

# Tung G Phan

## List of Publications by Year in descending order

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138  
papers

5,770  
citations

76326

40  
h-index

88630

70  
g-index

140  
all docs

140  
docs citations

140  
times ranked

6696  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic diversity and evolution of SARS-CoV-2. <i>Infection, Genetics and Evolution</i> , 2020, 81, 104260.	2.3	498
2	Detection of a novel circovirus PCV3 in pigs with cardiac and multi-systemic inflammation. <i>Virology Journal</i> , 2016, 13, 184.	3.4	309
3	The Fecal Viral Flora of Wild Rodents. <i>PLoS Pathogens</i> , 2011, 7, e1002218.	4.7	304
4	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.	3.4	216
5	Novel coronavirus: From discovery to clinical diagnostics. <i>Infection, Genetics and Evolution</i> , 2020, 79, 104211.	2.3	209
6	Divergent Astrovirus Associated with Neurologic Disease in Cattle. <i>Emerging Infectious Diseases</i> , 2013, 19, 1385-1392.	4.3	155
7	Acute Diarrhea in West African Children: Diverse Enteric Viruses and a Novel Parvovirus Genus. <i>Journal of Virology</i> , 2012, 86, 11024-11030.	3.4	120
8	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.	3.4	116
9	Genetic heterogeneity, evolution, and recombination in noroviruses. <i>Journal of Medical Virology</i> , 2007, 79, 1388-1400.	5.0	115
10	SARS-CoV-2 and COVID-19: A genetic, epidemiological, and evolutionary perspective. <i>Infection, Genetics and Evolution</i> , 2020, 84, 104384.	2.3	115
11	Discovery of a Novel Polyomavirus in Acute Diarrheal Samples from Children. <i>PLoS ONE</i> , 2012, 7, e49449.	2.5	110
12	The Viruses of Wild Pigeon Droppings. <i>PLoS ONE</i> , 2013, 8, e72787.	2.5	108
13	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.	3.9	99
14	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.	5.0	98
15	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.	2.1	95
16	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.	2.4	94
17	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.	5.0	91
18	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.	1.8	89

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19	Genetic heterogeneity, evolution and recombination in emerging G9 rotaviruses. <i>Infection, Genetics and Evolution</i> , 2007, 7, 656-663.	2.3	85
20	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.	3.4	82
21	Human astrovirus, norovirus (GI, GII), and sapovirus infections in Pakistani children with diarrhea. <i>Journal of Medical Virology</i> , 2004, 73, 256-261.	5.0	80
22	A third gyrovirus species in human faeces. <i>Journal of General Virology</i> , 2012, 93, 1356-1361.	2.9	72
23	Genetic Characterization and Classification of Human and Animal Sapoviruses. <i>PLoS ONE</i> , 2016, 11, e0156373.	2.5	71
24	Outbreaks of Neuroinvasive Astrovirus Associated with Encephalomyelitis, Weakness, and Paralysis among Weaned Pigs, Hungary. <i>Emerging Infectious Diseases</i> , 2017, 23, 1982-1993.	4.3	66
25	Molecular Epidemiology of Adenovirus Infection among Pediatric Population with Diarrhea in Asia. <i>Microbiology and Immunology</i> , 2005, 49, 121-128.	1.4	64
26	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.	2.3	59
27	Detection, genetic characterization, and quantification of norovirus RNA from sera of children with gastroenteritis. <i>Journal of Clinical Virology</i> , 2009, 44, 161-163.	3.1	57
28	Molecular and epidemiological trend of norovirus associated gastroenteritis in Dhaka City, Bangladesh. <i>Journal of Clinical Virology</i> , 2007, 40, 218-223.	3.1	51
29	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.	5.0	49
30	A new protoparvovirus in human fecal samples and cutaneous T cell lymphomas (mycosis fungoides). <i>Virology</i> , 2016, 496, 299-305.	2.4	49
31	One year into the pandemic: Short-term evolution of SARS-CoV-2 and emergence of new lineages. <i>Infection, Genetics and Evolution</i> , 2021, 92, 104869.	2.3	49
32	Bufavirus in Feces of Patients with Gastroenteritis, Finland. <i>Emerging Infectious Diseases</i> , 2014, 20, 1077-1079.	4.3	47
33	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.	2.3	45
34	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.	2.5	45
35	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.	3.9	44
36	Evidence of Intragenic Recombination in G1 Rotavirus VP7 Genes. <i>Journal of Virology</i> , 2007, 81, 10188-10194.	3.4	44

