

Tetsuya Mizutani

List of Publications by Year in descending order

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266

papers

7,031

citations

66343

42

h-index

95266

68

g-index

276

all docs

276

docs citations

276

times ranked

8409

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.	2.7	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.	0.9	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.	1.3	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, .	2.9	4
5	Genomic diversity and intragenic recombination of species C rotaviruses. <i>Journal of General Virology</i> , 2022, 103, .	2.9	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.	5.6	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.	2.3	2
8	Dispersal history of <i>Miniopterus fuliginosus</i> bats and their associated viruses in east Asia. <i>PLoS ONE</i> , 2021, 16, e0244006.	2.5	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.	0.9	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.	3.5	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.	3.3	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.	2.1	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.	3.7	6
15	Geographic Distribution and Phylogeny of Soricine Shrew-Borne Seewis Virus and Altai Virus in Russia. <i>Viruses</i> , 2021, 13, 1286.	3.3	5
16	Rodent-Borne Orthohantaviruses in Vietnam, Madagascar and Japan. <i>Viruses</i> , 2021, 13, 1343.	3.3	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.	2.1	19
18	Confirmation of <i>Oryctes rhinoceros</i> nudivirus infections in G-haplotype coconut rhinoceros beetles (<i>Oryctes rhinoceros</i>) from Palauan PCR-positive populations. <i>Scientific Reports</i> , 2021, 11, 18820.	3.3	14

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19	African pygmy hedgehog adenovirus: Virus replication, virus-induced cytopathogenesis and activation of mitogen-activated protein kinase signaling pathways in infected MDCK cells. Research in Veterinary Science, 2021, 139, 152-158.	1.9	0
20	Copper content in ascitic fluid is associated with angiogenesis and progression in ovarian cancer. Journal of Trace Elements in Medicine and Biology, 2021, 68, 126865.	3.0	15
21	Establishment of Intestinal Organoid from Rousettus leschenaultii and the Susceptibility to Bat-Associated Viruses, SARS-CoV-2 and Pteropine Orthoreovirus. International Journal of Molecular Sciences, 2021, 22, 10763.	4.1	14
22	Identification of G-quadruplex sequences in severe acute respiratory syndrome coronavirus 2. Translational and Regulatory Sciences, 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. Endocrine Journal, 2020, 67, 91-94.	1.6	5
24	Complete genome sequencing and genetic analysis of a Japanese porcine torovirus strain detected in swine feces. Archives of Virology, 2020, 165, 471-477.	2.1	7
25	Molecular characterization of feline paramyxovirus in Japanese cat populations. Archives of Virology, 2020, 165, 413-418.	2.1	19
26	Development of a one-run real-time PCR detection system for pathogens associated with porcine respiratory diseases. Journal of Veterinary Medical Science, 2020, 82, 217-223.	0.9	30
27	Genetic Diversity and Phylogeography of Thottapalayam thottimvirus (Hantaviridae) in Asian House Shrew (Suncus murinus) in Eurasia. Frontiers in Cellular and Infection Microbiology, 2020, 10, 438.	3.9	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. Archives of Virology, 2020, 165, 2909-2914.	2.1	3
29	Increased proportion of apoptotic cells in cat kidney tissues infected with feline morbillivirus. Archives of Virology, 2020, 165, 2647-2651.	2.1	0
30	Nutritional status and dietary diversity of school-age children living with HIV: a cross-sectional study in Phnom Penh, Cambodia. BMC Public Health, 2020, 20, 1181.	2.9	12
31	Draft Genome Sequence of Blood-Origin Streptococcus canis Strain FU149, Isolated from a Dog with Necrotizing Soft Tissue Infection. Microbiology Resource Announcements, 2020, 9, .	0.6	3
32	Draft Genome Sequences of Seven Streptococcus canis Strains Isolated from Diseased Companion Animals in Japan. Microbiology Resource Announcements, 2020, 9, .	0.6	6
33	Terminal Genome Sequences of the Soft Tick Bunyavirus. Microbiology Resource Announcements, 2020, 9, . Mosquito-borne viruses, insect-specific flaviviruses (family <i>Flaviviridae</i>, genus) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 16	0.6	2
34		0.9	10
35	mosquitoes. Journal of Veterinary Medical Science, 2020, 82, 1030-1041. ATeam technology for detecting early signs of viral cytopathic effect. Journal of Veterinary Medical Science, 2020, 82, 387-393.	0.9	2
36	Bacterial and protozoan pathogens/symbionts in ticks infecting wild grasscutters (Thryonomys) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 62	2.0	9

