Kwok-Yong Yuen

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3393051/publications.pdf

Version: 2024-02-01

872 papers

112,666 citations

126 h-index 297 g-index

912 all docs 912 docs citations

times ranked

912

118607 citing authors

#	Article	IF	CITATIONS
1	Clinical Characteristics of Coronavirus Disease 2019 in China. New England Journal of Medicine, 2020, 382, 1708-1720.	27.0	22,372
2	A familial cluster of pneumonia associated with the 2019 novel coronavirus indicating person-to-person transmission: a study of a family cluster. Lancet, The, 2020, 395, 514-523.	13.7	7,120
3	Temporal profiles of viral load in posterior oropharyngeal saliva samples and serum antibody responses during infection by SARS-CoV-2: an observational cohort study. Lancet Infectious Diseases, The, 2020, 20, 565-574.	9.1	2,704
4	Structural and Functional Basis of SARS-CoV-2 Entry by Using Human ACE2. Cell, 2020, 181, 894-904.e9.	28.9	2,443
5	Genomic characterization of the 2019 novel human-pathogenic coronavirus isolated from a patient with atypical pneumonia after visiting Wuhan. Emerging Microbes and Infections, 2020, 9, 221-236.	6.5	2,389
6	Consistent Detection of 2019 Novel Coronavirus in Saliva. Clinical Infectious Diseases, 2020, 71, 841-843.	5.8	1,423
7	Coronaviruses â€" drug discovery and therapeutic options. Nature Reviews Drug Discovery, 2016, 15, 327-347.	46.4	1,365
8	Role of lopinavir/ritonavir in the treatment of SARS: initial virological and clinical findings. Thorax, 2004, 59, 252-256.	5.6	1,361
9	Potent neutralizing antibodies against multiple epitopes on SARS-CoV-2 spike. Nature, 2020, 584, 450-456.	27.8	1,337
10	Severe acute respiratory syndrome coronavirus-like virus in Chinese horseshoe bats. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 14040-14045.	7.1	1,322
11	Characterization and Complete Genome Sequence of a Novel Coronavirus, Coronavirus HKU1, from Patients with Pneumonia. Journal of Virology, 2005, 79, 884-895.	3.4	1,269
12	Discovery of Seven Novel Mammalian and Avian Coronaviruses in the Genus Deltacoronavirus Supports Bat Coronaviruses as the Gene Source of Alphacoronavirus and Betacoronavirus and Avian Coronaviruses as the Gene Source of Gammacoronavirus and Deltacoronavirus. Journal of Virology, 2012, 86, 3995-4008.	3.4	1,244
13	Triple combination of interferon beta-1b, lopinavir–ritonavir, and ribavirin in the treatment of patients admitted to hospital with COVID-19: an open-label, randomised, phase 2 trial. Lancet, The, 2020, 395, 1695-1704.	13.7	1,244
14	Avian Influenza A (H5N1) Infection in Humans. New England Journal of Medicine, 2005, 353, 1374-1385.	27.0	1,235
15	Striking antibody evasion manifested by the Omicron variant of SARS-CoV-2. Nature, 2022, 602, 676-681.	27.8	1,038
16	Severe Acute Respiratory Syndrome Coronavirus as an Agent of Emerging and Reemerging Infection. Clinical Microbiology Reviews, 2007, 20, 660-694.	13.6	886
17	Simulation of the Clinical and Pathological Manifestations of Coronavirus Disease 2019 (COVID-19) in a Golden Syrian Hamster Model: Implications for Disease Pathogenesis and Transmissibility. Clinical Infectious Diseases, 2020, 71, 2428-2446.	5. 8	839
18	Human infections with the emerging avian influenza A H7N9 virus from wet market poultry: clinical analysis and characterisation of viral genome. Lancet, The, 2013, 381, 1916-1925.	13.7	781

