

Tung G Phan

List of Publications by Year in descending order

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Version: 2024-02-01

138
papers

5,770
citations

87401

40
h-index

100535

70
g-index

140
all docs

140
docs citations

140
times ranked

7213
citing authors

#	ARTICLE	IF	CITATIONS
1	Secondary syphilis as an initial presentation of HIV. Baylor University Medical Center Proceedings, 2022, 35, 1-2.	0.2	0
2	Development of a One-Step Qualitative RT-PCR Assay to Detect the SARS-CoV-2 Omicron (B.1.1.529) Variant in Respiratory Specimens. Journal of Clinical Microbiology, 2022, 60, jcm0002422.	1.8	22
3	Evaluation of Viral Loads in Patients With SARS-CoV-2 Delta Variant Infection: Higher Loads Do Not Translate Into Different Testing Scenarios. Microbiology Insights, 2022, 15, 117863612210875.	0.9	2
4	Evaluation of the Cepheid Xpert Xpress SARS-CoV-2 test for bronchoalveolar lavage. Journal of Clinical Virology Plus, 2022, 2, 100067.	0.4	3
5	Evaluation of the ePlex Respiratory pathogen panel 2 to detect viral and bacterial pathogens, including SARS-CoV-2 Omicron in nasopharyngeal swabs. Journal of Clinical Virology Plus, 2022, 2, 100072.	0.4	0
6	Clinical evaluation of the Cue's COVID-19 diagnostic test to detect SARS-CoV-2 in the upper respiratory tract. Journal of Medical Virology, 2022, 94, 3517-3519.	2.5	3
7	Whole genome sequencing and evolutionary analysis of G8P [8] rotaviruses emerging in Japan. VirusDisease, 2022, 33, 215-218.	1.0	6
8	Keratitis caused by Nocardia farcinica in a contact lens wearer. , 2022, , .		0
9	First detection of SARS-CoV-2 Omicron BA.4 variant in Western Pennsylvania, United States. Journal of Medical Virology, 2022, 94, 4053-4055.	2.5	17
10	Whole genome sequence of an uncommon G9P[4] species A rotavirus containing DS-1-like (genotype 2) genes in Japan. Archives of Virology, 2022, 167, 1603-1606.	0.9	1
11	Detection of Enterococcus avium in a case of urinary tract infection and haematuria. Access Microbiology, 2022, 4, .	0.2	2
12	Emergence of SARS-CoV-2 Omicron BA.5 variant of concern in Western Pennsylvania, United States. Journal of Medical Virology, 2022, 94, 4593-4594.	2.5	12
13	Diagnostic Tests for COVID-19. Advances in Experimental Medicine and Biology, 2021, 1318, 403-412.	0.8	1
14	Tuberculosis of the rare azygos lobe of the right lung. Respiratory Medicine Case Reports, 2021, 33, 101424.	0.2	1
15	Molecular epidemiology and genetic diversity of norovirus infection in children with acute gastroenteritis in Bangladesh, 2014-2019. Journal of Medical Virology, 2021, 93, 3564-3571.	2.5	8
16	A rare case of polymicrobial brain abscess involving Actinomyces. Radiology Case Reports, 2021, 16, 1123-1126.	0.2	4
17	Vibrio mimicus wound infection in a burn patient. Radiology Case Reports, 2021, 16, 1348-1351.	0.2	10
18	One year into the pandemic: Short-term evolution of SARS-CoV-2 and emergence of new lineages. Infection, Genetics and Evolution, 2021, 92, 104869.	1.0	49