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37	Efficacy of primary liver organoid culture from different stages of non-alcoholic steatohepatitis (NASH) mouse model. <i>Biomaterials</i> , 2020, 237, 119823.	11.4	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.	3.3	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.	0.9	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.	2.9	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.	3.6	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.	2.2	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.	2.3	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.	3.9	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.	2.3	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.	2.4	6
48	Gut microbiota confers host resistance to obesity by metabolizing dietary polyunsaturated fatty acids. <i>Nature Communications</i> , 2019, 10, 4007.	12.8	231
49	Phylogenetic analysis of novel posaviruses detected in feces of Japanese pigs with posaviruses and posa-like viruses of vertebrates and invertebrates. <i>Archives of Virology</i> , 2019, 164, 2147-2151.	2.1	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.	2.2	6
51	Association of feline morbillivirus infection with defined pathological changes in cat kidney tissues. <i>Veterinary Microbiology</i> , 2019, 228, 12-19.	1.9	24
52	Cloning of carrier cells infected with oncolytic adenovirus driven by <i>CMV</i> promoter and biosafety studies. <i>Journal of Gene Medicine</i> , 2019, 21, e3064.	2.8	5
53	The complete genomic sequence of <i>Rhinolophus</i> gammaherpesvirus 1 isolated from a greater horseshoe bat. <i>Archives of Virology</i> , 2019, 164, 317-319.	2.1	5
54	Complete genomic analysis and molecular characterization of Japanese porcine sapeloviruses. <i>Virus Genes</i> , 2019, 55, 198-208.	1.6	9

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55	Complete Genome Sequence of an Adenovirus-1 Isolate from an African Pygmy Hedgehog (<i>Atelerix</i>) Tj ETQq1 1 0.784314 rgBT /Ove 2019, 8, .	0.6	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, .	0.6	6
57	Epidemic nodular facial myxomatous dermatitis in juvenile Cranwell's horned frogs <i>Ceratophrys cranwelli</i> . <i>Diseases of Aquatic Organisms</i> , 2019, 134, 57-64.	1.0	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.	4.4	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.	2.2	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.	2.1	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.	2.2	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.	1.6	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.	2.9	13
65	Metagenomic identification and sequence analysis of a Teschovirus A-related virus in porcine feces in Japan, 2014-2016. <i>Infection, Genetics and Evolution</i> , 2018, 66, 210-216.	2.3	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.	1.3	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.	2.2	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.	2.5	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.	0.8	3
71	Luteinizing Hormone Facilitates Antral Follicular Maturation and Survival via Thecal Paracrine Signaling in Cattle. <i>Endocrinology</i> , 2018, 159, 2337-2347.	2.8	27
72	Molecular diversity of the faecal microbiota of Toy Poodles in Japan. <i>Journal of Veterinary Medical Science</i> , 2018, 80, 749-754.	0.9	23