#	Article	IF	CITATIONS
19	Improved Molecular Diagnosis of COVID-19 by the Novel, Highly Sensitive and Specific COVID-19-RdRp/Hel Real-Time Reverse Transcription-PCR Assay Validated <i>In Vitro</i> and with Clinical Specimens. Journal of Clinical Microbiology, 2020, 58, .	3.9	780
20	Anti–spike IgG causes severe acute lung injury by skewing macrophage responses during acute SARS-CoV infection. JCI Insight, 2019, 4, .	5.0	742
21	Middle East Respiratory Syndrome Coronavirus: Another Zoonotic Betacoronavirus Causing SARS-Like Disease. Clinical Microbiology Reviews, 2015, 28, 465-522.	13.6	703
22	Comparative tropism, replication kinetics, and cell damage profiling of SARS-CoV-2 and SARS-CoV with implications for clinical manifestations, transmissibility, and laboratory studies of COVID-19: an observational study. Lancet Microbe, The, 2020, 1, e14-e23.	7.3	683
23	Discovery of SARS-CoV-2 antiviral drugs through large-scale compound repurposing. Nature, 2020, 586, 113-119.	27.8	672
24	Coronavirus Genomics and Bioinformatics Analysis. Viruses, 2010, 2, 1804-1820.	3.3	650
25	Antibody evasion properties of SARS-CoV-2 Omicron sublineages. Nature, 2022, 604, 553-556.	27.8	649
26	Coronavirus Disease 2019 (COVID-19) Re-infection by a Phylogenetically Distinct Severe Acute Respiratory Syndrome Coronavirus 2 Strain Confirmed by Whole Genome Sequencing. Clinical Infectious Diseases, 2021, 73, e2946-e2951.	5.8	647
27	The role of community-wide wearing of face mask for control of coronavirus disease 2019 (COVID-19) epidemic due to SARS-CoV-2. Journal of Infection, 2020, 81, 107-114.	3.3	624
28	Convalescent Plasma Treatment Reduced Mortality in Patients With Severe Pandemic Influenza A (H1N1) 2009 Virus Infection. Clinical Infectious Diseases, 2011, 52, 447-456.	5.8	596
29	Treatment With Lopinavir/Ritonavir or Interferon- \hat{l}^21b Improves Outcome of MERS-CoV Infection in a Nonhuman Primate Model of Common Marmoset. Journal of Infectious Diseases, 2015, 212, 1904-1913.	4.0	572
30	Comparative Replication and Immune Activation Profiles of SARS-CoV-2 and SARS-CoV in Human Lungs: An Ex Vivo Study With Implications for the Pathogenesis of COVID-19. Clinical Infectious Diseases, 2020, 71, 1400-1409.	5.8	561
31	Coronavirus Diversity, Phylogeny and Interspecies Jumping. Experimental Biology and Medicine, 2009, 234, 1117-1127.	2.4	548
32	Design of Wide-Spectrum Inhibitors Targeting Coronavirus Main Proteases. PLoS Biology, 2005, 3, e324.	5.6	547
33	Surgical Mask Partition Reduces the Risk of Noncontact Transmission in a Golden Syrian Hamster Model for Coronavirus Disease 2019 (COVID-19). Clinical Infectious Diseases, 2020, 71, 2139-2149.	5.8	501
34	Interspecies transmission and emergence of novel viruses: lessons from bats and birds. Trends in Microbiology, 2013, 21, 544-555.	7.7	461
35	Attenuated replication and pathogenicity of SARS-CoV-2 B.1.1.529 Omicron. Nature, 2022, 603, 693-699.	27.8	460
36	In vitro susceptibility of 10 clinical isolates of SARS coronavirus to selected antiviral compounds. Journal of Clinical Virology, 2004, 31, 69-75.	3.1	459

#	Article	IF	CITATIONS
37	Acute SARS-CoV-2 Infection Impairs Dendritic Cell and T Cell Responses. Immunity, 2020, 53, 864-877.e5.	14.3	450
38	Infection of bat and human intestinal organoids by SARS-CoV-2. Nature Medicine, 2020, 26, 1077-1083.	30.7	441
39	SARS-CoV-2 nsp13, nsp14, nsp15 and orf6 function as potent interferon antagonists. Emerging Microbes and Infections, 2020, 9, 1418-1428.	6.5	439
40	Severe acute respiratory syndrome Coronavirus ORF3a protein activates the NLRP3 inflammasome by promoting TRAF3â€dependent ubiquitination of ASC. FASEB Journal, 2019, 33, 8865-8877.	0.5	434
41	Taxonomy of the order Mononegavirales: update 2016. Archives of Virology, 2016, 161, 2351-2360.	2.1	407
42	Delayed Clearance of Viral Load and Marked Cytokine Activation in Severe Cases of Pandemic H1N1 2009 Influenza Virus Infection. Clinical Infectious Diseases, 2010, 50, 850-859.	5.8	403
43	Cytokine Responses in Severe Acute Respiratory Syndrome Coronavirus-Infected Macrophages In Vitro: Possible Relevance to Pathogenesis. Journal of Virology, 2005, 79, 7819-7826.	3.4	394
44	Middle East Respiratory Syndrome Coronavirus Efficiently Infects Human Primary T Lymphocytes and Activates the Extrinsic and Intrinsic Apoptosis Pathways. Journal of Infectious Diseases, 2016, 213, 904-914.	4.0	379
45	Coronavirus HKU1 and Other Coronavirus Infections in Hong Kong. Journal of Clinical Microbiology, 2006, 44, 2063-2071.	3.9	370
46	Escalating infection control response to the rapidly evolving epidemiology of the coronavirus disease 2019 (COVID-19) due to SARS-CoV-2 in Hong Kong. Infection Control and Hospital Epidemiology, 2020, 41, 493-498.	1.8	370
47	Active Replication of Middle East Respiratory Syndrome Coronavirus and Aberrant Induction of Inflammatory Cytokines and Chemokines in Human Macrophages: Implications for Pathogenesis. Journal of Infectious Diseases, 2014, 209, 1331-1342.	4.0	369
48	Delayed induction of proinflammatory cytokines and suppression of innate antiviral response by the novel Middle East respiratory syndrome coronavirus: implications for pathogenesis and treatment. Journal of General Virology, 2013, 94, 2679-2690.	2.9	347
49	Possible Central Nervous System Infection by SARS Coronavirus. Emerging Infectious Diseases, 2004, 10, 342-344.	4.3	344
50	Structure-based discovery of Middle East respiratory syndrome coronavirus fusion inhibitor. Nature Communications, 2014, 5, 3067.	12.8	324
51	Neutralization of Severe Acute Respiratory Syndrome Coronavirus 2 Omicron Variant by Sera From BNT162b2 or CoronaVac Vaccine Recipients. Clinical Infectious Diseases, 2022, 75, e822-e826.	5.8	322
52	Human intestinal tract serves as an alternative infection route for Middle East respiratory syndrome coronavirus. Science Advances, 2017, 3, eaao4966.	10.3	317
53	Broad-spectrum antivirals for the emerging Middle East respiratory syndrome coronavirus. Journal of Infection, 2013, 67, 606-616.	3.3	314
54	Clinical Features and Complete Genome Characterization of a Distinct Human Rhinovirus (HRV) Genetic Cluster, Probably Representing a Previously Undetected HRV Species, HRV-C, Associated with Acute Respiratory Illness in Children. Journal of Clinical Microbiology, 2007, 45, 3655-3664.	3.9	313