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19	Detection of <i>Enterococcus hirae</i> in a case of acute osteomyelitis. <i>Radiology Case Reports</i> , 2021, 16, 2366-2369.	0.2	4
20	Cutavirus: A newly discovered parvovirus on the rise. <i>Infection, Genetics and Evolution</i> , 2020, 80, 104175.	1.0	6
21	Endophthalmitis resulting from gonococcal keratoconjunctivitis. <i>New Microbes and New Infections</i> , 2020, 36, 100724.	0.8	0
22	Tympanic membrane perforation secondary to <i>Aspergillus niger</i> otomycosis. <i>IDCases</i> , 2020, 22, e00944.	0.4	0
23	Unusual mono-reassortant of a Wa-like G1P[8] species A rotavirus containing a DS-1-like (genotype 2) NSP4 gene. <i>Virus Genes</i> , 2020, 56, 638-641.	0.7	5
24	MALDI-TOF vs. VITEK 2 for identification of <i>Aggregatibacter actinomycetemcomitans</i> chest wall abscess. <i>IDCases</i> , 2020, 20, e00749.	0.4	1
25	SARS-CoV-2 and COVID-19: A genetic, epidemiological, and evolutionary perspective. <i>Infection, Genetics and Evolution</i> , 2020, 84, 104384.	1.0	115
26	Unusual community-associated carbapenem-resistant <i>Acinetobacter baumannii</i> infection, Pennsylvania, USA. <i>IDCases</i> , 2020, 21, e00851.	0.4	2
27	Disseminated cryptococcosis in an immunocompetent patient. <i>Respiratory Medicine Case Reports</i> , 2020, 30, 101034.	0.2	8
28	Genetic diversity and evolution of SARS-CoV-2. <i>Infection, Genetics and Evolution</i> , 2020, 81, 104260.	1.0	498
29	Novel coronavirus: From discovery to clinical diagnostics. <i>Infection, Genetics and Evolution</i> , 2020, 79, 104211.	1.0	209
30	<i>Acinetobacter junii</i> as a rare pathogen of urinary tract infection. <i>Urology Case Reports</i> , 2020, 32, 101209.	0.1	7
31	Genomic characterization of Changuinola viruses from Panama: evidence for multiple genome segment reassortment. <i>Virus Genes</i> , 2020, 56, 527-530.	0.7	1
32	Molecular epidemiology and surveillance of circulating rotavirus among children with gastroenteritis in Bangladesh during 2014–2019. <i>PLoS ONE</i> , 2020, 15, e0242813.	1.1	12
33	<i>Mycobacterium marinum</i> infection of the hand presenting as a nodular skin lesion. <i>Journal of Clinical Tuberculosis and Other Mycobacterial Diseases</i> , 2020, 20, 100166.	0.6	3
34	Viral species richness and composition in young children with loose or watery stool in Ethiopia. <i>BMC Infectious Diseases</i> , 2019, 19, 53.	1.3	18
35	Enteric Virome and Bacterial Microbiota in Children With Ulcerative Colitis and Crohn Disease. <i>Journal of Pediatric Gastroenterology and Nutrition</i> , 2019, 68, 30-36.	0.9	89
36	Nasal virome of dogs with respiratory infection signs include novel taupapillomaviruses. <i>Virus Genes</i> , 2019, 55, 191-197.	0.7	14

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37	A novel passerivirus (family Picornaviridae) in an outbreak of enteritis with high mortality in estrildid finches (<i>Uraeginthus</i> sp.). <i>Archives of Virology</i> , 2018, 163, 1063-1071.	0.9	9
38	Genomic analysis of a novel picornavirus from a migratory waterfowl, greater white-fronted goose (<i>Anser albifrons</i>). <i>Archives of Virology</i> , 2018, 163, 1087-1090.	0.9	7
39	Sera of Peruvians with fever of unknown origins include viral nucleic acids from non-vertebrate hosts. <i>Virus Genes</i> , 2018, 54, 33-40.	0.7	19
40	Multiple divergent picobirnaviruses with functional prokaryotic Shine-Dalgarno ribosome binding sites present in cloacal sample of a diarrheic chicken. <i>Virology</i> , 2018, 525, 62-72.	1.1	26
41	Enteric virome of Ethiopian children participating in a clean water intervention trial. <i>PLoS ONE</i> , 2018, 13, e0202054.	1.1	29
42	Virome of US bovine calf serum. <i>Biologicals</i> , 2017, 46, 64-67.	0.5	39
43	Rotavirus I in feces of a cat with diarrhea. <i>Virus Genes</i> , 2017, 53, 487-490.	0.7	19
44	Genome characterization of a novel megrivirus-related avian picornavirus from a carnivorous wild bird, western marsh harrier (<i>Circus aeruginosus</i>). <i>Archives of Virology</i> , 2017, 162, 2781-2789.	0.9	7
45	Genetic characterization of a second novel picornavirus from an amphibian host, smooth newt (<i>Lissotriton vulgaris</i>). <i>Archives of Virology</i> , 2017, 162, 1043-1050.	0.9	12
46	Small Circular Rep-Encoding Single-Stranded DNA Genomes in Peruvian Diarrhea Virome. <i>Genome Announcements</i> , 2017, 5, .	0.8	15
47	Genomes of viral isolates derived from different mosquitos species. <i>Virus Research</i> , 2017, 242, 49-57.	1.1	40
48	A Naturally Transmitted Epitheliotropic Polyomavirus Pathogenic in Immunodeficient Rats: Characterization, Transmission, and Preliminary Epidemiologic Studies. <i>Toxicologic Pathology</i> , 2017, 45, 593-603.	0.9	10
49	Outbreaks of Neuroinvasive Astrovirus Associated with Encephalomyelitis, Weakness, and Paralysis among Weaned Pigs, Hungary. <i>Emerging Infectious Diseases</i> , 2017, 23, 1982-1993.	2.0	66
50	Genetic Characterization and Classification of Human and Animal Sapoviruses. <i>PLoS ONE</i> , 2016, 11, e0156373.	1.1	71
51	A new protoparvovirus in human fecal samples and cutaneous T cell lymphomas (mycosis fungoides). <i>Virology</i> , 2016, 496, 299-305.	1.1	49
52	A new densovirus in cerebrospinal fluid from a case of anti-NMDA-receptor encephalitis. <i>Archives of Virology</i> , 2016, 161, 3231-3235.	0.9	15
53	Detection of a novel circovirus PCV3 in pigs with cardiac and multi-systemic inflammation. <i>Virology Journal</i> , 2016, 13, 184.	1.4	309
54	A diarrheic chicken simultaneously co-infected with multiple picornaviruses: Complete genome analysis of avian picornaviruses representing up to six genera. <i>Virology</i> , 2016, 489, 63-74.	1.1	36

