

Tetsuya Mizutani

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8411634/publications.pdf>

Version: 2024-02-01

266
papers

7,031
citations

76031

42
h-index

107981

68
g-index

276
all docs

276
docs citations

276
times ranked

9015
citing authors

#	ARTICLE	IF	CITATIONS
1	Characterization of Keterah orthonairovirus and evaluation of therapeutic candidates against Keterah orthonairovirus infectious disease. <i>Ticks and Tick-borne Diseases</i> , 2022, 13, 101834.	1.1	4
2	Validation of the usefulness of 26S rDNA D1/D2, internal transcribed spacer, and intergenic spacer 1 for molecular epidemiological analysis of <i>Macrorhabdus ornithogaster</i> . <i>Journal of Veterinary Medical Science</i> , 2022, 84, 244-250.	0.3	1
3	Anti-malarial activity in a Chinese herbal supplement containing <i>Inonotus obliquus</i> and <i>Panax notoginseng</i> . <i>Parasitology International</i> , 2022, 87, 102532.	0.6	1
4	First isolation and genomic characterization of bovine parechovirus from faecal samples of cattle in Japan. <i>Journal of General Virology</i> , 2022, 103, .	1.3	4
5	Genomic diversity and intragenic recombination of species C rotaviruses. <i>Journal of General Virology</i> , 2022, 103, .	1.3	8
6	Establishment of an experimental model of normal dog bladder organoid using a three-dimensional culture method. <i>Biomedicine and Pharmacotherapy</i> , 2022, 151, 113105.	2.5	10
7	Metagenomic identification, sequencing, and genome analysis of porcine hepe-astroviruses (bastroviruses) in porcine feces in Japan. <i>Infection, Genetics and Evolution</i> , 2021, 88, 104664.	1.0	2
8	Dispersal history of <i>Miniopterus fuliginosus</i> bats and their associated viruses in east Asia. <i>PLoS ONE</i> , 2021, 16, e0244006.	1.1	6
9	Viral-derived DNA invasion and individual variation in an Indonesian population of large flying fox <i>Pteropus vampyrus</i> . <i>Journal of Veterinary Medical Science</i> , 2021, 83, 1068-1074.	0.3	1
10	Two Novel Endornaviruses Co-infecting a Phytophthora Pathogen of <i>Asparagus officinalis</i> Modulate the Developmental Stages and Fungicide Sensitivities of the Host Oomycete. <i>Frontiers in Microbiology</i> , 2021, 12, 633502.	1.5	20
11	Stochastic modelling of the effects of human-mobility restriction and viral infection characteristics on the spread of COVID-19. <i>Scientific Reports</i> , 2021, 11, 6856.	1.6	10
12	Know the facts of COVID-19. <i>Okayama Igakkai Zasshi</i> , 2021, 133, 62-67.	0.0	0
13	Isolation and characterization of mammalian orthoreovirus type 3 from a fecal sample from a wild boar in Japan. <i>Archives of Virology</i> , 2021, 166, 1671-1680.	0.9	6
14	Association between the blaCTX-M-14-harboring <i>Escherichia coli</i> Isolated from Weasels and Domestic Animals Reared on a University Campus. <i>Antibiotics</i> , 2021, 10, 432.	1.5	6
15	Geographic Distribution and Phylogeny of Soricine Shrew-Borne Seewis Virus and Altai Virus in Russia. <i>Viruses</i> , 2021, 13, 1286.	1.5	5
16	Rodent-Borne Orthohantaviruses in Vietnam, Madagascar and Japan. <i>Viruses</i> , 2021, 13, 1343.	1.5	3
17	Natto extract, a Japanese fermented soybean food, directly inhibits viral infections including SARS-CoV-2 in vitro. <i>Biochemical and Biophysical Research Communications</i> , 2021, 570, 21-25.	1.0	19
18	Confirmation of <i>Oryctes rhinoceros nudivirus</i> infections in G-haplotype coconut rhinoceros beetles (<i>Oryctes rhinoceros</i>) from Palauan PCR-positive populations. <i>Scientific Reports</i> , 2021, 11, 18820.	1.6	14

#	ARTICLE	IF	CITATIONS
19	African pygmy hedgehog adenovirus: Virus replication, virus-induced cytopathogenesis and activation of mitogen-activated protein kinase signaling pathways in infected MDCK cells. <i>Research in Veterinary Science</i> , 2021, 139, 152-158.	0.9	0
20	Copper content in ascitic fluid is associated with angiogenesis and progression in ovarian cancer. <i>Journal of Trace Elements in Medicine and Biology</i> , 2021, 68, 126865.	1.5	15
21	Establishment of Intestinal Organoid from <i>Rousettus leschenaultii</i> and the Susceptibility to Bat-Associated Viruses, SARS-CoV-2 and Pteropine Orthoreovirus. <i>International Journal of Molecular Sciences</i> , 2021, 22, 10763.	1.8	14
22	Identification of G-quadruplex sequences in severe acute respiratory syndrome coronavirus 2. <i>Translational and Regulatory Sciences</i> , 2021, 3, 89-92.	0.2	1
23	Development of a novel and rapid measurement system for growth differentiation factor-15, progranulin, and osteopontin in uterine sarcoma. <i>Endocrine Journal</i> , 2020, 67, 91-94.	0.7	5
24	Complete genome sequencing and genetic analysis of a Japanese porcine torovirus strain detected in swine feces. <i>Archives of Virology</i> , 2020, 165, 471-477.	0.9	7
25	Molecular characterization of feline paramyxovirus in Japanese cat populations. <i>Archives of Virology</i> , 2020, 165, 413-418.	0.9	19
26	Development of a one-run real-time PCR detection system for pathogens associated with porcine respiratory diseases. <i>Journal of Veterinary Medical Science</i> , 2020, 82, 217-223.	0.3	30
27	Genetic Diversity and Phylogeography of Thottapalayam thottivirus (Hantaviridae) in Asian House Shrew (<i>Suncus murinus</i>) in Eurasia. <i>Frontiers in Cellular and Infection Microbiology</i> , 2020, 10, 438.	1.8	5
28	Multiple genotypes of enterovirus G carrying a papain-like cysteine protease (PL-CP) sequence circulating on two pig farms in Japan: first identification of enterovirus G10 carrying a PL-CP sequence. <i>Archives of Virology</i> , 2020, 165, 2909-2914.	0.9	3
29	Increased proportion of apoptotic cells in cat kidney tissues infected with feline morbillivirus. <i>Archives of Virology</i> , 2020, 165, 2647-2651.	0.9	0
30	Nutritional status and dietary diversity of school-age children living with HIV: a cross-sectional study in Phnom Penh, Cambodia. <i>BMC Public Health</i> , 2020, 20, 1181.	1.2	12
31	Draft Genome Sequence of Blood-Origin <i>Streptococcus canis</i> Strain FU149, Isolated from a Dog with Necrotizing Soft Tissue Infection. <i>Microbiology Resource Announcements</i> , 2020, 9, .	0.3	3
32	Draft Genome Sequences of Seven <i>Streptococcus canis</i> Strains Isolated from Diseased Companion Animals in Japan. <i>Microbiology Resource Announcements</i> , 2020, 9, .	0.3	6
33	Terminal Genome Sequences of the Soft Tick Bunyavirus. <i>Microbiology Resource Announcements</i> , 2020, 9, .	0.3	2
34	Mosquito-borne viruses, insect-specific flaviviruses (family <i>Flaviviridae</i>, genus <i>Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 16</i>	0.3	10
35	mosquitoes. <i>Journal of Veterinary Medical Science</i> , 2020, 82, 1030-1041. ATeam technology for detecting early signs of viral cytopathic effect. <i>Journal of Veterinary Medical Science</i> , 2020, 82, 387-393.	0.3	2
36	Bacterial and protozoan pathogens/symbionts in ticks infecting wild grasscutters (<i>Thryonomys</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 62	0.9	9