#	Article	IF	Citations
55	SARS-CoV-2 Omicron variant shows less efficient replication and fusion activity when compared with Delta variant in TMPRSS2-expressed cells. Emerging Microbes and Infections, 2022, 11, 277-283.	6.5	308
56	Delayed antiviral plus immunomodulator treatment still reduces mortality in mice infected by high inoculum of influenza A/H5N1 virus. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 8091-8096.	7.1	280
57	Attenuated SARS-CoV-2 variants with deletions at the S1/S2 junction. Emerging Microbes and Infections, 2020, 9, 837-842.	6.5	270
58	Hyperimmune IV Immunoglobulin Treatment. Chest, 2013, 144, 464-473.	0.8	269
59	SARS-CoV-2 infects human neural progenitor cells and brain organoids. Cell Research, 2020, 30, 928-931.	12.0	267
60	Bats as a continuing source of emerging infections in humans. Reviews in Medical Virology, 2007, 17, 67-91.	8.3	266
61	Severe Acute Respiratory Syndrome Coronavirus M Protein Inhibits Type I Interferon Production by Impeding the Formation of TRAF3·TANK·TBK1/IKKϵ Complex. Journal of Biological Chemistry, 2009, 284, 16202-16209.	3.4	261
62	Molecular diversity of coronaviruses in bats. Virology, 2006, 351, 180-187.	2.4	256
63	Bat Origins of MERS-CoV Supported by Bat Coronavirus HKU4ÂUsage of Human Receptor CD26. Cell Host and Microbe, 2014, 16, 328-337.	11.0	252
64	Distribution of Amantadineâ€Resistant H5N1 Avian Influenza Variants in Asia. Journal of Infectious Diseases, 2006, 193, 1626-1629.	4.0	243
65	Ecoepidemiology and Complete Genome Comparison of Different Strains of Severe Acute Respiratory Syndrome-Related <i>Rhinolophus</i> Bat Coronavirus in China Reveal Bats as a Reservoir for Acute, Self-Limiting Infection That Allows Recombination Events. Journal of Virology, 2010, 84, 2808-2819.	3.4	242
66	Avian Influenza Virus Infections in Humans. Chest, 2006, 129, 156-168.	0.8	236
67	Identification of influenza A nucleoprotein as an antiviral target. Nature Biotechnology, 2010, 28, 600-605.	17.5	234
68	Comparative Analysis of Twelve Genomes of Three Novel Group 2c and Group 2d Coronaviruses Reveals Unique Group and Subgroup Features. Journal of Virology, 2007, 81, 1574-1585.	3.4	233
69	Characterization of the Lipidomic Profile of Human Coronavirus-Infected Cells: Implications for Lipid Metabolism Remodeling upon Coronavirus Replication. Viruses, 2019, 11, 73.	3.3	228
70	Genetic Characterization of Betacoronavirus Lineage C Viruses in Bats Reveals Marked Sequence Divergence in the Spike Protein of Pipistrellus Bat Coronavirus HKU5 in Japanese Pipistrelle: Implications for the Origin of the Novel Middle East Respiratory Syndrome Coronavirus. Journal of Virology, 2013, 87, 8638-8650.	3.4	225
71	Immunorestitution Disease Involving the Innate and Adaptive Response. Clinical Infectious Diseases, 2000, 30, 882-892.	5.8	221
72	Clinical and Molecular Epidemiological Features of Coronavirus HKU1–Associated Communityâ€Acquired Pneumonia. Journal of Infectious Diseases, 2005, 192, 1898-1907.	4.0	221

