Tetsuya Mizutani

List of Publications by Year
in descending order

266	7,031	66343	95266
papers	citations	42 h-index	68 g-index

276	276	276	8409
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	The First Identification and Retrospective Study of Severe Fever With Thrombocytopenia Syndrome in Japan. Journal of Infectious Diseases, 2014, 209, 816-827.	4.0	672
2	Direct Metagenomic Detection of Viral Pathogens in Nasal and Fecal Specimens Using an Unbiased High-Throughput Sequencing Approach. PLoS ONE, 2009, 4, e4219.	2.5	240
3	Gut microbiota confers host resistance to obesity by metabolizing dietary polyunsaturated fatty acids. Nature Communications, 2019, 10, 4007.	12.8	231
4	Bat Coronaviruses and Experimental Infection of Bats, the Philippines. Emerging Infectious Diseases, 2010, 16, 1217-1223.	4.3	177
5	Phosphorylation of p38 MAPK and its downstream targets in SARS coronavirus-infected cells. Biochemical and Biophysical Research Communications, 2004, 319, 1228-1234.	2.1	141
6	Vesicular stomatitis virus pseudotyped with severe acute respiratory syndrome coronavirus spike protein. Journal of General Virology, 2005, 86, 2269-2274.	2.9	133
7	Lethal Canine Distemper Virus Outbreak in Cynomolgus Monkeys in Japan in 2008. Journal of Virology, 2013, 87, 1105-1114.	3.4	112
8	Murine Coronavirus Replication-Induced p38 Mitogen-Activated Protein Kinase Activation Promotes Interleukin-6 Production and Virus Replication in Cultured Cells. Journal of Virology, 2002, 76, 5937-5948.	3.4	106
9	Species-independent detection of RNA virus by representational difference analysis using non-ribosomal hexanucleotides for reverse transcription. Nucleic Acids Research, 2005, 33, e65-e65.	14.5	105
10	LC16m8, a Highly Attenuated Vaccinia Virus Vaccine Lacking Expression of the Membrane Protein B5R, Protects Monkeys from Monkeypox. Journal of Virology, 2006, 80, 5179-5188.	3.4	101
11	Human hepatocyte clonal cell lines that support persistent replication of hepatitis C virus. Virus Research, 1998, 56, 157-167.	2.2	99
12	JNK and PI3k/Akt signaling pathways are required for establishing persistent SARS-CoV infection in Vero E6 cells. Biochimica Et Biophysica Acta - Molecular Basis of Disease, 2005, 1741, 4-10.	3.8	90
13	Virulence and pathophysiology of the Congo Basin and West African strains of monkeypox virus in non-human primates. Journal of General Virology, 2009, 90, 2266-2271.	2.9	86
14	Reston Ebolavirus Antibodies in Bats, the Philippines. Emerging Infectious Diseases, 2011, 17, 1559-60.	4.3	85
15	Distribution and characterization of tick-borne encephalitis viruses from Siberia and far-eastern Asia. Journal of General Virology, 2001, 82, 1319-1328.	2.9	84
16	Different chemokine expression in lethal and nonâ€lethal murine west nile virus infection. Journal of Medical Virology, 2004, 74, 507-513.	5.0	78
17	Isolation of Novel Adenovirus from Fruit Bat (Pteropus dasymallus yayeyamae). Emerging Infectious Diseases, 2008, 14, 347-349.	4.3	77
18	Establishment of a novel experimental model for muscleâ€invasive bladder cancer using a dog bladder cancer organoid culture. Cancer Science, 2019, 110, 2806-2821.	3.9	75