#	ARTICLE	IF	CITATIONS
37	Efficacy of primary liver organoid culture from different stages of non-alcoholic steatohepatitis (NASH) mouse model. <i>Biomaterials</i> , 2020, 237, 119823.	5.7	50
38	BoLA-DRB3 Polymorphism is Associated with Differential Susceptibility to Bovine Leukemia Virus-Induced Lymphoma and Proviral Load. <i>Viruses</i> , 2020, 12, 352.	1.5	51
39	Molecular characterization of full genome sequences of Newcastle disease viruses circulating among vaccinated chickens in Egypt during 2011–2013. <i>Journal of Veterinary Medical Science</i> , 2020, 82, 809-816.	0.3	10
40	Genetic diversity of enterovirus G detected in faecal samples of wild boars in Japan: identification of novel genotypes carrying a papain-like cysteine protease sequence. <i>Journal of General Virology</i> , 2020, 101, 840-852.	1.3	5
41	Comparative Analysis of Fecal Microbiota in Grasscutter (<i>Thryonomys swinderianus</i>) and Other Herbivorous Livestock in Ghana. <i>Microorganisms</i> , 2020, 8, 265.	1.6	5
42	Detection of Viral Infection and Subsequent Apoptosis in Cells by Raman Scattering Microspectroscopy. , 2020, , .		0
43	First identification of Sapoviruses in wild boar. <i>Virus Research</i> , 2019, 271, 197680.	1.1	7
44	Complete genome sequencing and genetic characterization of porcine sapovirus genogroup (G) X and GXI: GVI, GVII, GX, and GXI sapoviruses share common genomic features and form a unique porcine SaV clade. <i>Infection, Genetics and Evolution</i> , 2019, 75, 103959.	1.0	5
45	Establishment of a novel experimental model for muscle-invasive bladder cancer using a dog bladder cancer organoid culture. <i>Cancer Science</i> , 2019, 110, 2806-2821.	1.7	75
46	A novel defective recombinant porcine enterovirus G virus carrying a porcine torovirus papain-like cysteine protease gene and a putative anti-apoptosis gene in place of viral structural protein genes. <i>Infection, Genetics and Evolution</i> , 2019, 75, 103975.	1.0	14
47	END-phenomenon negative bovine viral diarrhea virus that induces the host's innate immune response supports propagation of BVDVs with different immunological properties. <i>Virology</i> , 2019, 538, 97-110.	1.1	6
48	Gut microbiota confers host resistance to obesity by metabolizing dietary polyunsaturated fatty acids. <i>Nature Communications</i> , 2019, 10, 4007.	5.8	231
49	Phylogenetic analysis of novel posaviruses detected in feces of Japanese pigs with posaviruses and posalike viruses of vertebrates and invertebrates. <i>Archives of Virology</i> , 2019, 164, 2147-2151.	0.9	8
50	Encephalomyocarditis virus is potentially derived from eastern bent-wing bats living in East Asian countries. <i>Virus Research</i> , 2019, 259, 62-67.	1.1	6
51	Association of feline morbillivirus infection with defined pathological changes in cat kidney tissues. <i>Veterinary Microbiology</i> , 2019, 228, 12-19.	0.8	24
52	Cloning of carrier cells infected with oncolytic adenovirus driven by <i>midkine</i> promoter and biosafety studies. <i>Journal of Gene Medicine</i> , 2019, 21, e3064.	1.4	5
53	The complete genomic sequence of <i>Rhinolophus gammaherpesvirus 1</i> isolated from a greater horseshoe bat. <i>Archives of Virology</i> , 2019, 164, 317-319.	0.9	5
54	Complete genomic analysis and molecular characterization of Japanese porcine sapeloviruses. <i>Virus Genes</i> , 2019, 55, 198-208.	0.7	9

#	ARTICLE	IF	CITATIONS
55	Complete Genome Sequence of an Adenovirus-1 Isolate from an African Pygmy Hedgehog (<i>Atelerix</i>) Tj ETQq1 1 0.784314 rgBT / Ome 2019, 8, .	0.3	7
56	Draft Genome Sequence of <i>Streptococcus canis</i> Clinical Strain OT1, Isolated from a Dog Owner with Invasive Infection without a Dog Bite in Japan. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.3	6
57	Epidemic nodular facial myxomatous dermatitis in juvenile Cranwellâ€™s horned frogs <i>Ceratophrys cranwelli</i> . <i>Diseases of Aquatic Organisms</i> , 2019, 134, 57-64.	0.5	0
58	Angiotensin II promotes pulmonary metastasis of melanoma through the activation of adhesion molecules in vascular endothelial cells. <i>Biochemical Pharmacology</i> , 2018, 154, 136-147.	2.0	24
59	Characterization of a novel thogotovirus isolated from <i>Amblyomma testudinarium</i> ticks in Ehime, Japan: A significant phylogenetic relationship to Bourbon virus. <i>Virus Research</i> , 2018, 249, 57-65.	1.1	30
60	A new comprehensive method for detection of livestock-related pathogenic viruses using a target enrichment system. <i>Biochemical and Biophysical Research Communications</i> , 2018, 495, 1871-1877.	1.0	7
61	Isolation and characterization of Kabuto Mountain virus, a new tick-borne phlebovirus from <i>Haemaphysalis flava</i> ticks in Japan. <i>Virus Research</i> , 2018, 244, 252-261.	1.1	24
62	Draft Genome Sequence of <i>Streptococcus canis</i> Clinical Strain TA4, Harboring the M-Like Protein Gene and Isolated in Japan from a Patient with Bacteremia. <i>Genome Announcements</i> , 2018, 6, .	0.8	8
63	Isolation of Pteropine orthoreovirus from <i>Pteropus vampyrus</i> in Garut, Indonesia. <i>Virus Genes</i> , 2018, 54, 823-827.	0.7	14
64	Malaria knowledge, preventive actions, and treatment-seeking behavior among ethnic minorities in Ratanakiri Province, Cambodia: a community-based cross-sectional survey. <i>BMC Public Health</i> , 2018, 18, 1206.	1.2	13
65	Metagenomic identification and sequence analysis of a Teschovirus A-related virus in porcine feces in Japan, 2014â€“2016. <i>Infection, Genetics and Evolution</i> , 2018, 66, 210-216.	1.0	10
66	Dembo polymerase chain reaction technique for detection of bovine abortion, diarrhea, and respiratory disease complex infectious agents in potential vectors and reservoirs. <i>Journal of Veterinary Science</i> , 2018, 19, 350.	0.5	5
67	Whole genome analysis of a novel picornavirus related to the Enterovirus/Sapelovirus supergroup from porcine feces in Japan. <i>Virus Research</i> , 2018, 257, 68-73.	1.1	3
68	Draft Genome Sequence of <i>Mycobacterium montefiorensis</i> Isolated from Japanese Black Salamander (<i>Hynobius nigrescens</i>). <i>Genome Announcements</i> , 2018, 6, .	0.8	5
69	Postnatal care could be the key to improving the continuum of care in maternal and child health in Ratanakiri, Cambodia. <i>PLoS ONE</i> , 2018, 13, e0198829.	1.1	47
70	Characteristics of Staphylococcal Enterotoxin A Production and Growth of <i>Staphylococcus aureus</i> in Shaking and Stationary Cultures. <i>Biocontrol Science</i> , 2018, 23, 207-214.	0.2	3
71	Luteinizing Hormone Facilitates Antral Follicular Maturation and Survival via Thecal Paracrine Signaling in Cattle. <i>Endocrinology</i> , 2018, 159, 2337-2347.	1.4	27
72	Molecular diversity of the faecal microbiota of Toy Poodles in Japan. <i>Journal of Veterinary Medical Science</i> , 2018, 80, 749-754.	0.3	23