#	Article	IF	CITATIONS
73	Molecular Epidemiology of Human Coronavirus OC43 Reveals Evolution of Different Genotypes over Time and Recent Emergence of a Novel Genotype due to Natural Recombination. Journal of Virology, 2011, 85, 11325-11337.	3.4	218
74	Soluble ACE2-mediated cell entry of SARS-CoV-2 via interaction with proteins related to the renin-angiotensin system. Cell, 2021, 184, 2212-2228.e12.	28.9	216
75	Differential maturation and subcellular localization of severe acute respiratory syndrome coronavirus surface proteins S, M and E. Journal of General Virology, 2005, 86, 1423-1434.	2.9	215
76	Zika fever and congenital Zika syndrome: An unexpected emerging arboviral disease. Journal of Infection, 2016, 72, 507-524.	3.3	215
77	Differentiated human airway organoids to assess infectivity of emerging influenza virus. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 6822-6827.	7.1	215
78	New Hepatitis E Virus Genotype in Camels, the Middle East. Emerging Infectious Diseases, 2014, 20, 1044-1048.	4.3	213
79	Exceptionally Potent Neutralization of Middle East Respiratory Syndrome Coronavirus by Human Monoclonal Antibodies. Journal of Virology, 2014, 88, 7796-7805.	3.4	212
80	Ciprofloxacin Decreased Polyoma BK Virus Load in Patients Who Underwent Allogeneic Hematopoietic Stem Cell Transplantation. Clinical Infectious Diseases, 2005, 40, 528-537.	5.8	207
81	Comparative Analysis of 22 Coronavirus HKU1 Genomes Reveals a Novel Genotype and Evidence of Natural Recombination in Coronavirus HKU1. Journal of Virology, 2006, 80, 7136-7145.	3.4	205
82	Modulation of the Unfolded Protein Response by the Severe Acute Respiratory Syndrome Coronavirus Spike Protein. Journal of Virology, 2006, 80, 9279-9287.	3.4	202
83	Lessons learned 1 year after SARS-CoV-2 emergence leading to COVID-19 pandemic. Emerging Microbes and Infections, 2021, 10, 507-535.	6.5	202
84	<i>Talaromyces (Penicillium) marneffei</i> infection in non-HIV-infected patients. Emerging Microbes and Infections, 2016, 5, 1-9.	6.5	201
85	Viral load in patients infected with pandemic H1N1 2009 influenza A virus. Journal of Medical Virology, 2010, 82, 1-7.	5.0	200
86	Comparative Analysis of Complete Genome Sequences of Three Avian Coronaviruses Reveals a Novel Group 3c Coronavirus. Journal of Virology, 2009, 83, 908-917.	3.4	196
87	Differential Cell Line Susceptibility to the Emerging Novel Human Betacoronavirus 2c EMC/2012: Implications for Disease Pathogenesis and Clinical Manifestation. Journal of Infectious Diseases, 2013, 207, 1743-1752.	4.0	195
88	Early diagnosis of SARS Coronavirus infection by real time RT-PCR. Journal of Clinical Virology, 2003, 28, 233-238.	3.1	194
89	Potent Neutralization of MERS-CoV by Human Neutralizing Monoclonal Antibodies to the Viral Spike Glycoprotein. Science Translational Medicine, 2014, 6, 234ra59.	12.4	194
90	SREBP-dependent lipidomic reprogramming as a broad-spectrum antiviral target. Nature Communications, 2019, 10, 120.	12.8	192