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55	The fecal virome of South and Central American children with diarrhea includes small circular DNA viral genomes of unknown origin. <i>Archives of Virology</i> , 2016, 161, 959-966.	0.9	36
56	Detection and genetic characterization of a novel parvovirus distantly related to human bufavirus in domestic pigs. <i>Archives of Virology</i> , 2016, 161, 1033-1037.	0.9	27
57	Novel picornavirus in domestic rabbits (<i>Oryctolagus cuniculus</i> var. <i>domestica</i>). <i>Infection, Genetics and Evolution</i> , 2016, 37, 117-122.	1.0	20
58	Absence of giant blood Marseille-like virus DNA detection by polymerase chain reaction in plasma from healthy US blood donors and serum from multiply transfused patients from Cameroon. <i>Transfusion</i> , 2015, 55, 1256-1262.	0.8	10
59	A highly divergent picornavirus in an amphibian, the smooth newt (<i>Lissotriton vulgaris</i>). <i>Journal of General Virology</i> , 2015, 96, 2607-2613.	1.3	19
60	Sequence and phylogenetic analysis identifies a putative novel gyrovirus 3 genotype in ferret feces. <i>Virus Genes</i> , 2015, 50, 137-141.	0.7	20
61	Secondary structure analysis of swine pasivirus (family Picornaviridae) RNA reveals a type-IV IRES and a parechovirus-like 3' UTR organization. <i>Archives of Virology</i> , 2015, 160, 1363-1366.	0.9	3
62	A gyrovirus infecting a sea bird. <i>Archives of Virology</i> , 2015, 160, 2105-2109.	0.9	25
63	Genome analysis of a novel, highly divergent picornavirus from common kestrel (<i>Falco tinnunculus</i>): The first non-enteroviral picornavirus with type-I-like IRES. <i>Infection, Genetics and Evolution</i> , 2015, 32, 425-431.	1.0	18
64	Human polyomavirus 6 DNA in the cerebrospinal fluid of an HIV-positive patient with leukoencephalopathy. <i>Journal of Clinical Virology</i> , 2015, 68, 24-27.	1.6	12
65	Small circular single stranded DNA viral genomes in unexplained cases of human encephalitis, diarrhea, and in untreated sewage. <i>Virology</i> , 2015, 482, 98-104.	1.1	94
66	A new gyrovirus in human feces. <i>Virus Genes</i> , 2015, 51, 132-135.	0.7	22
67	Sesavirus: prototype of a new parvovirus genus in feces of a sea lion. <i>Virus Genes</i> , 2015, 50, 134-136.	0.7	22
68	Bufavirus in Feces of Patients with Gastroenteritis, Finland. <i>Emerging Infectious Diseases</i> , 2014, 20, 1077-1079.	2.0	47
69	New Parvovirus in Child with Unexplained Diarrhea, Tunisia. <i>Emerging Infectious Diseases</i> , 2014, 20, 1911-1913.	2.0	29
70	Genome characterization of a novel chicken picornavirus distantly related to the members of genus Avihepatovirus with a single 2A protein and a megrivirus-like 3' UTR. <i>Infection, Genetics and Evolution</i> , 2014, 28, 333-338.	1.0	6
71	Genomic characterization of a rotavirus G8P[1] detected in a child with diarrhea reveal direct animal-to-human transmission. <i>Infection, Genetics and Evolution</i> , 2014, 27, 402-407.	1.0	14
72	New astrovirus in human feces from Burkina Faso. <i>Journal of Clinical Virology</i> , 2014, 60, 161-164.	1.6	32