#	ARTICLE	IF	CITATIONS
73	Application of the SureSelect target enrichment system for next-generation sequencing to obtain the complete genome sequence of bovine leukemia virus. <i>Archives of Virology</i> , 2018, 163, 3155-3159.	0.9	7
74	Whole genome analysis of a novel neurotropic bovine astrovirus detected in a Japanese black steer with non-suppurative encephalomyelitis in Japan. <i>Archives of Virology</i> , 2018, 163, 2805-2810.	0.9	15
75	Genetic diversity and recombination of enterovirus G strains in Japanese pigs: High prevalence of strains carrying a papain-like cysteine protease sequence in the enterovirus G population. <i>PLoS ONE</i> , 2018, 13, e0190819.	1.1	30
76	Barriers for pregnant women living in rural, agricultural villages to accessing antenatal care in Cambodia: A community-based cross-sectional study combined with a geographic information system. <i>PLoS ONE</i> , 2018, 13, e0194103.	1.1	29
77	An ascovirus isolated from <i>Spodoptera litura</i> (Noctuidae: Lepidoptera) transmitted by the generalist endoparasitoid <i>Meteorus pulchricornis</i> (Braconidae: Hymenoptera). <i>Journal of General Virology</i> , 2018, 99, 574-584.	1.3	12
78	Decreased expression of the immediate early protein, ICP4, by deletion of the tegument protein VP22 of equine herpesvirus type 1. <i>Journal of Veterinary Medical Science</i> , 2018, 80, 311-315.	0.3	2
79	Angiotensin II increase the pulmonary metastasis through the vascular endothelial cell adherent pathway in hematogenous metastasis of melanoma cells. <i>Proceedings for Annual Meeting of the Japanese Pharmacological Society</i> , 2018, WCP2018, PO1-8-11.	0.0	0
80	Diversity in VP3, NSP3, and NSP4 of rotavirus B detected from Japanese cattle. <i>Infection, Genetics and Evolution</i> , 2017, 49, 97-103.	1.0	10
81	Whole genome analysis of porcine astroviruses detected in Japanese pigs reveals genetic diversity and possible intra-genotypic recombination. <i>Infection, Genetics and Evolution</i> , 2017, 50, 38-48.	1.0	30
82	First isolation and characterization of pteropine orthoreoviruses in fruit bats in the Philippines. <i>Archives of Virology</i> , 2017, 162, 1529-1539.	0.9	26
83	Characterization of cross-clade monoclonal antibodies against H5N1 highly pathogenic avian influenza virus and their application to the antigenic analysis of diverse H5 subtype viruses. <i>Archives of Virology</i> , 2017, 162, 2257-2269.	0.9	6
84	Complete genome analysis of porcine kobuviruses from the feces of pigs in Japan. <i>Virus Genes</i> , 2017, 53, 593-602.	0.7	8
85	Characterization of avian paramyxovirus serotype 14, a novel serotype, isolated from a duck fecal sample in Japan. <i>Virus Research</i> , 2017, 228, 46-57.	1.1	34
86	<i>Mycobacterium avium</i> subsp. <i>hominissuis</i> meningoencephalitis in a cat. <i>Veterinary Microbiology</i> , 2017, 204, 43-45.	0.8	7
87	Genetic diversity and intergenogroup recombination events of sapoviruses detected from feces of pigs in Japan. <i>Infection, Genetics and Evolution</i> , 2017, 55, 209-217.	1.0	20
88	Isolation and characterization of Tatumizu tick virus: A new coltivirus from <i>Haemaphysalis flava</i> ticks in Japan. <i>Virus Research</i> , 2017, 242, 131-140.	1.1	34
89	Complete Genome Sequence of <i>Mycobacterium stephanolepidis</i> . <i>Genome Announcements</i> , 2017, 5, .	0.8	8
90	Generation of a novel live rabies vaccine strain with a high level of safety by introducing attenuating mutations in the nucleoprotein and glycoprotein. <i>Vaccine</i> , 2017, 35, 5622-5628.	1.7	8