#	Article	IF	CITATIONS
91	Recombinant Modified Vaccinia Virus Ankara Expressing the Spike Glycoprotein of Severe Acute Respiratory Syndrome Coronavirus Induces Protective Neutralizing Antibodies Primarily Targeting the Receptor Binding Region. Journal of Virology, 2005, 79, 2678-2688.	3.4	188
92	Two Years after Pandemic Influenza A/2009/H1N1: What Have We Learned?. Clinical Microbiology Reviews, 2012, 25, 223-263.	13.6	182
93	Antibody responses against SARS coronavirus are correlated with disease outcome of infected individuals. Journal of Medical Virology, 2006, 78, 1-8.	5.0	180
94	Initial viral load and the outcomes of SARS. Cmaj, 2004, 171, 1349-1352.	2.0	179
95	Clinical and Molecular Epidemiology of Human Bocavirus in Respiratory and Fecal Samples from Children in Hong Kong. Journal of Infectious Diseases, 2007, 196, 986-993.	4.0	172
96	Severe Acute Respiratory Syndrome (SARS) Coronavirus ORF8 Protein Is Acquired from SARS-Related Coronavirus from Greater Horseshoe Bats through Recombination. Journal of Virology, 2015, 89, 10532-10547.	3.4	172
97	Middle East Respiratory Syndrome Coronavirus 4a Protein Is a Double-Stranded RNA-Binding Protein That Suppresses PACT-Induced Activation of RIG-I and MDA5 in the Innate Antiviral Response. Journal of Virology, 2014, 88, 4866-4876.	3.4	171
98	Identification of <i>TMPRSS2</i> as a Susceptibility Gene for Severe 2009 Pandemic A(H1N1) Influenza and A(H7N9) Influenza. Journal of Infectious Diseases, 2015, 212, 1214-1221.	4.0	170
99	Rat Hepatitis E Virus as Cause of Persistent Hepatitis after Liver Transplant. Emerging Infectious Diseases, 2018, 24, 2241-2250.	4.3	167
100	Antigenic Crossâ€Reactivity between Severe Acute Respiratory Syndrome–Associated Coronavirus and Human Coronaviruses 229E and OC43. Journal of Infectious Diseases, 2005, 191, 2033-2037.	4.0	165
101	Infectious diseases emerging from Chinese wet-markets: zoonotic origins of severe respiratory viral infections. Current Opinion in Infectious Diseases, 2006, 19, 401-407.	3.1	165
102	Attenuated Interferon and Proinflammatory Response in SARS-CoV-2–Infected Human Dendritic Cells Is Associated With Viral Antagonism of STAT1 Phosphorylation. Journal of Infectious Diseases, 2020, 222, 734-745.	4.0	165
103	Usefulness of the MicroSeq 500 16S Ribosomal DNA-Based Bacterial Identification System for Identification of Clinically Significant Bacterial Isolates with Ambiguous Biochemical Profiles. Journal of Clinical Microbiology, 2003, 41, 1996-2001.	3.9	162
104	Complete genome sequence of bat coronavirus HKU2 from Chinese horseshoe bats revealed a much smaller spike gene with a different evolutionary lineage from the rest of the genome. Virology, 2007, 367, 428-439.	2.4	162
105	Comparative genomic analysis of pre-epidemic and epidemic Zika virus strains for virological factors potentially associated with the rapidly expanding epidemic. Emerging Microbes and Infections, 2016, 5, 1-12.	6.5	162
106	Longitudinal Profile of Immunoglobulin G (IgG), IgM, and IgA Antibodies against the Severe Acute Respiratory Syndrome (SARS) Coronavirus Nucleocapsid Protein in Patients with Pneumonia Due to the SARS Coronavirus. Vaccine Journal, 2004, 11, 665-668.	2.6	158
107	Cross-reactive antibodies in convalescent SARS patients' sera against the emerging novel human coronavirus EMC (2012) by both immunofluorescent and neutralizing antibody tests. Journal of Infection, 2013, 67, 130-140.	3.3	158
108	The K526R substitution in viral protein PB2 enhances the effects of E627K on influenza virus replication. Nature Communications, 2014, 5, 5509.	12.8	155

#	Article	IF	Citations
109	New Hepatitis E Virus Genotype in Bactrian Camels, Xinjiang, China, 2013. Emerging Infectious Diseases, 2016, 22, 2219-2221.	4.3	153
110	Middle East respiratory syndrome coronavirus and bat coronavirus HKU9 both can utilize GRP78 for attachment onto host cells. Journal of Biological Chemistry, 2018, 293, 11709-11726.	3 . 4	153
111	Air and environmental sampling for SARS-CoV-2 around hospitalized patients with coronavirus disease 2019 (COVID-19). Infection Control and Hospital Epidemiology, 2020, 41, 1258-1265.	1.8	153
112	Clofazimine broadly inhibits coronaviruses including SARS-CoV-2. Nature, 2021, 593, 418-423.	27.8	151
113	Feline morbillivirus, a previously undescribed paramyxovirus associated with tubulointerstitial nephritis in domestic cats. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 5435-5440.	7.1	150
114	Productive replication of Middle East respiratory syndrome coronavirus in monocyte-derived dendritic cells modulates innate immune response. Virology, 2014, 454-455, 197-205.	2.4	149
115	Identification of novel porcine and bovine parvoviruses closely related to human parvovirus 4. Journal of General Virology, 2008, 89, 1840-1848.	2.9	148
116	Is the discovery of the novel human betacoronavirus 2c EMC/2012 (HCoV-EMC) the beginning of another SARS-like pandemic?. Journal of Infection, 2012, 65, 477-489.	3.3	147
117	Discovery of a Novel Coronavirus, China Rattus Coronavirus HKU24, from Norway Rats Supports the Murine Origin of Betacoronavirus 1 and Has Implications for the Ancestor of Betacoronavirus Lineage A. Journal of Virology, 2015, 89, 3076-3092.	3.4	147
118	Identification of Novel Small-Molecule Inhibitors of Severe Acute Respiratory Syndrome-Associated Coronavirus by Chemical Genetics. Chemistry and Biology, 2004, 11, 1293-1299.	6.0	141
119	Human enterovirus 71 epidemics: what's next?. Emerging Health Threats Journal, 2013, 6, 19780.	3.0	141
120	MERS coronavirus induces apoptosis in kidney and lung by upregulating Smad7 and FGF2. Nature Microbiology, 2016, 1, 16004.	13.3	140
121	Metallodrug ranitidine bismuth citrate suppresses SARS-CoV-2 replication and relieves virus-associated pneumonia in Syrian hamsters. Nature Microbiology, 2020, 5, 1439-1448.	13.3	140
122	Differential cell line susceptibility to the emerging Zika virus: implications for disease pathogenesis, non-vector-borne human transmission and animal reservoirs. Emerging Microbes and Infections, 2016, 5, 1-12.	6.5	139
123	Adoptive transfer of autologous Epstein-Barr virus-specific cytotoxic T cells for nasopharyngeal carcinoma. International Journal of Cancer, 2001, 94, 73-80.	5.1	137
124	A humanized neutralizing antibody against MERS-CoV targeting the receptor-binding domain of the spike protein. Cell Research, 2015, 25, 1237-1249.	12.0	137
125	Cytotoxic T Lymphocytes Established by Seasonal Human Influenza Cross-React against 2009 Pandemic H1N1 Influenza Virus. Journal of Virology, 2010, 84, 6527-6535.	3.4	136
126	Emergence of scarlet fever Streptococcus pyogenes emm12 clones in Hong Kong is associated with toxin acquisition and multidrug resistance. Nature Genetics, 2015, 47, 84-87.	21.4	135