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19	Inhibitory effect of mizoribine and ribavirin on the replication of severe acute respiratory syndrome (SARS)-associated coronavirus. Antiviral Research, 2005, 66, 159-163.	4.1	73
20	Replication of Hepatitis C Virus in Cultured Non-neoplastic Human Hepatocytes. Japanese Journal of Cancer Research, 1996, 87, 787-792.	1.7	70
21	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
22	Importance of Akt signaling pathway for apoptosis in SARS-CoV-infected Vero E6 cells. Virology, 2004, 327, 169-174.	2.4	68
23	Tyrosine dephosphorylation of STAT3 in SARS coronavirus-infected Vero E6 cells. FEBS Letters, 2004, 577, 187-192.	2.8	65
24	Full genome analysis of bovine astrovirus from fecal samples of cattle in Japan: identification of possible interspecies transmission of bovine astrovirus. Archives of Virology, 2015, 160, 2491-2501.	2.1	65
25	Phylogenetic and virulence analysis of tick-borne encephalitis viruses from Japan and far-eastern Russia. Journal of General Virology, 1999, 80, 3127-3135.	2.9	65
26	Epidemic Myalgia in Adults Associated with Human Parechovirus Type 3 Infection, Yamagata, Japan, 2008. Emerging Infectious Diseases, 2012, 18, 1787-1793.	4.3	65
27	Identification and molecular characterization of a new nonsegmented double-stranded RNA virus isolated from Culex mosquitoes in Japan. Virus Research, 2011, 155, 147-155.	2.2	62
28	Canine Distemper Virus Associated with a Lethal Outbreak in Monkeys Can Readily Adapt To Use Human Receptors. Journal of Virology, 2013, 87, 7170-7175.	3.4	60
29	Role of the N-linked glycans of the prM and E envelope proteins in tick-borne encephalitis virus particle secretion. Vaccine, 2005, 23, 3043-3052.	3.8	59
30	Loopâ€mediated isothermal amplificationâ€based diagnostic assay for monkeypox virus infections. Journal of Medical Virology, 2009, 81, 1102-1108.	5.0	59
31	Existence of feline morbillivirus infection in Japanese cat populations. Archives of Virology, 2014, 159, 371-373.	2.1	57
32	Protection against tick-borne encephalitis virus isolated in Japan by active and passive immunization. Vaccine, 1999, 17, 1532-1539.	3.8	55
33	A BHK-21 cell culture-adapted tick-borne encephalitis virus mutant is attenuated for neuroinvasiveness. Vaccine, 2003, 21, 4043-4051.	3.8	51
34	BoLA-DRB3 Polymorphism is Associated with Differential Susceptibility to Bovine Leukemia Virus-Induced Lymphoma and Proviral Load. Viruses, 2020, 12, 352.	3.3	51
35	Efficacy of primary liver organoid culture from different stages of non-alcoholic steatohepatitis (NASH) mouse model. Biomaterials, 2020, 237, 119823.	11.4	50
36	Evaluation of European tick-borne encephalitis virus vaccine against recent Siberian and far-eastern subtype strains. Vaccine, 2001, 19, 4774-4779.	3.8	47

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37	Postnatal care could be the key to improving the continuum of care in maternal and child health in Ratanakiri, Cambodia. PLoS ONE, 2018, 13, e0198829.	2.5	47
38	Analysis of the cell tropism of HCV by using in vitro HCV-infected human lymphocytes and hepatocytes. Journal of Hepatology, 1997, 27, 445-454.	3.7	45
39	Pathogenicity of tick-borne encephalitis virus isolated in Hokkaido, Japan in mouse model. Vaccine, 1999, 17, 779-787.	3.8	45
40	Single point mutation in tick-borne encephalitis virus prM protein induces a reduction of virus particle secretion. Journal of General Virology, 2004, 85, 3049-3058.	2.9	45
41	Novel DNA virus isolated from samples showing endothelial cell necrosis in the Japanese eel, Anguilla japonica. Virology, 2011, 412, 179-187.	2.4	45
42	Imported Case of Acute Respiratory Tract Infection Associated with a Member of Species Nelson Bay Orthoreovirus. PLoS ONE, 2014, 9, e92777.	2.5	44
43	Characterization of in vitro and in vivo Antiviral Activity of Lactoferrin and Ribavirin upon Hantavirus.. Journal of Veterinary Medical Science, 2001, 63, 637-645.	0.9	43
44	Identification, characterization and full-length sequence analysis of a novel dsRNA virus isolated from the arboreal ant Camponotus yamaokai. Journal of General Virology, 2015, 96, 1930-1937.	2.9	43
45	Amino acid changes responsible for attenuation of virus neurovirulence in an infectious cDNA clone of the Oshima strain of Tick-borne encephalitis virus. Journal of General Virology, 2004, 85, 1007-1018.	2.9	42
46	Long-Term Human T-Cell Culture System Supporting Hepatitis C Virus Replication. Biochemical and Biophysical Research Communications, 1996, 227, 822-826.	2.1	41
47	Rapid Genome Sequencing of RNA Viruses. Emerging Infectious Diseases, 2007, 13, 322-324.	4.3	41
48	Development of Recombinant Nucleoprotein-Based Diagnostic Systems for Lassa Fever. Vaccine Journal, 2007, 14, 1182-1189.	3.1	40
49	Evaluation of a novel vesicular stomatitis virus pseudotype-based assay for detection of neutralizing antibody responses to SARS-CoV. Journal of Medical Virology, 2006, 78, 1509-1512.	5.0	39
50	Novel virus discovery in field-collected mosquito larvae using an improved system for rapid determination of viral RNA sequences (RDV ver4.0). Archives of Virology, 2009, 154, 153-158.	2.1	35
51	Novel Betaherpesvirus in Bats. Emerging Infectious Diseases, 2010, 16, 986-988.	4.3	35
52	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. Veterinary Microbiology, 2014, 171, 66-73.	1.9	35
53	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
54	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