#	ARTICLE	IF	CITATIONS
91	Molecular characteristics and prevalence of small ruminant lentiviruses in goats in Japan. Archives of Virology, 2017, 162, 3007-3015.	0.9	5
92	Identification of a novel bovine enterovirus possessing highly divergent amino acid sequences in capsid protein. BMC Microbiology, 2017, 17, 18.	1.3	17
93	The full genome sequences of 8 equine herpesvirus type 4 isolates from horses in Japan. Journal of Veterinary Medical Science, 2017, 79, 206-212.	0.3	6
94	Development of a one-run real-time PCR detection system for pathogens associated with bovine respiratory disease complex. Journal of Veterinary Medical Science, 2017, 79, 517-523.	0.3	70
95	Discovery of fur seal feces-associated circular DNA virus in swine feces in Japan. Journal of Veterinary Medical Science, 2017, 79, 1664-1666.	0.3	5
96	Isolation and characterization of a new iflavirus from Armigeres spp. mosquitoes in the Philippines. Journal of General Virology, 2017, 98, 2876-2881.	1.3	16
97	The Window of Implantation Is Closed by Estrogen via Insulin-Like Growth Factor 1 Pathway. Journal of Reproduction and Infertility, 2017, 18, 231-241.	1.0	7
98	Detection of Japanese Eel Endothelial Cells-infecting Virus (JEECV) in Mature Japanese Eel &i>&i>Anguilla japonica&i>&i>; Caught from Their Spawning Area. Fish Pathology, 2016, 51, 64-66.	0.4	5
99	Involvement of the 3' UTR in Encapsidation of the Hepatitis C Virus. PLoS Pathogens, 2016, 12, e1005441.	2.1	21
100	Genome Sequences of Rotavirus A Strains Ty-1 and Ty-3, Isolated from Turkeys in Ireland in 1979. Genome Announcements, 2016, 4, .	0.8	13
101	Genome sequence of a novel mitovirus identified in the phytopathogenic fungus Alternaria arborescens. Archives of Virology, 2016, 161, 2627-2631.	0.9	21
102	Whole genome sequences of Japanese porcine species C rotaviruses reveal a high diversity of genotypes of individual genes and will contribute to a comprehensive, generally accepted classification system. Infection, Genetics and Evolution, 2016, 44, 106-113.	1.0	17
103	Detection of skunk adenovirus 1 (SkAdV-1) in an African pigmy hedgehog (Atelerix albiventris). Veterinary Record Case Reports, 2016, 4, e000321.	0.1	10
104	Characterization and phylogenetic analysis of a novel picornavirus from swine feces in Japan. Archives of Virology, 2016, 161, 1685-1690.	0.9	9
105	Genome sequence of a novel victorivirus identified in the phytopathogenic fungus Alternaria arborescens. Archives of Virology, 2016, 161, 1701-1704.	0.9	25
106	Development of a novel detection system for microbes from bovine diarrhea by real-time PCR. Journal of Veterinary Medical Science, 2016, 78, 383-389.	0.3	26
107	Detection of Japanese eel endothelial cells-infecting virus in &i>&i>Anguilla japonica&i>&i>; elvers. Journal of Veterinary Medical Science, 2016, 78, 705-707.	0.3	13
108	Detection of &i>&i>Campylobacter jejuni&i>&i>; in rectal swab samples from &i>&i>Rousettus amplexicaudatus&i>&i>; in the Philippines. Journal of Veterinary Medical Science, 2016, 78, 1347-1350.	0.3	13

#	ARTICLE	IF	CITATIONS
109	Detection of novel kobu-like viruses in Japanese black cattle in Japan. <i>Journal of Veterinary Medical Science</i> , 2016, 78, 321-324.	0.3	12
110	A novel Bunyavirus from the soft tick, <i>Argas vespertilionis</i> , in Japan. <i>Journal of Veterinary Medical Science</i> , 2016, 78, 443-445.	0.3	26
111	Molecular epidemiological study of adenovirus infecting western lowland gorillas and humans in and around Moukalaba-Doudou National Park (Gabon). <i>Virus Genes</i> , 2016, 52, 671-678.	0.7	8
112	Identification of further diversity among posaviruses. <i>Archives of Virology</i> , 2016, 161, 3541-3548.	0.9	12
113	Complete Genome Sequencing and Phylogenetic Analysis of a Getah Virus Strain (Genus Alphavirus,) Tj ETQq1 1 0.784314 rgBT /Over Vector-Borne and Zoonotic Diseases, 2016, 16, 769-776.	0.6	10
114	Activation of c-Jun N-terminal kinase by Akabane virus is required for apoptosis. <i>Research in Veterinary Science</i> , 2016, 107, 147-151.	0.9	11
115	Complete Genome Sequencing of Bovine Viral Diarrhea Virus 1, Subgenotypes 1n and 1o. <i>Genome Announcements</i> , 2016, 4, .	0.8	12
116	Whole genome analysis of Japanese bovine toroviruses reveals natural recombination between porcine and bovine toroviruses. <i>Infection, Genetics and Evolution</i> , 2016, 38, 90-95.	1.0	13
117	Isolation of a sp. nov. Ljungan virus from wild birds in Japan. <i>Journal of General Virology</i> , 2016, 97, 1818-1822.	1.3	6
118	Comprehensive pathogen detection associated with four recurrent episodes of Kawasaki disease in a patient during a single year using next-generation sequencing. <i>JMM Case Reports</i> , 2016, 3, e005019.	1.3	11
119	Characterization of Venom and Oviduct Components of Parasitoid Wasp <i>Asobara japonica</i> . <i>PLoS ONE</i> , 2016, 11, e0160210.	1.1	8
120	Development of one-step real-time reverse transcriptase-PCR-based assays for the rapid and simultaneous detection of four viruses causing porcine diarrhea. <i>Japanese Journal of Veterinary Research</i> , 2016, 64, 5-14.	0.7	16
121	Use of S1 nuclease in deep sequencing for detection of double-stranded RNA viruses. <i>Journal of Veterinary Medical Science</i> , 2015, 77, 1163-1166.	0.3	5
122	Genetic and antigenic analysis of <i>Chlamydia pecorum</i> strains isolated from calves with diarrhea. <i>Journal of Veterinary Medical Science</i> , 2015, 77, 777-782.	0.3	2
123	Transcriptional regulation of genes related to progesterone production [Review]. <i>Endocrine Journal</i> , 2015, 62, 757-763.	0.7	28
124	Quantitative PCR detection of feline morbillivirus in cat urine samples. <i>Journal of Veterinary Medical Science</i> , 2015, 77, 1701-1703.	0.3	16
125	Emergence of infectious malignant thrombocytopenia in Japanese macaques (<i>Macaca fuscata</i>) by SRV-4 after transmission to a novel host. <i>Scientific Reports</i> , 2015, 5, 8850.	1.6	14
126	Ovarian mucinous adenocarcinoma with functioning stroma in postmenopausal women: aromatase and SF-1 expressions. <i>Journal of Ovarian Research</i> , 2015, 8, 73.	1.3	8