#	Article	IF	Citations
127	Relative rates of non-pneumonic SARS coronavirus infection and SARS coronavirus pneumonia. Lancet, The, 2004, 363, 841-845.	13.7	134
128	Diagnosis and spectrum of melamine-related renal disease: Plausible mechanism of stone formation in humans. Clinica Chimica Acta, 2009, 402, 150-155.	1.1	133
129	A novel peptide with potent and broad-spectrum antiviral activities against multiple respiratory viruses. Scientific Reports, 2016, 6, 22008.	3.3	133
130	Direct Bacterial Identification in Positive Blood Cultures by Use of Two Commercial Matrix-Assisted Laser Desorption Ionization–Time of Flight Mass Spectrometry Systems. Journal of Clinical Microbiology, 2013, 51, 1733-1739.	3.9	132
131	The emergence of influenza A H7N9 in human beings 16 years after influenza A H5N1: a tale of two cities. Lancet Infectious Diseases, The, 2013, 13, 809-821.	9.1	129
132	High neutralizing antibody titer in intensive care unit patients with COVID-19. Emerging Microbes and Infections, 2020, 9, 1664-1670.	6.5	129
133	Rapid Diagnosis of a Coronavirus Associated with Severe Acute Respiratory Syndrome (SARS). Clinical Chemistry, 2003, 49, 953-955.	3.2	128
134	Emerging SARS-CoV-2 variants expand species tropism to murines. EBioMedicine, 2021, 73, 103643.	6.1	127
135	Invasive Streptococcus iniae Infections Outside North America. Journal of Clinical Microbiology, 2003, 41, 1004-1009.	3.9	126
136	Sensitive and Specific Monoclonal Antibody-Based Capture Enzyme Immunoassay for Detection of Nucleocapsid Antigen in Sera from Patients with Severe Acute Respiratory Syndrome. Journal of Clinical Microbiology, 2004, 42, 2629-2635.	3.9	126
137	Discovery of a Novel Bottlenose Dolphin Coronavirus Reveals a Distinct Species of Marine Mammal Coronavirus in Gammacoronavirus. Journal of Virology, 2014, 88, 1318-1331.	3.4	126
138	Clinical and Molecular Epidemiology of Human Rhinovirus C in Children and Adults in Hong Kong Reveals a Possible Distinct Human Rhinovirus C Subgroup. Journal of Infectious Diseases, 2009, 200, 1096-1103.	4.0	125
139	Cytokine Profiles Induced by the Novel Swineâ€Origin Influenza A/H1N1 Virus: Implications for Treatment Strategies. Journal of Infectious Diseases, 2010, 201, 346-353.	4.0	125
140	The Effect of Pseudomonas aeruginosa Infection on Clinical Parameters in Steady-State Bronchiectasis. Chest, 1998, 114, 1594-1598.	0.8	124
141	Intranasal Vaccination of Recombinant Adeno-Associated Virus Encoding Receptor-Binding Domain of Severe Acute Respiratory Syndrome Coronavirus (SARS-CoV) Spike Protein Induces Strong Mucosal Immune Responses and Provides Long-Term Protection against SARS-CoV Infection. Journal of Immunology. 2008. 180. 948-956.	0.8	124
142	Nucleocapsid Protein as Early Diagnostic Marker for SARS. Emerging Infectious Diseases, 2004, 10, 1947-1949.	4.3	123
143	Clinical management and infection control of SARS: Lessons learned. Antiviral Research, 2013, 100, 407-419.	4.1	122
144	Detection of SARS Coronavirus in Patients with Severe Acute Respiratory Syndrome by Conventional and Real-Time Quantitative Reverse Transcription-PCR Assays. Clinical Chemistry, 2004, 50, 67-72.	3.2	121