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55	Rapid determination of viral RNA sequences in mosquitoes collected in the field. Journal of Virological Methods, 2007, 146, 372-374.	2.1	33
56	Sequence and phylogenetic analyses of Saffold cardiovirus from children with exudative tonsillitis in Yamagata, Japan. Scandinavian Journal of Infectious Diseases, 2010, 42, 950-952.	1.5	33
57	Antigen-capture ELISA for the detection of Rift Valley fever virus nucleoprotein using new monoclonal antibodies. Journal of Virological Methods, 2012, 180, 68-74.	2.1	32
58	Identification of a natural recombination in the F and H genes of feline morbillivirus. Virology, 2014, 468-470, 524-531.	2.4	32
59	Mechanisms of establishment of persistent SARS-CoV-infected cells. Biochemical and Biophysical Research Communications, 2006, 347, 261-265.	2.1	31
60	Detection of a new bat gammaherpesvirus in the Philippines. Virus Genes, 2009, 39, 90-93.	1.6	30
61	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
62	Characterization of a novel thogotovirus isolated from Amblyomma testudinarium ticks in Ehime, Japan: A significant phylogenetic relationship to Bourbon virus. Virus Research, 2018, 249, 57-65.	2.2	30
63	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
64	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
65	Characterization of Dak Nong virus, an insect nidovirus isolated from Culex mosquitoes in Vietnam. Archives of Virology, 2013, 158, 2273-2284.	2.1	29
66	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
67	Genetic and antigenic characterization of the Amur virus associated with hemorrhagic fever with renal syndrome. Virus Research, 2004, 101, 127-134.	2.2	28
68	Transcriptional regulation of genes related to progesterone production [Review]. Endocrine Journal, 2015, 62, 757-763.	1.6	28
69	Borna Disease Virus Infection in Domestic Cats: Evaluation by RNA and Antibody Detection. Journal of Veterinary Medical Science, 1999, 61, 1167-1170.	0.9	27
70	Luteinizing Hormone Facilitates Antral Follicular Maturation and Survival via Thecal Paracrine Signaling in Cattle. Endocrinology, 2018, 159, 2337-2347.	2.8	27
71	Hepatitis G Virus Replication in Human Cultured Cells Displaying Susceptibility to Hepatitis C Virus Infection. Biochemical and Biophysical Research Communications, 1997, 235, 505-508.	2.1	26
72	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