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73	Application of the SureSelect target enrichment system for next-generation sequencing to obtain the complete genome sequence of bovine leukemia virus. Archives of Virology, 2018, 163, 3155-3159.	2.1	7
74	Whole genome analysis of a novel neurotropic bovine astrovirus detected in a Japanese black steer with non-suppurative encephalomyelitis in Japan. Archives of Virology, 2018, 163, 2805-2810.	2.1	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. PLoS ONE, 2018, 13, e0190819.	2.5	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. PLoS ONE, 2018, 13, e0194103.	2.5	29
77	An ascovirus isolated from Spodoptera litura (Noctuidae: Lepidoptera) transmitted by the generalist endoparasitoid Meteorus pulchricornis (Braconidae: Hymenoptera). Journal of General Virology, 2018, 99, 574-584.	2.9	12
78	Decreased expression of the immediate early protein, ICP4, by deletion of the tegument protein VP22 of equine herpesvirus type 1. Journal of Veterinary Medical Science, 2018, 80, 311-315.	0.9	2
79	Angiotensin II increase the pulmonary metastasis through the vascular endothelial cell adherent pathway in hematogenous metastasis of melanoma cells. Proceedings for Annual Meeting of the Japanese Pharmacological Society, 2018, WCP2018, PO1-8-11.	0.0	0
80	Diversity in VP3, NSP3, and NSP4 of rotavirus B detected from Japanese cattle. Infection, Genetics and Evolution, 2017, 49, 97-103.	2.3	10
81	Whole genome analysis of porcine astroviruses detected in Japanese pigs reveals genetic diversity and possible intra-genotypic recombination. Infection, Genetics and Evolution, 2017, 50, 38-48.	2.3	30
82	First isolation and characterization of pteropine orthoreoviruses in fruit bats in the Philippines. Archives of Virology, 2017, 162, 1529-1539.	2.1	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. Archives of Virology, 2017, 162, 2257-2269.	2.1	6
84	Complete genome analysis of porcine kobuviruses from the feces of pigs in Japan. Virus Genes, 2017, 53, 593-602.	1.6	8
85	Characterization of avian paramyxovirus serotype 14, a novel serotype, isolated from a duck fecal sample in Japan. Virus Research, 2017, 228, 46-57.	2.2	34
86	Mycobacterium avium subsp. hominissuis meningoencephalitis in a cat. Veterinary Microbiology, 2017, 204, 43-45.	1.9	7
87	Genetic diversity and intergenogroup recombination events of sapoviruses detected from feces of pigs in Japan. Infection, Genetics and Evolution, 2017, 55, 209-217.	2.3	20
88	Isolation and characterization of Tarumizu tick virus: A new coltivirus from Haemaphysalis flava ticks in Japan. Virus Research, 2017, 242, 131-140.	2.2	34
89	Complete Genome Sequence of Mycobacterium stephanolepidis. Genome Announcements, 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. Vaccine, 2017, 35, 5622-5628.	3.8	8

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91	Molecular characteristics and prevalence of small ruminant lentiviruses in goats in Japan. Archives of Virology, 2017, 162, 3007-3015.	2.1	5
92	Identification of a novel bovine enterovirus possessing highly divergent amino acid sequences in capsid protein. BMC Microbiology, 2017, 17, 18.	3.3	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.	0.9	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.	0.9	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.	0.9	5
96	Isolation and characterization of a new iflavivirus from Armigeres spp. mosquitoes in the Philippines. Journal of General Virology, 2017, 98, 2876-2881.	2.9	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 & Anguilla japonica & Caught from Their Spawning Area. Fish Pathology, 2016, 51, 64-66.	0.7	5
99	Involvement of the 3' UTR Untranslated Region in Encapsidation of the Hepatitis C Virus. PLoS Pathogens, 2016, 12, e1005441.	4.7	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.	2.1	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.	2.3	17
103	Detection of skunk adenovirus 1 (SkAdV-1) in an African pigmy hedgehog (Atelerix albiventris). Veterinary Record Case Reports, 2016, 4, e000321.	0.2	10
104	Characterization and phylogenetic analysis of a novel picornavirus from swine feces in Japan. Archives of Virology, 2016, 161, 1685-1690.	2.1	9
105	Genome sequence of a novel victorivirus identified in the phytopathogenic fungus Alternaria arborescens. Archives of Virology, 2016, 161, 1701-1704.	2.1	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.	0.9	26
107	Detection of Japanese eel endothelial cells-infecting virus in & Anguilla japonica & elvers. Journal of Veterinary Medical Science, 2016, 78, 705-707.	0.9	13
108	Detection of & Campylobacter jejuni & in rectal swab samples from & Rousettus amplexicaudatus & in the Philippines. Journal of Veterinary Medical Science, 2016, 78, 1347-1350.	0.9	13