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37	Development of a rapid immunochromatographic test for noroviruses genogroups I and II. <i>Journal of Virological Methods</i> , 2008, 148, 1-8.	2.1	44
38	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.	3.4	43
39	Novel Human Adenovirus Strain, Bangladesh. <i>Emerging Infectious Diseases</i> , 2012, 18, 846-848.	4.3	43
40	Cyclovirus in nasopharyngeal aspirates of Chilean children with respiratory infections. <i>Journal of General Virology</i> , 2014, 95, 922-927.	2.9	43
41	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.	2.1	42
42	Genomes of viral isolates derived from different mosquitos species. <i>Virus Research</i> , 2017, 242, 49-57.	2.2	40
43	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.	2.1	39
44	Virome of US bovine calf serum. <i>Biologicals</i> , 2017, 46, 64-67.	1.4	39
45	Genomic characterization of a novel human adenovirus type 31 recombinant in the hexon gene. <i>Journal of General Virology</i> , 2011, 92, 2770-2775.	2.9	37
46	Sequence analysis of vietnamese P[6] rotavirus strains suggests evidence of interspecies transmission. <i>Journal of Medical Virology</i> , 2007, 79, 1959-1965.	5.0	36
47	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.	2.4	36
48	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.	2.1	36
49	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.	2.9	35
50	Molecular epidemiology of viral gastroenteritis in Asia. <i>Pediatrics International</i> , 2004, 46, 245-252.	0.5	34
51	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.	5.0	34
52	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.	5.0	34
53	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
54	New astrovirus in human feces from Burkina Faso. <i>Journal of Clinical Virology</i> , 2014, 60, 161-164.	3.1	32

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55	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.	2.1	30
56	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.	5.0	30
57	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.	5.0	29
58	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.	5.0	29
59	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.	3.9	29
60	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.	5.0	29
61	New Parvovirus in Child with Unexplained Diarrhea, Tunisia. <i>Emerging Infectious Diseases</i> , 2014, 20, 1911-1913.	4.3	29
62	Enteric virome of Ethiopian children participating in a clean water intervention trial. <i>PLoS ONE</i> , 2018, 13, e0202054.	2.5	29
63	Identification of sapovirus infection among Japanese infants in a day care center. <i>Journal of Medical Virology</i> , 2005, 77, 595-601.	5.0	28
64	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.	2.1	27
65	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.	2.4	26
66	A gyrovirus infecting a sea bird. <i>Archives of Virology</i> , 2015, 160, 2105-2109.	2.1	25
67	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.	5.0	24
68	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.	2.3	22
69	A new gyrovirus in human feces. <i>Virus Genes</i> , 2015, 51, 132-135.	1.6	22
70	Sesavirus: prototype of a new parvovirus genus in feces of a sea lion. <i>Virus Genes</i> , 2015, 50, 134-136.	1.6	22
71	Development of a One-Step Qualitative RT-PCR Assay to Detect the SARS-CoV-2 Omicron (B.1.1.529) Variant in Respiratory Specimens. <i>Journal of Clinical Microbiology</i> , 2022, 60, jcm0002422.	3.9	22
72	Emergence of intragenotype recombinant sapovirus in Japan. <i>Infection, Genetics and Evolution</i> , 2007, 7, 542-546.	2.3	21