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73	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
74	First isolation and characterization of pteropine orthoreoviruses in fruit bats in the Philippines. Archives of Virology, 2017, 162, 1529-1539.	2.1	26
75	Genetic Characterization of Hantaviruses Transmitted by the Korean Field Mouse (<i>Apodemus) Tj ETQq1 1 0.784314 rgBT /Overloc	4.3	25
76	Recombinant nucleocapsid protein-based IgG enzyme-linked immunosorbent assay for the serological diagnosis of SARS. Journal of Virological Methods, 2005, 125, 181-186.	2.1	25
77	Synthesis and biological evaluation of nucleoside analogues having 6-chloropurine as anti-SARS-CoV agents. Bioorganic and Medicinal Chemistry Letters, 2007, 17, 2470-2473.	2.2	25
78	Genome sequence of a novel victorivirus identified in the phytopathogenic fungus Alternaria arborescens. Archives of Virology, 2016, 161, 1701-1704.	2.1	25
79	Enzyme-linked immunosorbent assay using recombinant antigens expressed in mammalian cells for serodiagnosis of tick-borne encephalitis. Journal of Virological Methods, 2003, 108, 171-179.	2.1	24
80	Genomic and serological detection of bat coronavirus from bats in the Philippines. Archives of Virology, 2012, 157, 2349-2355.	2.1	24
81	Angiotensin II promotes pulmonary metastasis of melanoma through the activation of adhesion molecules in vascular endothelial cells. Biochemical Pharmacology, 2018, 154, 136-147.	4.4	24
82	Isolation and characterization of Kabuto Mountain virus, a new tick-borne phlebovirus from Haemaphysalis flava ticks in Japan. Virus Research, 2018, 244, 252-261.	2.2	24
83	Association of feline morbillivirus infection with defined pathological changes in cat kidney tissues. Veterinary Microbiology, 2019, 228, 12-19.	1.9	24
84	Inhibition of cell proliferation by SARS-CoV infection in Vero E6 cells. FEMS Immunology and Medical Microbiology, 2006, 46, 236-243.	2.7	23
85	Detection of bat coronaviruses from Miniopterus fuliginosus in Japan. Virus Genes, 2012, 44, 40-44.	1.6	23
86	Molecular diversity of the faecal microbiota of Toy Poodles in Japan. Journal of Veterinary Medical Science, 2018, 80, 749-754.	0.9	23
87	Hepatitis C Virus in Pelvic Lymph Nodes and Female Reproductive Organs. Japanese Journal of Cancer Research, 1997, 88, 925-927.	1.7	21
88	A novel sapelovirus-like virus isolation from wild boar. Virus Genes, 2011, 43, 243-248.	1.6	21
89	Transcriptional regulation of human ferredoxin reductase through an intronic enhancer in steroidogenic cells. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2014, 1839, 33-42.	1.9	21
90	Involvement of the 3â€™ Untranslated Region in Encapsidation of the Hepatitis C Virus. PLoS Pathogens, 2016, 12, e1005441.	4.7	21

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91	Genome sequence of a novel mitovirus identified in the phytopathogenic fungus Alternaria arborescens. Archives of Virology, 2016, 161, 2627-2631.	2.1	21
92	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. Journal of Medical Virology, 2005, 75, 504-512.	5.0	20
93	A seroepidemiologic study of Reston ebolavirus in swine in the Philippines. BMC Veterinary Research, 2012, 8, 82.	1.9	20
94	Characterization of Self-Assembled Virus-Like Particles of Merkel Cell Polyomavirus. PLoS ONE, 2015, 10, e0115646.	2.5	20
95	Identification and complete genome analysis of a novel bovine picornavirus in Japan. Virus Research, 2015, 210, 205-212.	2.2	20
96	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
97	Two Novel Endornaviruses Co-infecting a Phytophthora Pathogen of Asparagus officinalis Modulate the Developmental Stages and Fungicide Sensitivities of the Host Oomycete. Frontiers in Microbiology, 2021, 12, 633502.	3.5	20
98	A Serosurvey of Borna Disease Virus Infection in Wild Rats by a Capture ELISA. Journal of Veterinary Medical Science, 1999, 61, 113-117.	0.9	19
99	Detection of a novel herpesvirus from bats in the Philippines. Virus Genes, 2015, 51, 136-139.	1.6	19
100	Molecular characterization of feline paramyxovirus in Japanese cat populations. Archives of Virology, 2020, 165, 413-418.	2.1	19
101	Natto extract, a Japanese fermented soybean food, directly inhibits viral infections including SARS-CoV-2 inÂvitro. Biochemical and Biophysical Research Communications, 2021, 570, 21-25.	2.1	19
102	Sequencing and Phylogenetic Analyses of Saffold Cardiovirus (SAFV) Genotype 3 Isolates from Children with Upper Respiratory Infection in Gunma, Japan. Japanese Journal of Infectious Diseases, 2010, 63, 378-380.	1.2	19
103	Detection of Hantaviral Antibodies among Patients with Hepatitis of Unknown Etiology in Japan. Microbiology and Immunology, 2000, 44, 357-362.	1.4	18
104	Induction of protective immunity against severe acute respiratory syndrome coronavirus (SARS-CoV) infection using highly attenuated recombinant vaccinia virus DIs. Virology, 2006, 351, 368-380.	2.4	18
105	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. Journal of Biological Chemistry, 2010, 285, 28240-28251.	3.4	18
106	Epizootiological and Epidemiological Study of Hantavirus Infection in Japan. Microbiology and Immunology, 2004, 48, 843-851.	1.4	17
107	Regulation of p90RSK phosphorylation by SARS-CoV infection in Vero E6 cells. FEBS Letters, 2006, 580, 1417-1424.	2.8	17
108	Characterization of Monoclonal Antibodies to Junin Virus Nucleocapsid Protein and Application to the Diagnosis of Hemorrhagic Fever Caused by South American Arenaviruses. Vaccine Journal, 2009, 16, 1132-1138.	3.1	17