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73	Cyclovirus in nasopharyngeal aspirates of Chilean children with respiratory infections. <i>Journal of General Virology</i> , 2014, 95, 922-927.	1.3	43
74	Rosavirus: the prototype of a proposed new genus of the Picornaviridae family. <i>Virus Genes</i> , 2013, 47, 556-558.	0.7	10
75	Novel Human Gammapapillomavirus Species in a Nasal Swab. <i>Genome Announcements</i> , 2013, 1, e0002213.	0.8	10
76	The Perils of Pathogen Discovery: Origin of a Novel Parvovirus-Like Hybrid Genome Traced to Nucleic Acid Extraction Spin Columns. <i>Journal of Virology</i> , 2013, 87, 11966-11977.	1.5	216
77	Genome Sequence of a Novel Virus of the Species Human Adenovirus D Associated with Acute Gastroenteritis. <i>Genome Announcements</i> , 2013, 1, .	0.8	33
78	The Viruses of Wild Pigeon Droppings. <i>PLoS ONE</i> , 2013, 8, e72787.	1.1	108
79	Divergent Astrovirus Associated with Neurologic Disease in Cattle. <i>Emerging Infectious Diseases</i> , 2013, 19, 1385-1392.	2.0	155
80	Genome Sequence of an Unusual Human G10P[8] Rotavirus Detected in Vietnam. <i>Journal of Virology</i> , 2012, 86, 10236-10237.	1.5	13
81	Acute Diarrhea in West African Children: Diverse Enteric Viruses and a Novel Parvovirus Genus. <i>Journal of Virology</i> , 2012, 86, 11024-11030.	1.5	120
82	Genetic Diversity of the Genus Cosavirus in the Family Picornaviridae: A New Species, Recombination, and 26 New Genotypes. <i>PLoS ONE</i> , 2012, 7, e36685.	1.1	45
83	Discovery of a Novel Polyomavirus in Acute Diarrheal Samples from Children. <i>PLoS ONE</i> , 2012, 7, e49449.	1.1	110
84	Novel Human Adenovirus Strain, Bangladesh. <i>Emerging Infectious Diseases</i> , 2012, 18, 846-848.	2.0	43
85	A third gyrovirus species in human faeces. <i>Journal of General Virology</i> , 2012, 93, 1356-1361.	1.3	72
86	Detection and genetic characterization of rotavirus infections in non-hospitalized children with acute gastroenteritis in Japan, 2007-2009. <i>Infection, Genetics and Evolution</i> , 2011, 11, 415-422.	1.0	22
87	Genomic characterization of a novel human adenovirus type 31 recombinant in the hexon gene. <i>Journal of General Virology</i> , 2011, 92, 2770-2775.	1.3	37
88	The Acidic Domain of Hepatitis C Virus NS4A Contributes to RNA Replication and Virus Particle Assembly. <i>Journal of Virology</i> , 2011, 85, 1193-1204.	1.5	43
89	The Fecal Viral Flora of Wild Rodents. <i>PLoS Pathogens</i> , 2011, 7, e1002218.	2.1	304
90	Novel recombinant norovirus in Japan. <i>Virus Genes</i> , 2010, 40, 362-364.	0.7	11