#	Article	IF	CITATIONS
145	Serotype 1-Specific Monoclonal Antibody-Based Antigen Capture Immunoassay for Detection of Circulating Nonstructural Protein NS1: Implications for Early Diagnosis and Serotyping of Dengue Virus Infections. Journal of Clinical Microbiology, 2006, 44, 2872-2878.	3.9	121
146	Identification of MicroRNA-Like RNAs in Mycelial and Yeast Phases of the Thermal Dimorphic Fungus Penicillium marneffei. PLoS Neglected Tropical Diseases, 2013, 7, e2398.	3.0	121
147	Oral SARS-CoV-2 Inoculation Establishes Subclinical Respiratory Infection with Virus Shedding in Golden Syrian Hamsters. Cell Reports Medicine, 2020, 1, 100121.	6.5	121
148	Transmission of Rat Hepatitis E Virus Infection to Humans in Hong Kong: A Clinical and Epidemiological Analysis. Hepatology, 2021, 73, 10-22.	7.3	121
149	Molecular diagnostics in tuberculosis. European Journal of Clinical Microbiology and Infectious Diseases, 2005, 24, 711-720.	2.9	120
150	Laribacter hongkongensis gen. nov., sp. nov., a Novel Gram-Negative Bacterium Isolated from a Cirrhotic Patient with Bacteremia and Empyema. Journal of Clinical Microbiology, 2001, 39, 4227-4232.	3.9	119
151	Prevention of Acute Myocardial Infarction and Stroke among Elderly Persons by Dual Pneumococcal and Influenza Vaccination: A Prospective Cohort Study. Clinical Infectious Diseases, 2010, 51, 1007-1016.	5.8	119
152	Differential Sensitivities of Severe Acute Respiratory Syndrome (SARS) Coronavirus Spike Polypeptide Enzyme-Linked Immunosorbent Assay (ELISA) and SARS Coronavirus Nucleocapsid Protein ELISA for Serodiagnosis of SARS Coronavirus Pneumonia. Journal of Clinical Microbiology, 2005, 43, 3054-3058.	3.9	118
153	Eggerthella hongkongensis sp. nov. and eggerthella sinensis sp. nov., two novel Eggerthella species, account for half of the cases of Eggerthella bacteremia. Diagnostic Microbiology and Infectious Disease, 2004, 49, 255-263.	1.8	116
154	Cleavage of spike protein of SARS coronavirus by protease factor Xa is associated with viral infectivity. Biochemical and Biophysical Research Communications, 2007, 359, 174-179.	2.1	116
155	A peptide-based viral inactivator inhibits Zika virus infection in pregnant mice and fetuses. Nature Communications, 2017, 8, 15672.	12.8	115
156	Detection of Specific Antibodies to Severe Acute Respiratory Syndrome (SARS) Coronavirus Nucleocapsid Protein for Serodiagnosis of SARS Coronavirus Pneumonia. Journal of Clinical Microbiology, 2004, 42, 2306-2309.	3.9	114
157	Isolation and Characterization of a Novel Betacoronavirus Subgroup A Coronavirus, Rabbit Coronavirus HKU14, from Domestic Rabbits. Journal of Virology, 2012, 86, 5481-5496.	3.4	114
158	Detection of Severe Acute Respiratory Syndrome (SARS) Coronavirus Nucleocapsid Protein in SARS Patients by Enzyme-Linked Immunosorbent Assay. Journal of Clinical Microbiology, 2004, 42, 2884-2889.	3.9	113
159	Pathogenicity, transmissibility, and fitness of SARS-CoV-2 Omicron in Syrian hamsters. Science, 2022, 377, 428-433.	12.6	113
160	Susceptibility testing of Clostridium difficile against metronidazole and vancomycin by disk diffusion and Etest. Diagnostic Microbiology and Infectious Disease, 1999, 34, 1-6.	1.8	112
161	Host and viral determinants for efficient SARS-CoV-2 infection of the human lung. Nature Communications, 2021, 12, 134.	12.8	112
162	Epidemiology characteristics of human coronaviruses in patients with respiratory infection symptoms and phylogenetic analysis of HCoV-OC43 during 2010-2015 in Guangzhou. PLoS ONE, 2018, 13, e0191789.	2.5	112