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109	Molecular epidemiology of avian bornavirus from pet birds in Japan. Virus Genes, 2013, 47, 173-177.	1.6	17
110	H2 genotypes of G4P[6], G5P[7], and G9[23] porcine rotaviruses show super-short RNA electropherotypes. Veterinary Microbiology, 2015, 176, 250-256.	1.9	17
111	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
112	Identification of a novel bovine enterovirus possessing highly divergent amino acid sequences in capsid protein. BMC Microbiology, 2017, 17, 18.	3.3	17
113	Nascent Synthesis of Leader Sequence-Containing Subgenomic mRNAs in Coronavirus Genome-Length Replicative Intermediate RNA. Virology, 2000, 275, 238-243.	2.4	16
114	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. Endocrine Journal, 2004, 51, 569-577.	1.6	16
115	Development of an enzyme-linked immunosorbent assay for serological diagnosis of tick-borne encephalitis using subviral particles. Journal of Virological Methods, 2006, 134, 55-60.	2.1	16
116	Application of quantitative gene expression analysis for pertussis vaccine safety control. Vaccine, 2008, 26, 4686-4696.	3.8	16
117	Detection of enterovirus genome sequence from diarrheal feces of goat. Virus Genes, 2014, 48, 550-552.	1.6	16
118	Quantitative PCR detection of feline morbillivirus in cat urine samples. Journal of Veterinary Medical Science, 2015, 77, 1701-1703.	0.9	16
119	Isolation and characterization of a new iflavirus from Armigeres spp. mosquitoes in the Philippines. Journal of General Virology, 2017, 98, 2876-2881.	2.9	16
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	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
122	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
123	Characterization of Monoclonal Antibodies against Hokkaido Strain Tickâ€Borne Encephalitis Virus. Microbiology and Immunology, 2000, 44, 533-536.	1.4	14
124	Ligation-mediated amplification for effective rapid determination of viral RNA sequences (RDV). Journal of Clinical Virology, 2008, 43, 56-59.	3.1	14
125	Molecular, biological, and antigenic characterization of a Border disease virus isolated from a pig during classical swine fever surveillance in Japan. Journal of Veterinary Diagnostic Investigation, 2014, 26, 547-552.	1.1	14
126	Emergence of infectious malignant thrombocytopenia in Japanese macaques (Macaca fuscata) by SRV-4 after transmission to a novel host. Scientific Reports, 2015, 5, 8850.	3.3	14

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127	Isolation of Pteropine orthoreovirus from Pteropus vampyrus in Garut, Indonesia. Virus Genes, 2018, 54, 823-827.	1.6	14
128	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. Infection, Genetics and Evolution, 2019, 75, 103975.	2.3	14
129	Confirmation of Oryctes rhinoceros nudivirus infections in G-haplotype coconut rhinoceros beetles (Oryctes rhinoceros) from Palauan PCR-positive populations. Scientific Reports, 2021, 11, 18820.	3.3	14
130	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
131	Expression of apoptosis on rat liver by hepatic vagus hyperactivity after ventromedial hypothalamic lesioning. American Journal of Physiology - Renal Physiology, 2001, 280, G958-G967.	3.4	13
132	Discrimination of West Nile Virus and Japanese Encephalitis Virus Strains Using RTâ€PCR RFLP Analysis. Microbiology and Immunology, 2003, 47, 439-445.	1.4	13
133	Amino Acid Substitutions in the S2 Region Enhance Severe Acute Respiratory Syndrome Coronavirus Infectivity in Rat Angiotensin-Converting Enzyme 2-Expressing Cells. Journal of Virology, 2007, 81, 10831-10834.	3.4	13
134	Characterization of a genetic and antigenic variant of avian paramyxovirus 6 isolated from a migratory wild bird, the red-necked stint (Calidris ruficollis). Archives of Virology, 2014, 159, 3101-3105.	2.1	13
135	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
136	Detection of Japanese eel endothelial cells-infecting virus in <i>Anguilla japonica</i> elvers. Journal of Veterinary Medical Science, 2016, 78, 705-707.	0.9	13
137	Detection of <i>Campylobacter jejuni</i> in rectal swab samples from <i>Rousettus amplexicaudatus</i> in the Philippines. Journal of Veterinary Medical Science, 2016, 78, 1347-1350.	0.9	13
138	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
139	Malaria knowledge, preventive actions, and treatment-seeking behavior among ethnic minorities in Ratanakiri Province, Cambodia: a community-based cross-sectional survey. BMC Public Health, 2018, 18, 1206.	2.9	13
140	Novel reovirus isolation from an Ostrich (Struthio camelus) in Japan. Veterinary Microbiology, 2009, 134, 227-232.	1.9	12
141	Detection of novel kobu-like viruses in Japanese black cattle in Japan. Journal of Veterinary Medical Science, 2016, 78, 321-324.	0.9	12
142	Identification of further diversity among posaviruses. Archives of Virology, 2016, 161, 3541-3548.	2.1	12
143	Complete Genome Sequencing of Bovine Viral Diarrhea Virus 1, Subgenotypes 1n and 1o. Genome Announcements, 2016, 4, .	0.8	12
144	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

