, 2015, 32, 51-59.	1.0	19
54	First Isolation and Characterization of a Group C Banna Virus (BAV) from Anopheles sinensis Mosquitoes in Hubei, China. Viruses, 2018, 10, 555.	1.5	19

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55	Isolation and characterization of a novel mesonivirus from Culex mosquitoes in China. Virus Research, 2017, 240, 130-139.	1.1	18
56	Proteolytic Stability of Insecticidal Toxins Expressed in Recombinant Bacilli. Applied and Environmental Microbiology, 2007, 73, 218-225.	1.4	17
57	Mosquitoes of Etiological Concern in Kenya and Possible Control Strategies. Insects, 2019, 10, 173.	1.0	17
58	A dataset of distribution and diversity of mosquito-associated viruses and their mosquito vectors in China. Scientific Data, 2020, 7, 342.	2.4	17
59	Biochemical and antigenic characterization of the structural proteins and their post-translational modifications in purified SARS-CoV-2 virions of an inactivated vaccine candidate. Emerging Microbes and Infections, 2020, 9, 2653-2662.	3.0	17
60	Protective Efficacy of Inactivated Vaccine against SARS-CoV-2 Infection in Mice and Non-Human Primates. Virologica Sinica, 2021, 36, 879-889.	1.2	17
61	Inhibition of japanese encephalitis virus infection by flavivirus recombinant e protein domain III. Virologica Sinica, 2013, 28, 152-160.	1.2	16
62	CcpA-Mediated Enhancement of Sugar and Amino Acid Metabolism in <i>Lysinibacillus sphaericus</i> by NMR-Based Metabolomics. Journal of Proteome Research, 2012, 11, 4654-4661.	1.8	15
63	A replication-defective Japanese encephalitis virus (JEV) vaccine candidate with NS1 deletion confers dual protection against JEV and West Nile virus in mice. Npj Vaccines, 2020, 5, 73.	2.9	15
64	Phylogenetic Analysis and Heterologous Expression of Surface Layer Protein SlpC ofBacillus sphaericusC3-41. Bioscience, Biotechnology and Biochemistry, 2008, 72, 1257-1263.	0.6	14
65	Biodegradation of Methyl tert-Butyl Ether by Enriched Bacterial Culture. Current Microbiology, 2009, 59, 30-34.	1.0	14
66	Characterization of Ebinur Lake Virus and Its Human Seroprevalence at the China–Kazakhstan Border. Frontiers in Microbiology, 2019, 10, 3111.	1.5	14
67	A Novel Transcriptional Activator, tubX , Is Required for the Stability of Bacillus sphaericus Mosquitocidal Plasmid pBsph. Journal of Bacteriology, 2014, 196, 4304-4314.	1.0	13
68	Polyphosphate kinase of Lysinibacillus sphaericus and its effects on accumulation of polyphosphate and bacterial growth. Microbiological Research, 2015, 172, 41-47.	2.5	13
69	Generation and characterization of West Nile pseudo-infectious reporter virus for antiviral screening. Antiviral Research, 2017, 141, 38-47.	1.9	13
70	A human antibody of potent efficacy against SARS-CoV-2 in rhesus macaques showed strong blocking activity to B.1.351. MAbs, 2021, 13, 1930636.	2.6	13
71	The residual occurrences of Bacillus thuringiensis biopesticides in food and beverages. International Journal of Food Microbiology, 2008, 127, 68-72.	2.1	12
72	Characterization of three autolysins with activity against cereulide-producing Bacillus isolates in food matrices. International Journal of Food Microbiology, 2017, 241, 291-297.	2.1	12

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73	Homologous RNA secondary structure duplications in $3\hat{a}\in^2$ untranslated region influence subgenomic RNA production and replication of dengue virus. Virology, 2018, 524, 114-126.	1.1	12