#	ARTICLE	IF	CITATIONS
127	Complete Genome Sequences of Two Japanese Eel Endothelial Cell-Infecting Virus Strains Isolated in Japan. <i>Genome Announcements</i> , 2015, 3, .	0.8	4
128	Two types of genetic carrier, the <i>IncP</i> genomic island and the novel <i>IncP</i> plasmid, for the <i>aac(2)</i> gene that confers kasugamycin resistance in <i>Acidovorax avenae</i> ssp. <i>avenae</i> . <i>Molecular Plant Pathology</i> , 2015, 16, 288-300.	2.0	10
129	Characterization of Self-Assembled Virus-Like Particles of Merkel Cell Polyomavirus. <i>PLoS ONE</i> , 2015, 10, e0115646.	1.1	20
130	Complete Genome Sequence of Bovine Viral Diarrhea Virus 2 Japanese Reference and Vaccine Strain KZ-91CP. <i>Genome Announcements</i> , 2015, 3, .	0.8	2
131	H2 genotypes of G4P[6], G5P[7], and G9[23] porcine rotaviruses show super-short RNA electropherotypes. <i>Veterinary Microbiology</i> , 2015, 176, 250-256.	0.8	17
132	Establishment and characterization of a novel orthotopic mouse model for human uterine sarcoma with different metastatic potentials. <i>Cancer Letters</i> , 2015, 366, 182-190.	3.2	7
133	Full genome analysis of bovine astrovirus from fecal samples of cattle in Japan: identification of possible interspecies transmission of bovine astrovirus. <i>Archives of Virology</i> , 2015, 160, 2491-2501.	0.9	65
134	Isolation and characterization of a novel Rhabdovirus from a wild boar (<i>Sus scrofa</i>) in Japan. <i>Veterinary Microbiology</i> , 2015, 179, 197-203.	0.8	5
135	Identification of novel steroidogenic factor 1 (SF-1)-target genes and components of the SF-1 nuclear complex. <i>Molecular and Cellular Endocrinology</i> , 2015, 408, 133-137.	1.6	7
136	Whole-genome sequence analysis of G3 and G14 equine group A rotaviruses isolated in the late 1990s and 2009-2010. <i>Archives of Virology</i> , 2015, 160, 1171-1179.	0.9	11
137	Identification, characterization and full-length sequence analysis of a novel dsRNA virus isolated from the arboreal ant <i>Camponotus yamaokai</i> . <i>Journal of General Virology</i> , 2015, 96, 1930-1937.	1.3	43
138	Parrot bornavirus-2 and -4 RNA detected in wild bird samples in Japan are phylogenetically adjacent to those found in pet birds in Japan. <i>Virus Genes</i> , 2015, 51, 234-243.	0.7	6
139	Detection of a novel herpesvirus from bats in the Philippines. <i>Virus Genes</i> , 2015, 51, 136-139.	0.7	19
140	Identification and complete genome analysis of a novel bovine picornavirus in Japan. <i>Virus Research</i> , 2015, 210, 205-212.	1.1	20
141	Design of Goat Farming Support System Adapted for Small and Remote Islands using ICT, And its Application on Branding of Agricultural Products. <i>IEEJ Transactions on Industry Applications</i> , 2015, 135, 107-112.	0.1	0
142	Imported Case of Acute Respiratory Tract Infection Associated with a Member of Species Nelson Bay Orthoreovirus. <i>PLoS ONE</i> , 2014, 9, e92777.	1.1	44
143	Characterization of a genetic and antigenic variant of avian paramyxovirus 6 isolated from a migratory wild bird, the red-necked stint (<i>Calidris ruficollis</i>). <i>Archives of Virology</i> , 2014, 159, 3101-3105.	0.9	13
144	Molecular, biological, and antigenic characterization of a Border disease virus isolated from a pig during classical swine fever surveillance in Japan. <i>Journal of Veterinary Diagnostic Investigation</i> , 2014, 26, 547-552.	0.5	14

#	ARTICLE	IF	CITATIONS
145	Detection of a pneumonia virus of mice (PVM) in an African hedgehog (<i>Atelerix arbiventris</i>) with suspected wobbly hedgehog syndrome (WHS). <i>Veterinary Microbiology</i> , 2014, 173, 136-140.	0.8	9
146	Existence of feline morbillivirus infection in Japanese cat populations. <i>Archives of Virology</i> , 2014, 159, 371-373.	0.9	57
147	Transcriptional regulation of human ferredoxin reductase through an intronic enhancer in steroidogenic cells. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2014, 1839, 33-42.	0.9	21
148	The First Identification and Retrospective Study of Severe Fever With Thrombocytopenia Syndrome in Japan. <i>Journal of Infectious Diseases</i> , 2014, 209, 816-827.	1.9	672
149	Detection of enterovirus genome sequence from diarrheal feces of goat. <i>Virus Genes</i> , 2014, 48, 550-552.	0.7	16
150	Identification of a natural recombination in the F and H genes of feline morbillivirus. <i>Virology</i> , 2014, 468-470, 524-531.	1.1	32
151	Identification of novel bovine group A rotavirus G15P[14] strain from epizootic diarrhea of adult cows by de novo sequencing using a next-generation sequencer. <i>Veterinary Microbiology</i> , 2014, 171, 66-73.	0.8	35
152	A Stability Analysis of Mathematical Model on Caprine Arthritis Encephalitis Virus Transmission. <i>Proceedings of the ISCTE International Symposium on Stochastic Systems Theory and Its Applications</i> , 2014, 2014, 265-271.	0.1	1
153	Pregnancy-associated plasma protein (PAPP)-A expressed in the mammary gland controls epithelial cell proliferation and differentiation. <i>Endocrine</i> , 2013, 43, 387-393.	1.1	4
154	Characterization of Dak Nong virus, an insect nidovirus isolated from <i>Culex</i> mosquitoes in Vietnam. <i>Archives of Virology</i> , 2013, 158, 2273-2284.	0.9	29
155	Molecular epidemiology of avian bornavirus from pet birds in Japan. <i>Virus Genes</i> , 2013, 47, 173-177.	0.7	17
156	Lethal Canine Distemper Virus Outbreak in <i>Cynomolgus</i> Monkeys in Japan in 2008. <i>Journal of Virology</i> , 2013, 87, 1105-1114.	1.5	112
157	Canine Distemper Virus Associated with a Lethal Outbreak in Monkeys Can Readily Adapt To Use Human Receptors. <i>Journal of Virology</i> , 2013, 87, 7170-7175.	1.5	60
158	Degenerate polymerase chain reaction strategy with <scp>DNA</scp> microarray for detection of multiple and various subtypes of virus during blood screening. <i>Transfusion</i> , 2013, 53, 2545-2555.	0.8	3
159	Detection of Bovine Group A Rotavirus Using Rapid Antigen Detection Kits, RT-PCR and Next-Generation DNA Sequencing. <i>Journal of Veterinary Medical Science</i> , 2013, 75, 1651-1655.	0.3	10
160	Genomic and serological detection of bat coronavirus from bats in the Philippines. <i>Archives of Virology</i> , 2012, 157, 2349-2355.	0.9	24
161	Analysis of the humoral immune responses among <i>cynomolgus</i> macaque naturally infected with Reston virus during the 1996 outbreak in the Philippines. <i>BMC Veterinary Research</i> , 2012, 8, 189.	0.7	6
162	A seroepidemiologic study of Reston ebolavirus in swine in the Philippines. <i>BMC Veterinary Research</i> , 2012, 8, 82.	0.7	20