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73	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.	5.0	21
74	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.	2.1	20
75	Sequence and phylogenetic analysis identifies a putative novel gyrovirus 3 genotype in ferret feces. <i>Virus Genes</i> , 2015, 50, 137-141.	1.6	20
76	Novel picornavirus in domestic rabbits ( <i>Oryctolagus cuniculus</i> var. <i>domestica</i> ). <i>Infection, Genetics and Evolution</i> , 2016, 37, 117-122.	2.3	20
77	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.	5.0	19
78	A highly divergent picornavirus in an amphibian, the smooth newt ( <i>Lissotriton vulgaris</i> ). <i>Journal of General Virology</i> , 2015, 96, 2607-2613.	2.9	19
79	Rotavirus I in feces of a cat with diarrhea. <i>Virus Genes</i> , 2017, 53, 487-490.	1.6	19
80	Sera of Peruvians with fever of unknown origins include viral nucleic acids from non-vertebrate hosts. <i>Virus Genes</i> , 2018, 54, 33-40.	1.6	19
81	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.	2.3	18
82	Viral species richness and composition in young children with loose or watery stool in Ethiopia. <i>BMC Infectious Diseases</i> , 2019, 19, 53.	2.9	18
83	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.	5.0	17
84	First detection of SARS-CoV-2 Omicron BA.4 variant in Western Pennsylvania, United States. <i>Journal of Medical Virology</i> , 2022, 94, 4053-4055.	5.0	17
85	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.	2.1	16
86	Anti-norovirus polyclonal antibody and its potential for development of an antigen-ELISA. <i>Journal of Medical Virology</i> , 2007, 79, 1180-1186.	5.0	15
87	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.	2.3	15
88	A new densovirus in cerebrospinal fluid from a case of anti-NMDA-receptor encephalitis. <i>Archives of Virology</i> , 2016, 161, 3231-3235.	2.1	15
89	Small Circular Rep-Encoding Single-Stranded DNA Genomes in Peruvian Diarrhea Virome. <i>Genome Announcements</i> , 2017, 5, .	0.8	15
90	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.	2.3	14

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91	Nasal virome of dogs with respiratory infection signs include novel taupapillomaviruses. <i>Virus Genes</i> , 2019, 55, 191-197.	1.6	14
92	Molecular and epidemiological trend of rotavirus infection among infants and children in Japan. <i>Infection, Genetics and Evolution</i> , 2009, 9, 955-961.	2.3	13
93	Genome Sequence of an Unusual Human G10P[8] Rotavirus Detected in Vietnam. <i>Journal of Virology</i> , 2012, 86, 10236-10237.	3.4	13
94	Novel Recombinant Norovirus in China. <i>Emerging Infectious Diseases</i> , 2006, 12, 857-858.	4.3	12
95	Molecular and Epidemiological Trend of Sapovirus, and Astrovirus Infection in Japan. <i>Journal of Tropical Pediatrics</i> , 2010, 56, 205-207.	1.5	12
96	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.	3.1	12
97	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.	2.1	12
98	Molecular epidemiology and surveillance of circulating rotavirus among children with gastroenteritis in Bangladesh during 2014â€“2019. <i>PLoS ONE</i> , 2020, 15, e0242813.	2.5	12
99	Emergence of SARSâ€“CoVâ€“2 Omicron BA.5 variant of concern in Western Pennsylvania, United States. <i>Journal of Medical Virology</i> , 2022, 94, 4593-4594.	5.0	12
100	Novel recombinant norovirus in Japan. <i>Virus Genes</i> , 2010, 40, 362-364.	1.6	11
101	Novel Recombinant Sapovirus, Japan. <i>Emerging Infectious Diseases</i> , 2006, 12, 865-867.	4.3	10
102	Rosavirus: the prototype of a proposed new genus of the Picornaviridae family. <i>Virus Genes</i> , 2013, 47, 556-558.	1.6	10
103	Novel Human Gammapapillomavirus Species in a Nasal Swab. <i>Genome Announcements</i> , 2013, 1, e0002213.	0.8	10
104	Absence of giant blood Marseilleâ€“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.	1.6	10
105	A Naturally Transmitted Epitheliotropic Polyomavirus Pathogenic in Immunodeficient Rats: Characterization, Transmission, and Preliminary Epidemiologic Studies. <i>Toxicologic Pathology</i> , 2017, 45, 593-603.	1.8	10
106	<i>Vibrio mimicus</i> wound infection in a burn patient. <i>Radiology Case Reports</i> , 2021, 16, 1348-1351.	0.6	10
107	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.	2.1	9
108	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.	2.1	9