74	Visualization of chikungunya virus infection <i>in vitro</i> and <i>in vivo</i> . Emerging Microbes and Infections, 2019, 8, 1574-1583.	3.0	12
75	Characterization of a Novel Tanay Virus Isolated From Anopheles sinensis Mosquitoes in Yunnan, China. Frontiers in Microbiology, 2019, 10, 1963.	1.5	12
76	SARS-CoV-2 Does Not Replicate in Aedes Mosquito Cells nor Present in Field-Caught Mosquitoes from Wuhan. Virologica Sinica, 2020, 35, 355-358.	1.2	12
77	Cloning and Expression of the Binary Toxin Gene from Bacillus sphaericus IAB872 in a Crystal-Minus Bacillus thuringiensis subsp. israelensis. Current Microbiology, 2001, 43, 21-25.	1.0	11
78	Generation of mariner-based transposon insertion mutant library of Bacillus sphaericus 2297 and investigation of genes involved in sporulation and mosquito-larvicidal crystal protein synthesis. FEMS Microbiology Letters, 2012, 330, 105-112.	0.7	11
79	Short Direct Repeats in the 3′ Untranslated Region Are Involved in Subgenomic Flaviviral RNA Production. Journal of Virology, 2020, 94, .	1.5	11
80	Different Degrees of 5'-to-3' DAR Interactions Modulate Zika Virus Genome Cyclization and Host-Specific Replication. Journal of Virology, 2020, 94, .	1.5	11
81	Transmission competence of a new mesonivirus, Yichang virus, in mosquitoes and its interference with representative flaviviruses. PLoS Neglected Tropical Diseases, 2020, 14, e0008920.	1.3	11
82	Coexpression of cyt1Aa of Bacillus thuringiensis subsp. israelensis with Bacillus sphaericus Binary Toxin Gene in Acrystalliferous Strain of B. thuringiensis. Current Microbiology, 2000, 40, 322-326.	1.0	10
83	Development and evaluation of oneâ€step multiplex realâ€time RTâ€PCR assay for simultaneous detection of Zika virus and Chikungunya virus. Journal of Medical Virology, 2018, 90, 389-396.	2.5	10
84	Increased morbidity of obese mice infected with mouse-adapted SARS-CoV-2. Cell Discovery, 2021, 7, 74.	3.1	10
85	Detection of Enterotoxin Genes in Mosquito-Larvicidal Bacillus Species. Current Microbiology, 2002, 45, 221-225.	1.0	9
86	Nucleoprotein-based indirect enzyme-linked immunosorbent assay (indirect ELISA) for detecting antibodies specific to Ebola virus and Marbug virus. Virologica Sinica, 2014, 29, 372-380.	1.2	9
87	The LspC3–41I restriction-modification system is the major determinant for genetic manipulations of Lysinibacillus sphaericus C3–41. BMC Microbiology, 2017, 17, 116.	1.3	9
88	Rapid detection of Banna virus by reverse transcription-loop-mediated isothermal amplification (RT-LAMP). International Journal of Infectious Diseases, 2019, 78, 93-98.	1.5	9
89	Replication-Defective West Nile Virus with NS1 Deletion as a New Vaccine Platform for Flavivirus. Journal of Virology, 2019, 93, .	1.5	9
90	Pathogenesis and Immune Response of Ebinur Lake Virus: A Newly Identified Orthobunyavirus That Exhibited Strong Virulence in Mice. Frontiers in Microbiology, 2020, 11, 625661.	1.5	9

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91	Evaluating the virucidal activity of four disinfectants against SARS-CoV-2. American Journal of Infection Control, 2022, 50, 319-324.	1.1	9
92	mRNA based vaccines provide broad protection against different SARS-CoV-2 variants of concern. Emerging Microbes and Infections, 2022, 11, 1550-1553.	3.0	9
93	Species-Specific Cell Wall Binding Affinity of the S-Layer Proteins of Mosquitocidal Bacterium <i>Bacillus sphaericus </i> C3-41. Applied and Environmental Microbiology, 2009, 75, 3891-3895.	1.4	8