#	ARTICLE	IF	CITATIONS
163	Antigen-capture ELISA for the detection of Rift Valley fever virus nucleoprotein using new monoclonal antibodies. <i>Journal of Virological Methods</i> , 2012, 180, 68-74.	1.0	32
164	Detection of bat coronaviruses from <i>Miniopterus fuliginosus</i> in Japan. <i>Virus Genes</i> , 2012, 44, 40-44.	0.7	23
165	Epidemic Myalgia in Adults Associated with Human Parechovirus Type 3 Infection, Yamagata, Japan, 2008. <i>Emerging Infectious Diseases</i> , 2012, 18, 1787-1793.	2.0	65
166	Pathogenic characterization of a cervical lymph node derived from a patient with Kawasaki disease. <i>International Journal of Clinical and Experimental Pathology</i> , 2012, 5, 814-23.	0.5	12
167	Identification and molecular characterization of a new nonsegmented double-stranded RNA virus isolated from <i>Culex</i> mosquitoes in Japan. <i>Virus Research</i> , 2011, 155, 147-155.	1.1	62
168	Novel DNA virus isolated from samples showing endothelial cell necrosis in the Japanese eel, <i>Anguilla japonica</i> . <i>Virology</i> , 2011, 412, 179-187.	1.1	45
169	A novel sapelovirus-like virus isolation from wild boar. <i>Virus Genes</i> , 2011, 43, 243-248.	0.7	21
170	Reston Ebolavirus Antibodies in Bats, the Philippines. <i>Emerging Infectious Diseases</i> , 2011, 17, 1559-60.	2.0	85
171	Viruses Isolated from Bats and Their Importance as Emerging Infectious Diseases. <i>Journal of Veterinary Epidemiology</i> , 2011, 15, 88-93.	0.2	0
172	West Nile Virus : Understanding its Past, Present, and Future. <i>Journal of Disaster Research</i> , 2011, 6, 413-420.	0.4	0
173	Bat Coronaviruses and Experimental Infection of Bats, the Philippines. <i>Emerging Infectious Diseases</i> , 2010, 16, 1217-1223.	2.0	177
174	Identification of a Novel Distal Control Region Upstream of the Human Steroidogenic Acute Regulatory Protein (StAR) Gene That Participates in SF-1-dependent Chromatin Architecture. <i>Journal of Biological Chemistry</i> , 2010, 285, 28240-28251.	1.6	18
175	Novel Betaherpesvirus in Bats. <i>Emerging Infectious Diseases</i> , 2010, 16, 986-988.	2.0	35
176	Sequence and phylogenetic analyses of Saffold cardiovirus from children with exudative tonsillitis in Yamagata, Japan. <i>Scandinavian Journal of Infectious Diseases</i> , 2010, 42, 950-952.	1.5	33
177	Signaling Pathways of SARS-CoV In Vitro and In Vivo. , 2010, , 305-322.		6
178	Development of a Method to Detect Viral RNA Sequences From Cultured Cells by Combining Size Fraction and a Rapid Determination System for Viral RNA Sequences (RDV). <i>Journal of Veterinary Science & Technology</i> , 2010, 01, .	0.3	1
179	Sequencing and phylogenetic analyses of Saffold cardiovirus (SAFV) genotype 3 isolates from children with upper respiratory infection in Gunma, Japan. <i>Japanese Journal of Infectious Diseases</i> , 2010, 63, 378-80.	0.5	7
180	Sequencing and Phylogenetic Analyses of Saffold Cardiovirus (SAFV) Genotype 3 Isolates from Children with Upper Respiratory Infection in Gunma, Japan. <i>Japanese Journal of Infectious Diseases</i> , 2010, 63, 378-380.	0.5	19

#	ARTICLE	IF	CITATIONS
181	Characterization of Monoclonal Antibodies to Junin Virus Nucleocapsid Protein and Application to the Diagnosis of Hemorrhagic Fever Caused by South American Arenaviruses. <i>Vaccine Journal</i> , 2009, 16, 1132-1138.	3.2	17
182	Virulence and pathophysiology of the Congo Basin and West African strains of monkeypox virus in non-human primates. <i>Journal of General Virology</i> , 2009, 90, 2266-2271.	1.3	86
183	Direct Metagenomic Detection of Viral Pathogens in Nasal and Fecal Specimens Using an Unbiased High-Throughput Sequencing Approach. <i>PLoS ONE</i> , 2009, 4, e4219.	1.1	240
184	Novel reovirus isolation from an Ostrich (<i>Struthio camelus</i>) in Japan. <i>Veterinary Microbiology</i> , 2009, 134, 227-232.	0.8	12
185	Loop-mediated isothermal amplification-based diagnostic assay for monkeypox virus infections. <i>Journal of Medical Virology</i> , 2009, 81, 1102-1108.	2.5	59
186	Novel virus discovery in field-collected mosquito larvae using an improved system for rapid determination of viral RNA sequences (RDV ver4.0). <i>Archives of Virology</i> , 2009, 154, 153-158.	0.9	35
187	Detection of a new bat gammaherpesvirus in the Philippines. <i>Virus Genes</i> , 2009, 39, 90-93.	0.7	30
188	Ligation-mediated amplification for effective rapid determination of viral RNA sequences (RDV). <i>Journal of Clinical Virology</i> , 2008, 43, 56-59.	1.6	14
189	Application of quantitative gene expression analysis for pertussis vaccine safety control. <i>Vaccine</i> , 2008, 26, 4686-4696.	1.7	16
190	Isolation of Novel Adenovirus from Fruit Bat (<i>Pteropus dasymallus yayeyamae</i>). <i>Emerging Infectious Diseases</i> , 2008, 14, 347-349.	2.0	77
191	Diagnosis and assessment of monkeypox virus (MPXV) infection by quantitative PCR assay: differentiation of Congo Basin and West African MPXV strains. <i>Japanese Journal of Infectious Diseases</i> , 2008, 61, 140-2.	0.5	12
192	Development of Recombinant Nucleoprotein-Based Diagnostic Systems for Lassa Fever. <i>Vaccine Journal</i> , 2007, 14, 1182-1189.	3.2	40
193	Amino Acid Substitutions in the S2 Region Enhance Severe Acute Respiratory Syndrome Coronavirus Infectivity in Rat Angiotensin-Converting Enzyme 2-Expressing Cells. <i>Journal of Virology</i> , 2007, 81, 10831-10834.	1.5	13
194	Rapid Genome Sequencing of RNA Viruses. <i>Emerging Infectious Diseases</i> , 2007, 13, 322-324.	2.0	41
195	Synthesis and biological evaluation of nucleoside analogues having 6-chloropurine as anti-SARS-CoV agents. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2007, 17, 2470-2473.	1.0	25
196	Rapid determination of viral RNA sequences in mosquitoes collected in the field. <i>Journal of Virological Methods</i> , 2007, 146, 372-374.	1.0	33
197	Regulation of p90RSK phosphorylation by SARS-CoV infection in Vero E6 cells. <i>FEBS Letters</i> , 2006, 580, 1417-1424.	1.3	17
198	Mechanisms of establishment of persistent SARS-CoV-infected cells. <i>Biochemical and Biophysical Research Communications</i> , 2006, 347, 261-265.	1.0	31