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109	Detection of novel kobu-like viruses in Japanese black cattle in Japan. Journal of Veterinary Medical Science, 2016, 78, 321-324.	0.9	12
110	A novel Bunyavirus from the soft tick, <i>Argas vespertilionis</i>, in Japan. Journal of Veterinary Medical Science, 2016, 78, 443-445.	0.9	26
111	Molecular epidemiological study of adenovirus infecting western lowland gorillas and humans in and around Moukalaba-Doudou National Park (Gabon). Virus Genes, 2016, 52, 671-678.	1.6	8
112	Identification of further diversity among posaviruses. Archives of Virology, 2016, 161, 3541-3548.	2.1	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.	1.5	10
114	Activation of c-Jun N-terminal kinase by Akabane virus is required for apoptosis. Research in Veterinary Science, 2016, 107, 147-151.	1.9	11
115	Complete Genome Sequencing of Bovine Viral Diarrhea Virus 1, Subgenotypes 1n and 1o. Genome Announcements, 2016, 4, .	0.8	12
116	Whole genome analysis of Japanese bovine toroviruses reveals natural recombination between porcine and bovine toroviruses. Infection, Genetics and Evolution, 2016, 38, 90-95.	2.3	13
117	Isolation of a sp. nov. Ljungan virus from wild birds in Japan. Journal of General Virology, 2016, 97, 1818-1822.	2.9	6
118	Comprehensive pathogen detection associated with four recurrent episodes of Kawasaki disease in a patient during a single year using next-generation sequencing. JMM Case Reports, 2016, 3, e005019.	1.3	11
119	Characterization of Venom and Oviduct Components of Parasitoid Wasp <i>Asobara japonica</i> . PLoS ONE, 2016, 11, e0160210.	2.5	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. Japanese Journal of Veterinary Research, 2016, 64, 5-14.	0.7	16
121	Use of S1 nuclease in deep sequencing for detection of double-stranded RNA viruses. Journal of Veterinary Medical Science, 2015, 77, 1163-1166.	0.9	5
122	Genetic and antigenic analysis of <i>Chlamydia pecorum</i> strains isolated from calves with diarrhea. Journal of Veterinary Medical Science, 2015, 77, 777-782.	0.9	2
123	Transcriptional regulation of genes related to progesterone production [Review]. Endocrine Journal, 2015, 62, 757-763.	1.6	28
124	Quantitative PCR detection of feline morbillivirus in cat urine samples. Journal of Veterinary Medical Science, 2015, 77, 1701-1703.	0.9	16
125	Emergence of infectious malignant thrombocytopenia in Japanese macaques (<i>Macaca fuscata</i>) by SRV-4 after transmission to a novel host. Scientific Reports, 2015, 5, 8850.	3.3	14
126	Ovarian mucinous adenocarcinoma with functioning stroma in postmenopausal women: aromatase and SF-1 expressions. Journal of Ovarian Research, 2015, 8, 73.	3.0	8

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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)-scpIIa</i> gene that confers kasugamycin resistance in <i>Acidovorax avenae</i> ssp. <i>avenae</i> . <i>Molecular Plant Pathology</i> , 2015, 16, 288-300.	4.2	10
129	Characterization of Self-Assembled Virus-Like Particles of Merkel Cell Polyomavirus. <i>PLoS ONE</i> , 2015, 10, e0115646.	2.5	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.	1.9	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.	7.2	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.	2.1	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.	1.9	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.	3.2	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.	2.1	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.	2.9	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.	1.6	6
139	Detection of a novel herpesvirus from bats in the Philippines. <i>Virus Genes</i> , 2015, 51, 136-139.	1.6	19
140	Identification and complete genome analysis of a novel bovine picornavirus in Japan. <i>Virus Research</i> , 2015, 210, 205-212.	2.2	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.	0.2	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.	2.5	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.	2.1	13
144	Molecular, biological, and antigenic characterization of a <i>Border disease virus</i> isolated from a pig during classical swine fever surveillance in Japan. <i>Journal of Veterinary Diagnostic Investigation</i> , 2014, 26, 547-552.	1.1	14

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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.	1.9	9
146	Existence of feline morbillivirus infection in Japanese cat populations. <i>Archives of Virology</i> , 2014, 159, 371-373.	2.1	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.	1.9	21
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