94	Regulator DegU is required for multicellular behavior in Lysinibacillus sphaericus. Research in Microbiology, 2018, 169, 177-187.	1.0	8
95	CesH Represses Cereulide Synthesis as an Alpha/Beta Fold Hydrolase in Bacillus cereus. Toxins, 2019, 11, 231.	1.5	8
96	Rational design of West Nile virus vaccine through large replacement of 3′ UTR with internal poly(A). EMBO Molecular Medicine, 2021, 13, e14108.	3.3	8
97	Dynamic Surveillance of Mosquitoes and Their Viromes in Wuhan During 2020. Zoonoses, 2021, 1, .	0.5	8
98	Detection and phylogenic analysis of one anthrax virulence plasmid pXO1 conservative open reading frame ubiquitous presented within Bacillus cereus group strains. Biochemical and Biophysical Research Communications, 2006, 349, 1214-1219.	1.0	7
99	Construction and characterization of the interdomain chimeras using CryllAa and CryllBa from Bacillus thuringiensis and identification of a possible novel toxic chimera. Biotechnology Letters, 2014, 36, 105-111.	1.1	7
100	Development and characterization of West Nile virus replicon expressing secreted Gaussia Luciferase. Virologica Sinica, 2013, 28, 161-166.	1.2	6
101	Collagen-Like Glycoprotein BclS Is Involved in the Formation of Filamentous Structures of the Lysinibacillus sphaericus Exosporium. Applied and Environmental Microbiology, 2014, 80, 6656-6663.	1.4	6
102	The 2014 Ebola virus outbreak in West Africa highlights no evidence of rapid evolution or adaptation to humans. Scientific Reports, 2016, 6, 35822.	1.6	6
103	Influences of magnetic powder addition on the anaerobic digestion of municipal dewatered sludge. Environmental Progress and Sustainable Energy, 2019, 38, 374-379.	1.3	6
104	Characterization of two newly isolated bacteriophages PW2 and PW4 and derived endolysins with lysis activity against Bacillus cereus group strains. Virus Research, 2021, 302, 198489.	1.1	6
105	Title is missing!. World Journal of Microbiology and Biotechnology, 2001, 17, 385-389.	1.7	5
106	Biosafety Level 4 Laboratory User Training Program, China. Emerging Infectious Diseases, 2019, 25, .	2.0	5
107	The Impact of Exogenous Aerobic Bacteria on Sustainable Methane Production Associated with Municipal Solid Waste Biodegradation: Revealed by High-Throughput Sequencing. Sustainability, 2020, 12, 1815.	1.6	5
108	Horizontal transfer of large plasmid with type IV secretion system and mosquitocidal genomic island with excision and integration capabilities in Lysinibacillus sphaericus. Environmental Microbiology, 2021, 23, 5131-5146.	1.8	5

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109	Characterization of a novel reassortment Tibet orbivirus isolated from Culicoides spp. in Yunnan, PR China. Journal of General Virology, 2021, 102, .	1.3	5
110	Mutagenesis of D80-82 and G83 residues in West Nile Virus NS2B: Effects on NS2B-NS3 activity and viral replication. Virologica Sinica, 2013, 28, 16-23.	1.2	4
111	Mapping the virome in lab-reared and wild-caught Aedes albopictus mosquitoes. Access Microbiology, $2019,1,\ldots$	0.2	4
112	Substrate preference of 5′-methylthioadenosine/S-adenosylhomocysteine nucleosidase inBurkholderia thailandensis. FEMS Microbiology Letters, 2013, 339, 110-116.	0.7	4
113	High-containment facilities and the role they play in global health security. Journal of Biosafety and Biosecurity, 2022, 4, 1-4.	1.4	4