#	ARTICLE	IF	CITATIONS
199	Inhibition of cell proliferation by SARS-CoV infection in Vero E6 cells. <i>FEMS Immunology and Medical Microbiology</i> , 2006, 46, 236-243.	2.7	23
200	Induction of protective immunity against severe acute respiratory syndrome coronavirus (SARS-CoV) infection using highly attenuated recombinant vaccinia virus DIs. <i>Virology</i> , 2006, 351, 368-380.	1.1	18
201	Development of an enzyme-linked immunosorbent assay for serological diagnosis of tick-borne encephalitis using subviral particles. <i>Journal of Virological Methods</i> , 2006, 134, 55-60.	1.0	16
202	Evaluation of a novel vesicular stomatitis virus pseudotype-based assay for detection of neutralizing antibody responses to SARS-CoV. <i>Journal of Medical Virology</i> , 2006, 78, 1509-1512.	2.5	39
203	LC16m8, a Highly Attenuated Vaccinia Virus Vaccine Lacking Expression of the Membrane Protein B5R, Protects Monkeys from Monkeypox. <i>Journal of Virology</i> , 2006, 80, 5179-5188.	1.5	101
204	Pseudotyped Vesicular Stomatitis Virus for Functional Analysis of Sars Coronavirus Spike Protein. <i>Advances in Experimental Medicine and Biology</i> , 2006, 581, 293-296.	0.8	2
205	Characterization of Persistent SARS-CoV Infection in Vero E6 Cells. <i>Advances in Experimental Medicine and Biology</i> , 2006, 581, 323-326.	0.8	5
206	Highly Attenuated Vaccinia Virus Dis as a Potential Sars Vaccine. <i>Advances in Experimental Medicine and Biology</i> , 2006, 581, 593-596.	0.8	4
207	Marburgvirus nucleoprotein-capture enzyme-linked immunosorbent assay using monoclonal antibodies to recombinant nucleoprotein: detection of authentic Marburgvirus. <i>Japanese Journal of Infectious Diseases</i> , 2006, 59, 323-5.	0.5	12
208	Species-independent detection of RNA virus by representational difference analysis using non-ribosomal hexanucleotides for reverse transcription. <i>Nucleic Acids Research</i> , 2005, 33, e65-e65.	6.5	105
209	Inhibitory effect of mizoribine and ribavirin on the replication of severe acute respiratory syndrome (SARS)-associated coronavirus. <i>Antiviral Research</i> , 2005, 66, 159-163.	1.9	73
210	Recombinant nucleocapsid protein-based IgG enzyme-linked immunosorbent assay for the serological diagnosis of SARS. <i>Journal of Virological Methods</i> , 2005, 125, 181-186.	1.0	25
211	Mutations in the nonstructural region 5B of hepatitis C virus genotype 1b: Their relation to viral load, response to interferon, and the nonstructural region 5A. <i>Journal of Medical Virology</i> , 2005, 75, 504-512.	2.5	20
212	Vesicular stomatitis virus pseudotyped with severe acute respiratory syndrome coronavirus spike protein. <i>Journal of General Virology</i> , 2005, 86, 2269-2274.	1.3	133
213	JNK and PI3k/Akt signaling pathways are required for establishing persistent SARS-CoV infection in Vero E6 cells. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2005, 1741, 4-10.	1.8	90
214	Role of the N-linked glycans of the prM and E envelope proteins in tick-borne encephalitis virus particle secretion. <i>Vaccine</i> , 2005, 23, 3043-3052.	1.7	59
215	Single point mutation in tick-borne encephalitis virus prM protein induces a reduction of virus particle secretion. <i>Journal of General Virology</i> , 2004, 85, 3049-3058.	1.3	45
216	Amino acid changes responsible for attenuation of virus neurovirulence in an infectious cDNA clone of the Oshima strain of Tick-borne encephalitis virus. <i>Journal of General Virology</i> , 2004, 85, 1007-1018.	1.3	42

#	ARTICLE	IF	CITATIONS
217	Importance of Akt signaling pathway for apoptosis in SARS-CoV-infected Vero E6 cells. <i>Virology</i> , 2004, 327, 169-174.	1.1	68
218	Different chemokine expression in lethal and non-lethal murine west nile virus infection. <i>Journal of Medical Virology</i> , 2004, 74, 507-513.	2.5	78
219	The fluctuations of viral load and serum alanine aminotransferase levels in chronic hepatitis C. <i>Hepatology Research</i> , 2004, 30, 11-17.	1.8	11
220	Epizootiological and Epidemiological Study of Hantavirus Infection in Japan. <i>Microbiology and Immunology</i> , 2004, 48, 843-851.	0.7	17
221	Tyrosine dephosphorylation of STAT3 in SARS coronavirus-infected Vero E6 cells. <i>FEBS Letters</i> , 2004, 577, 187-192.	1.3	65
222	Phosphorylation of p38 MAPK and its downstream targets in SARS coronavirus-infected cells. <i>Biochemical and Biophysical Research Communications</i> , 2004, 319, 1228-1234.	1.0	141
223	Genetic and antigenic characterization of the Amur virus associated with hemorrhagic fever with renal syndrome. <i>Virus Research</i> , 2004, 101, 127-134.	1.1	28
224	Identification of 2,3,7,8-Tetrachlorodibenzo-p-dioxin (TCDD)-inducible and -suppressive Genes in the Rat Placenta: Induction of Interferon-regulated Genes with Possible Inhibitory Roles for Angiogenesis in the Placenta. <i>Endocrine Journal</i> , 2004, 51, 569-577.	0.7	16
225	Enzyme-linked immunosorbent assay using recombinant antigens expressed in mammalian cells for serodiagnosis of tick-borne encephalitis. <i>Journal of Virological Methods</i> , 2003, 108, 171-179.	1.0	24
226	A BHK-21 cell culture-adapted tick-borne encephalitis virus mutant is attenuated for neuroinvasiveness. <i>Vaccine</i> , 2003, 21, 4043-4051.	1.7	51
227	Discrimination of West Nile Virus and Japanese Encephalitis Virus Strains Using RT-PCR RFLP Analysis. <i>Microbiology and Immunology</i> , 2003, 47, 439-445.	0.7	13
228	Development of an Efficient Method for Recovery of Puumala and Puumala-Related Viruses by Inoculation of Mongolian Gerbils. <i>Journal of Veterinary Medical Science</i> , 2003, 65, 1189-1194.	0.3	6
229	Murine Coronavirus Replication-Induced p38 Mitogen-Activated Protein Kinase Activation Promotes Interleukin-6 Production and Virus Replication in Cultured Cells. <i>Journal of Virology</i> , 2002, 76, 5937-5948.	1.5	106
230	Genetic Characterization of Hantaviruses Transmitted by the Korean Field Mouse (<i>Apodemus</i>) Tj ETQq0 0 0 rgBT/Overlock, 10 Tf 50 2	2.0	25
231	Genetic and biological comparison of tick-borne encephalitis viruses from Hokkaido and far-eastern Russia. <i>Japanese Journal of Veterinary Research</i> , 2002, 49, 297-307.	0.7	9
232	Evaluation of European tick-borne encephalitis virus vaccine against recent Siberian and far-eastern subtype strains. <i>Vaccine</i> , 2001, 19, 4774-4779.	1.7	47
233	Expression of apoptosis on rat liver by hepatic vagus hyperactivity after ventromedial hypothalamic lesioning. <i>American Journal of Physiology - Renal Physiology</i> , 2001, 280, G958-G967.	1.6	13
234	Characterization of in vitro and in vivo Antiviral Activity of Lactoferrin and Ribavirin upon Hantavirus.. <i>Journal of Veterinary Medical Science</i> , 2001, 63, 637-645.	0.3	43