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91	Sequence analysis of the VP7 gene of human rotaviruses G2 and G4 isolated in Japan, China, Thailand, and Vietnam during 2001–2003. <i>Journal of Medical Virology</i> , 2010, 82, 878-885.	2.5	17
92	Molecular and Epidemiological Trend of Sapovirus, and Astrovirus Infection in Japan. <i>Journal of Tropical Pediatrics</i> , 2010, 56, 205-207.	0.7	12
93	Hepatitis C Virus NS2 Protein Contributes to Virus Particle Assembly via Opposing Epistatic Interactions with the E1-E2 Glycoprotein and NS3-NS4A Enzyme Complexes. <i>Journal of Virology</i> , 2009, 83, 8379-8395.	1.5	116
94	Development of genotype-specific primers for differentiation of genotypes A and B of Aichi viruses. <i>Journal of Virological Methods</i> , 2009, 156, 107-110.	1.0	9
95	Molecular and epidemiological trend of rotavirus infection among infants and children in Japan. <i>Infection, Genetics and Evolution</i> , 2009, 9, 955-961.	1.0	13
96	Molecular epidemiology of adenovirus infection among infants and children with acute gastroenteritis in Dhaka City, Bangladesh. <i>Infection, Genetics and Evolution</i> , 2009, 9, 518-522.	1.0	45
97	Molecular analysis of G3 rotavirus among infants and children in Dhaka City, Bangladesh after 1993. <i>Infection, Genetics and Evolution</i> , 2009, 9, 983-986.	1.0	8
98	Detection, genetic characterization, and quantification of norovirus RNA from sera of children with gastroenteritis. <i>Journal of Clinical Virology</i> , 2009, 44, 161-163.	1.6	57
99	Sequence analysis of the capsid gene of Aichi viruses detected from Japan, Bangladesh, Thailand, and Vietnam. <i>Journal of Medical Virology</i> , 2008, 80, 1222-1227.	2.5	21
100	Evaluation of immunochromatography and commercial enzyme-linked immunosorbent assay for rapid detection of norovirus antigen in stool samples. <i>Journal of Virological Methods</i> , 2008, 147, 360-363.	1.0	42
101	Development of a rapid immunochromatographic test for noroviruses genogroups I and II. <i>Journal of Virological Methods</i> , 2008, 148, 1-8.	1.0	44
102	Evidence of Intragenic Recombination in G1 Rotavirus VP7 Genes. <i>Journal of Virology</i> , 2007, 81, 10188-10194.	1.5	44
103	Possible Misidentification of GSP[6] Rotavirus as a Novel Strain Detected in Humans for the First Time. <i>Journal of Clinical Microbiology</i> , 2007, 45, 2098-2099.	1.8	0
104	Isolation and Molecular Characterization of Aichi Viruses from Fecal Specimens Collected in Japan, Bangladesh, Thailand, and Vietnam. <i>Journal of Clinical Microbiology</i> , 2007, 45, 2287-2288.	1.8	99
105	Detection and Genetic Characterization of Group A Rotavirus Strains Circulating among Children with Acute Gastroenteritis in Japan. <i>Journal of Virology</i> , 2007, 81, 4645-4653.	1.5	82
106	Molecular and epidemiological trend of norovirus associated gastroenteritis in Dhaka City, Bangladesh. <i>Journal of Clinical Virology</i> , 2007, 40, 218-223.	1.6	51
107	Molecular characterization of rare G3P[9] rotavirus strains isolated from children hospitalized with acute gastroenteritis. <i>Journal of Medical Virology</i> , 2007, 79, 843-851.	2.5	34
108	Diversity of viruses associated with acute gastroenteritis in children hospitalized with diarrhea in Ho Chi Minh City, Vietnam. <i>Journal of Medical Virology</i> , 2007, 79, 582-590.	2.5	98