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109	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.	2.3	8
110	Disseminated cryptococcosis in an immunocompetent patient. <i>Respiratory Medicine Case Reports</i> , 2020, 30, 101034.	0.4	8
111	Molecular epidemiology and genetic diversity of norovirus infection in children with acute gastroenteritis in Bangladesh, 2014–2019. <i>Journal of Medical Virology</i> , 2021, 93, 3564-3571.	5.0	8
112	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.	2.1	7
113	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.	2.1	7
114	<i>Acinetobacter junii</i> as a rare pathogen of urinary tract infection. <i>Urology Case Reports</i> , 2020, 32, 101209.	0.3	7
115	Genome characterization of a novel chicken picornavirus distantly related to the members of genus <i>Avihepatovirus</i> with a single 2A protein and a megrivirus-like 3' UTR. <i>Infection, Genetics and Evolution</i> , 2014, 28, 333-338.	2.3	6
116	Cutavirus: A newly discovered parvovirus on the rise. <i>Infection, Genetics and Evolution</i> , 2020, 80, 104175.	2.3	6
117	Whole genome sequencing and evolutionary analysis of G8P [8] rotaviruses emerging in Japan. <i>VirusDisease</i> , 2022, 33, 215-218.	2.0	6
118	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.	1.6	5
119	A rare case of polymicrobial brain abscess involving <i>Actinomyces</i> . <i>Radiology Case Reports</i> , 2021, 16, 1123-1126.	0.6	4
120	Detection of <i>Enterococcus hirae</i> in a case of acute osteomyelitis. <i>Radiology Case Reports</i> , 2021, 16, 2366-2369.	0.6	4
121	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.	2.1	3
122	<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.	1.3	3
123	Evaluation of the Cepheid Xpert Xpress SARS-CoV-2 test for bronchoalveolar lavage. <i>Journal of Clinical Virology Plus</i> , 2022, 2, 100067.	1.0	3
124	Clinical evaluation of the Cue's COVID-19 diagnostic test to detect SARS-CoV-2 in the upper respiratory tract. <i>Journal of Medical Virology</i> , 2022, 94, 3517-3519.	5.0	3
125	Unusual community-associated carbapenem-resistant <i>Acinetobacter baumannii</i> infection, Pennsylvania, USA. <i>IDCases</i> , 2020, 21, e00851.	0.9	2
126	Evaluation of Viral Loads in Patients With SARS-CoV-2 Delta Variant Infection: Higher Loads Do Not Translate Into Different Testing Scenarios. <i>Microbiology Insights</i> , 2022, 15, 117863612210875.	2.0	2



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127	Detection of <i>Enterococcus avium</i> in a case of urinary tract infection and haematuria. <i>Access Microbiology</i> , 2022, 4, .	0.5	2
128	MALDI-TOF vs. VITEK 2 for identification of <i>Aggregatibacter actinomycetemcomitans</i> chest wall abscess. <i>IDCases</i> , 2020, 20, e00749.	0.9	1
129	Genomic characterization of Changuinola viruses from Panama: evidence for multiple genome segment reassortment. <i>Virus Genes</i> , 2020, 56, 527-530.	1.6	1
130	Diagnostic Tests for COVID-19. <i>Advances in Experimental Medicine and Biology</i> , 2021, 1318, 403-412.	1.6	1
131	Tuberculosis of the rare azygos lobe of the right lung. <i>Respiratory Medicine Case Reports</i> , 2021, 33, 101424.	0.4	1
132	Whole genome sequence of an uncommon G9P[4] species A rotavirus containing DS-1-like (genotype 2) genes in Japan. <i>Archives of Virology</i> , 2022, 167, 1603-1606.	2.1	1
133	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.	3.9	0
134	Endophthalmitis resulting from gonococcal keratoconjunctivitis. <i>New Microbes and New Infections</i> , 2020, 36, 100724.	1.6	0
135	Tympanic membrane perforation secondary to <i>Aspergillus niger</i> otomycosis. <i>IDCases</i> , 2020, 22, e00944.	0.9	0
136	Secondary syphilis as an initial presentation of HIV. <i>Baylor University Medical Center Proceedings</i> , 2022, 35, 1-2.	0.5	0
137	Evaluation of the ePlex Respiratory pathogen panel 2 to detect viral and bacterial pathogens, including SARS-CoV-2 Omicron in nasopharyngeal swabs. <i>Journal of Clinical Virology Plus</i> , 2022, 2, 100072.	1.0	0
138	Keratitis caused by <i>Nocardia farcinica</i> in a contact lens wearer. , 2022, , .		0