#	ARTICLE	IF	CITATIONS
235	Distribution and characterization of tick-borne encephalitis viruses from Siberia and far-eastern Asia. <i>Journal of General Virology</i> , 2001, 82, 1319-1328.	1.3	84
236	The Mechanism of Actinomycin D-Mediated Increase of Borna Disease Virus (BDV) RNA in Cells Persistently Infected by BDV. <i>Microbiology and Immunology</i> , 2000, 44, 597-603.	0.7	6
237	Characterization of Monoclonal Antibodies against Hokkaido Strain Tick-Borne Encephalitis Virus. <i>Microbiology and Immunology</i> , 2000, 44, 533-536.	0.7	14
238	Detection of Hantaviral Antibodies among Patients with Hepatitis of Unknown Etiology in Japan. <i>Microbiology and Immunology</i> , 2000, 44, 357-362.	0.7	18
239	Nascent Synthesis of Leader Sequence-Containing Subgenomic mRNAs in Coronavirus Genome-Length Replicative Intermediate RNA. <i>Virology</i> , 2000, 275, 238-243.	1.1	16
240	Pathogenicity of tick-borne encephalitis virus isolated in Hokkaido, Japan in mouse model. <i>Vaccine</i> , 1999, 17, 779-787.	1.7	45
241	Protection against tick-borne encephalitis virus isolated in Japan by active and passive immunization. <i>Vaccine</i> , 1999, 17, 1532-1539.	1.7	55
242	A Serosurvey of Borna Disease Virus Infection in Wild Rats by a Capture ELISA. <i>Journal of Veterinary Medical Science</i> , 1999, 61, 113-117.	0.3	19
243	Borna Disease Virus Infection in Domestic Cats: Evaluation by RNA and Antibody Detection. <i>Journal of Veterinary Medical Science</i> , 1999, 61, 1167-1170.	0.3	27
244	Reverse Transcription Nested Polymerase Chain Reaction for Detecting p40 RNA of Borna Disease Virus, without Risk of Plasmid Contamination.. <i>Journal of Veterinary Medical Science</i> , 1999, 61, 77-80.	0.3	4
245	Phylogenetic and virulence analysis of tick-borne encephalitis viruses from Japan and far-eastern Russia. <i>Journal of General Virology</i> , 1999, 80, 3127-3135.	1.3	65
246	Characterization of the progressive sublines derived from a weakly malignant cloned cell line, ER-1, co-inoculated subcutaneously with a foreign body. <i>Clinical and Experimental Metastasis</i> , 1998, 16, 291-298.	1.7	4
247	Detection of negative-stranded hepatitis C virus RNA using a novel strand-specific reverse transcription-polymerase chain reaction. <i>Virus Research</i> , 1998, 53, 209-214.	1.1	9
248	Human hepatocyte clonal cell lines that support persistent replication of hepatitis C virus. <i>Virus Research</i> , 1998, 56, 157-167.	1.1	99
249	Single-Step Reverse Transcription-Polymerase Chain Reaction for the Detection of Hepatitis C Virus RNA. <i>Microbiology and Immunology</i> , 1998, 42, 549-553.	0.7	4
250	Evaluation of Serological Diagnosis of Borna Disease Virus Infection Using Recombinant Proteins in Experimentally Infected Rats.. <i>Journal of Veterinary Medical Science</i> , 1998, 60, 531-534.	0.3	11
251	Hepatitis G Virus Replication in Human Cultured Cells Displaying Susceptibility to Hepatitis C Virus Infection. <i>Biochemical and Biophysical Research Communications</i> , 1997, 235, 505-508.	1.0	26
252	Analysis of the cell tropism of HCV by using in vitro HCV-infected human lymphocytes and hepatocytes. <i>Journal of Hepatology</i> , 1997, 27, 445-454.	1.8	45

#	ARTICLE	IF	CITATIONS
253	Hepatitis C Virus in Pelvic Lymph Nodes and Female Reproductive Organs. Japanese Journal of Cancer Research, 1997, 88, 925-927.	1.7	21
254	Long-Term Human T-Cell Culture System Supporting Hepatitis C Virus Replication. Biochemical and Biophysical Research Communications, 1996, 227, 822-826.	1.0	41
255	Replication of Hepatitis C Virus in Cultured Non-neoplastic Human Hepatocytes. Japanese Journal of Cancer Research, 1996, 87, 787-792.	1.7	70
256	Cytolytic Activity Induced by Intramuscular Injection of Plasmid DNA Expressing the Nucleocapsid Protein of the JHM Strain of Mouse Hepatitis Virus into C57BL/6 Mice.. Journal of Veterinary Medical Science, 1996, 58, 731-735.	0.3	1
257	Inhibition of Viral Multiplication in Cells Chronically Infected with Mouse Hepatitis Virus by Antisense RNA against the Polymerase Gene.. Journal of Veterinary Medical Science, 1995, 57, 563-565.	0.3	4
258	Characterization of DBT Cell Clones Derived from Cells Persistently Infected with the JHM Strain of Mouse Hepatitis Virus.. Journal of Veterinary Medical Science, 1995, 57, 813-817.	0.3	8
259	In Vivo and In Vitro Transcription of Small mRNAs Containing a Leader Sequence from Mouse Hepatitis Virus Strain JHM.. Journal of Veterinary Medical Science, 1994, 56, 1013-1015.	0.3	0
260	Inhibition of Viral Multiplication by Hammerhead Ribozymes Targeted against the Polymerase Gene of Mouse Hepatitis Virus.. Journal of Veterinary Medical Science, 1994, 56, 939-945.	0.3	8
261	Both Antisense and Sense RNAs against the Nucleocapsid Protein Gene Inhibit the Multiplication of Mouse Hepatitis Virus.. Journal of Veterinary Medical Science, 1994, 56, 211-215.	0.3	7
262	Application of DNA Fingerprinting in the Hokkaido Brown Bear (<i>Ursus arctos yesoensis</i>).. Journal of Veterinary Medical Science, 1994, 56, 887-890.	0.3	4
263	Inhibition of Mouse Hepatitis Virus Multiplication by Antisense Oligonucleotide, Antisense RNA, Sense RNA and Ribozyme. Advances in Experimental Medicine and Biology, 1994, 342, 129-135.	0.8	9
264	Inhibition of Mouse Hepatitis Virus Multiplication by an Oligonucleotide Complementary to the Leader RNA.. Journal of Veterinary Medical Science, 1992, 54, 465-472.	0.3	11
265	Exposure of pSV2-Neo Plasmid DNA to X-Rays Enhances Its Ability to Transform Mouse DBT Cells.. Journal of Veterinary Medical Science, 1991, 53, 941-943.	0.3	0
266	Laboratory Diagnosis. , 0, , 45-85.		1