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109	Prevalence of sapovirus infection among infants and children with acute gastroenteritis in Dhaka City, Bangladesh during 2004–2005. <i>Journal of Medical Virology</i> , 2007, 79, 633-638.	2.5	29
110	Anti-norovirus polyclonal antibody and its potential for development of an antigen-ELISA. <i>Journal of Medical Virology</i> , 2007, 79, 1180-1186.	2.5	15
111	Sequence analysis of the VP7 gene of human rotavirus G1 isolated in Japan, China, Thailand, and Vietnam in the context of changing distribution of rotavirus G-types. <i>Journal of Medical Virology</i> , 2007, 79, 1009-1016.	2.5	30
112	Genetic heterogeneity, evolution, and recombination in noroviruses. <i>Journal of Medical Virology</i> , 2007, 79, 1388-1400.	2.5	115
113	Amino acid substitutions in the VP7 protein of human rotavirus G3 isolated in China, Russia, Thailand, and Vietnam during 2001–2004. <i>Journal of Medical Virology</i> , 2007, 79, 1611-1616.	2.5	19
114	Sequence analysis of vietnamese P[6] rotavirus strains suggests evidence of interspecies transmission. <i>Journal of Medical Virology</i> , 2007, 79, 1959-1965.	2.5	36
115	Genetic heterogeneity, evolution and recombination in emerging G9 rotaviruses. <i>Infection, Genetics and Evolution</i> , 2007, 7, 656-663.	1.0	85
116	Genetic characterization of group A rotavirus strains circulating among children with acute gastroenteritis in Japan in 2004–2005. <i>Infection, Genetics and Evolution</i> , 2007, 7, 247-253.	1.0	15
117	An outbreak of adenovirus serotype 41 infection in infants and children with acute gastroenteritis in Maizuru City, Japan. <i>Infection, Genetics and Evolution</i> , 2007, 7, 279-284.	1.0	59
118	Emergence of intragenotype recombinant sapovirus in Japan. <i>Infection, Genetics and Evolution</i> , 2007, 7, 542-546.	1.0	21
119	Emergence of rare sapovirus genotype among infants and children with acute gastroenteritis in Japan. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2007, 26, 21-27.	1.3	35
120	Novel Recombinant Norovirus in China. <i>Emerging Infectious Diseases</i> , 2006, 12, 857-858.	2.0	12
121	Novel Recombinant Sapovirus, Japan. <i>Emerging Infectious Diseases</i> , 2006, 12, 865-867.	2.0	10
122	Changing distribution of group A rotavirus G-types and genetic analysis of G9 circulating in Japan. <i>Archives of Virology</i> , 2006, 151, 183-192.	0.9	39
123	Outbreak of sapovirus infection among infants and children with acute gastroenteritis in Osaka City, Japan during 2004–2005. <i>Journal of Medical Virology</i> , 2006, 78, 839-846.	2.5	29
124	Changing distribution of norovirus genotypes and genetic analysis of recombinant GIIb among infants and children with diarrhea in Japan. <i>Journal of Medical Virology</i> , 2006, 78, 971-978.	2.5	91
125	Existence of multiple genotypes associated with acute gastroenteritis during 6-year survey of norovirus infection in Japan. <i>Journal of Medical Virology</i> , 2006, 78, 1318-1324.	2.5	29
126	Detection of Norovirus Antigens from Recombinant Virus-Like Particles and Stool Samples by a Commercial Norovirus Enzyme-Linked Immunosorbent Assay Kit. <i>Journal of Clinical Microbiology</i> , 2006, 44, 3784-3786.	1.8	29

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127	Outbreak of acute gastroenteritis associated with group A rotavirus and genogroup I sapovirus among adults in a mental health care facility in Japan. <i>Journal of Medical Virology</i> , 2005, 75, 475-481.	2.5	24
128	Identification of enteroviral infection among infants and children admitted to hospital with acute gastroenteritis in Ho Chi Minh City, Vietnam. <i>Journal of Medical Virology</i> , 2005, 77, 257-264.	2.5	34
129	Identification of sapovirus infection among Japanese infants in a day care center. <i>Journal of Medical Virology</i> , 2005, 77, 595-601.	2.5	28
130	Genetic diversity of sapovirus in fecal specimens from infants and children with acute gastroenteritis in Pakistan. <i>Archives of Virology</i> , 2005, 150, 371-377.	0.9	20
131	A novel RT-multiplex PCR for enteroviruses, hepatitis A and E viruses and influenza A virus among infants and children with diarrhea in Vietnam. <i>Archives of Virology</i> , 2005, 150, 1175-1185.	0.9	16
132	Etiologic agents of acute gastroenteritis among Japanese infants and children: Virus diversity and genetic analysis of sapovirus. <i>Archives of Virology</i> , 2005, 150, 1415-1424.	0.9	30
133	Existence of multiple outbreaks of viral gastroenteritis among infants in a day care center in Japan. <i>Archives of Virology</i> , 2005, 150, 2061-2075.	0.9	95
134	Molecular Epidemiology of Adenovirus Infection among Pediatric Population with Diarrhea in Asia. <i>Microbiology and Immunology</i> , 2005, 49, 121-128.	0.7	64
135	Characterizations of Adenovirus Type 41 Isolates from Children with Acute Gastroenteritis in Japan, Vietnam, and Korea. <i>Journal of Clinical Microbiology</i> , 2004, 42, 4032-4039.	1.8	44
136	Molecular epidemiology of viral gastroenteritis in Asia. <i>Pediatrics International</i> , 2004, 46, 245-252.	0.2	34
137	Human astrovirus, norovirus (GI, GII), and sapovirus infections in Pakistani children with diarrhea. <i>Journal of Medical Virology</i> , 2004, 73, 256-261.	2.5	80
138	Virus diversity and an outbreak of group C rotavirus among infants and children with diarrhea in Maizuru city, Japan during 2002-2003. <i>Journal of Medical Virology</i> , 2004, 74, 173-179.	2.5	49