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145	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
146	Pathogenic characterization of a cervical lymph node derived from a patient with Kawasaki disease. International Journal of Clinical and Experimental Pathology, 2012, 5, 814-23.	0.5	12
147	Marburgvirus nucleoprotein-capture enzyme-linked immunosorbent assay using monoclonal antibodies to recombinant nucleoprotein: detection of authentic Marburgvirus. Japanese Journal of Infectious Diseases, 2006, 59, 323-5.	1.2	12
148	Diagnosis and assessment of monkeypox virus (MPXV) infection by quantitative PCR assay: differentiation of Congo Basin and West African MPXV strains. Japanese Journal of Infectious Diseases, 2008, 61, 140-2.	1.2	12
149	Inhibition of Mouse Hepatitis Virus Multiplication by an Oligonucleotide Complementary to the Leader RNA.. Journal of Veterinary Medical Science, 1992, 54, 465-472.	0.9	11
150	Evaluation of Serological Diagnosis of Borna Disease Virus Infection Using Recombinant Proteins in Experimentally Infected Rats.. Journal of Veterinary Medical Science, 1998, 60, 531-534.	0.9	11
151	The fluctuations of viral load and serum alanine aminotransferase levels in chronic hepatitis C. Hepatology Research, 2004, 30, 11-17.	3.4	11
152	Whole-genome sequence analysis of G3 and G14 equine group A rotaviruses isolated in the late 1990s and 2009-2010. Archives of Virology, 2015, 160, 1171-1179.	2.1	11
153	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
154	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
155	Detection of Bovine Group A Rotavirus Using Rapid Antigen Detection Kits, RT-PCR and Next-Generation DNA Sequencing. Journal of Veterinary Medical Science, 2013, 75, 1651-1655.	0.9	10
156	Two types of genetic carrier, the <scp>IncP</scp> genomic island and the novel <scp>IncP</scp>â€1Î² plasmid, for the aac(2â€²)â€<scp>Ila</scp> gene that confers kasugamycin resistance in <scp>A</scp>cidovorax avenae ssp. avenae. Molecular Plant Pathology, 2015, 16, 288-300.	4.2	10
157	Detection of skunk adenovirus 1 (SkAdVâ€1) in an African pigmy hedgehog (Atelerix albiventris). Veterinary Record Case Reports, 2016, 4, e000321.	0.2	10
158	Complete Genome Sequencing and Phylogenetic Analysis of a Getah Virus Strain (GenusAlphavirus,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5 Vector-Borne and Zoonotic Diseases, 2016, 16, 769-776.	1.5	10
159	Diversity in VP3, NSP3, and NSP4 of rotavirus B detected from Japanese cattle. Infection, Genetics and Evolution, 2017, 49, 97-103.	2.3	10
160	Metagenomic identification and sequence analysis of a Teschovirus A-related virus in porcine feces in Japan, 2014â€“2016. Infection, Genetics and Evolution, 2018, 66, 210-216.	2.3	10
161	Mosquito-borne viruses, insect-specific flaviviruses (family <i>Flaviviridae</i>, genus) Tj ETQq1 1 0.784314 rgBT /Overlock 10	0.9	10
162	Molecular characterization of full genome sequences of Newcastle disease viruses circulating among vaccinated chickens in Egypt during 2011â€“2013. Journal of Veterinary Medical Science, 2020, 82, 809-816.	0.9	10

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163	Stochastic modelling of the effects of human-mobility restriction and viral infection characteristics on the spread of COVID-19. Scientific Reports, 2021, 11, 6856.	3.3	10
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