114	Infection and pathogenesis of the Delta variant of SARS-CoV-2 in Rhesus macaque. Virologica Sinica, 2022, , .	1.2	4
115	Surface display of domain III of Japanese encephalitis virus E protein on Salmonella typhimurium by using an ice nucleation protein. Virologica Sinica, 2011, 26, 409-417.	1,2	3
116	Nanoscale mono- and multi-layer cylinder structures formed by recombinant S-layer proteins of mosquitocidal Bacillus sphaericus C3-41. Applied Microbiology and Biotechnology, 2013, 97, 7275-7283.	1.7	3
117	Complete Genome Sequence of a New Strain of Tanay Virus, Isolate YN15_103_01, from Yunnan, China. Genome Announcements, 2018, 6, .	0.8	3
118	Effects of Propoxur Exposure on Insecticidal Susceptibility and Developmental Traits in Culex pipiens quinquefasciatus. Insects, 2019, 10, 288.	1.0	3
119	Networking for training Level 3/4 biosafety laboratory staff. Journal of Biosafety and Biosecurity, 2019, 1, 46-49.	1.4	3
120	Efficacy of disinfectants for inactivation of Ebola virus in suspension by integrated cell culture coupled with real-time RT–PCR. Journal of Hospital Infection, 2022, 125, 67-74.	1.4	3
121	A toxin-antitoxin system is essential for the stability of mosquitocidal plasmid pBsph of Lysinibacillus sphaericus. Microbiological Research, 2018, 214, 114-122.	2.5	2
122	Application of Bacillus thuringiensis strains with conjugal and mobilizing capability drives gene transmissibility within Bacillus cereus group populations in confined habitats. BMC Microbiology, 2020, 20, 363.	1.3	2
123	Comparison and Evaluation of Real-Time Taqman PCR for Detection and Quantification of Ebolavirus. Viruses, 2021, 13, 1575.	1.5	2
124	Bacterial cell surface display: a method for studying Japanese encephalitis virus pathogenicity. Japanese Journal of Infectious Diseases, 2009, 62, 402-8.	0.5	2
125	Vector competence and immune response of Aedes aegypti for Ebinur Lake virus, a newly classified mosquito-borne orthobunyavirus. PLoS Neglected Tropical Diseases, 2022, 16, e0010642.	1.3	2
126	A new strategy for full-length Ebola virus glycoprotein expression in E.coli. Virologica Sinica, 2016, 31, 500-508.	1,2	1

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127	Investigation of Viral Pathogen Profiles in Some Natural Hosts and Vectors in China. Virologica Sinica, 2018, 33, 1-4.	1.2	1
128	vB_LspM-01: a novel myovirus displaying pseudolysogeny in Lysinibacillus sphaericus C3-41. Applied Microbiology and Biotechnology, 2018, 102, 10691-10702.	1.7	1
129	Network for safe and secure labs. Science, 2018, 362, 267-267.	6.0	1
130	Inaugural editorial: Towards evidence-based biosafety and biosecurity. Journal of Biosafety and Biosecurity, 2019, 1, 1-2.	1.4	1
131	Increased morbidity of obese mice infected with mouse-adapted SARS-CoV-2. Cell Discovery, 2021, 7, 74.	3.1	1
132	In Vitro and In Vivo Characterization of a New Strain of Mosquito Flavivirus Derived from Culicoides. Viruses, 2022, 14, 1298.	1.5	1
133	Evaluation of MICRO-CHEM PLUS as a Disinfectant for Biosafety Level 4 Laboratory in China. Applied Biosafety, 2018, 23, 32-38.	0.2	0
134	Development and Characterization of SYBR Green I Based RT-PCR Assay for Detection of Omsk Hemorrhagic Fever Virus. Virologica Sinica, 2021, , 1.	1.2	0
135	Title is missing!. , 2020, 14, e0008920.		0
136	Title is missing!. , 2020, 14, e0008920.		0
137	Title is missing!. , 2020, 14, e0008920.		0
138	Title is missing!. , 2020, 14, e0008920.		0