#	Article	IF	Citations
163	Rapid Detection of the Severe Acute Respiratory Syndrome (SARS) Coronavirus by a Loop-Mediated Isothermal Amplification Assay. Clinical Chemistry, 2004, 50, 1050-1052.	3.2	111
164	An Animal Model of MERS Produced by Infection of Rhesus Macaques With MERS Coronavirus. Journal of Infectious Diseases, 2014, 209, 236-242.	4.0	111
165	Explicit hypoxia targeting with tumor suppression by creating an "obligate―anaerobic Salmonella Typhimurium strain. Scientific Reports, 2012, 2, 436.	3.3	110
166	Evaluating the use of posterior oropharyngeal saliva in a point-of-care assay for the detection of SARS-CoV-2. Emerging Microbes and Infections, 2020, 9, 1356-1359.	6.5	109
167	Middle East respiratory syndrome coronavirus M protein suppresses type I interferon expression through the inhibition of TBK1-dependent phosphorylation of IRF3. Emerging Microbes and Infections, 2016, 5, 1-9.	6.5	108
168	SARS-CoV-2 shedding and seroconversion among passengers quarantined after disembarking a cruise ship: a case series. Lancet Infectious Diseases, The, 2020, 20, 1051-1060.	9.1	107
169	Emergence in China of human disease due to avian influenza A(H10N8) – Cause for concern?. Journal of Infection, 2014, 68, 205-215.	3.3	106
170	Severe Acute Respiratory Syndrome Coronavirus 2 Infects and Damages the Mature and Immature Olfactory Sensory Neurons of Hamsters. Clinical Infectious Diseases, 2021, 73, e503-e512.	5.8	106
171	Group G Beta-Hemolytic Streptococcal Bacteremia Characterized by 16S Ribosomal RNA Gene Sequencing. Journal of Clinical Microbiology, 2001, 39, 3147-3155.	3.9	105
172	High diversity of polyketide synthase genes and the melanin biosynthesis gene cluster in ⟨i⟩Penicilliumâ€∫marneffei⟨ i⟩. FEBS Journal, 2010, 277, 3750-3758.	4.7	105
173	Coexistence of Different Genotypes in the Same Bat and Serological Characterization of <i>Rousettus</i> Bat Coronavirus HKU9 Belonging to a Novel <i>Betacoronavirus</i> Subgroup. Journal of Virology, 2010, 84, 11385-11394.	3.4	102
174	Clinical, Virological, and Histopathological Manifestations of Fatal Human Infections by Avian Influenza A(H7N9) Virus. Clinical Infectious Diseases, 2013, 57, 1449-1457.	5.8	102
175	Leptin Mediates the Pathogenesis of Severe 2009 Pandemic Influenza A(H1N1) Infection Associated With Cytokine Dysregulation in Mice With Diet-Induced Obesity. Journal of Infectious Diseases, 2013, 207, 1270-1280.	4.0	102
176	Novel antiviral activity and mechanism of bromocriptine as a Zika virus NS2B-NS3 protease inhibitor. Antiviral Research, 2017, 141, 29-37.	4.1	102
177	Additional molecular testing of saliva specimens improves the detection of respiratory viruses. Emerging Microbes and Infections, 2017, 6, 1-7.	6.5	101
178	Avian-Origin Influenza A(H7N9) Infection in Influenza A(H7N9)–Affected Areas of China: A Serological Study. Journal of Infectious Diseases, 2014, 209, 265-269.	4.0	100
179	Mining of epitopes on spike protein of SARS-CoV-2 from COVID-19 patients. Cell Research, 2020, 30, 702-704.	12.0	100
180	Viral Replication in the Nasopharynx Is Associated with Diarrhea in Patients with Severe Acute Respiratory Syndrome. Clinical Infectious Diseases, 2004, 38, 467-475.	5.8	99

#	Article	IF	CITATIONS
181	The Natural Viral Load Profile of Patients With Pandemic 2009 Influenza A(H1N1) and the Effect of Oseltamivir Treatment. Chest, 2010, 137, 759-768.	0.8	99
182	Quasispecies of the D225G Substitution in the Hemagglutinin of Pandemic Influenza A(H1N1) 2009 Virus from Patients with Severe Disease in Hong Kong, China. Journal of Infectious Diseases, 2010, 201, 1517-1521.	4.0	99
183	D225G mutation in hemagglutinin of pandemic influenza H1N1 (2009) virus enhances virulence in mice. Experimental Biology and Medicine, 2010, 235, 981-988.	2.4	99
184	Mutations outside the rifampicin resistance-determining region associated with rifampicin resistance in Mycobacterium tuberculosis. Journal of Antimicrobial Chemotherapy, 2011, 66, 730-733.	3.0	99
185	Structure-based discovery of clinically approved drugs as Zika virus NS2B-NS3 protease inhibitors that potently inhibit Zika virus infection inÂvitro and inÂvivo. Antiviral Research, 2017, 145, 33-43.	4.1	99
186	Comparative Host Gene Transcription by Microarray Analysis Early after Infection of the Huh7 Cell Line by Severe Acute Respiratory Syndrome Coronavirus and Human Coronavirus 229E. Journal of Virology, 2005, 79, 6180-6193.	3 . 4	97
187	Bisphosphonates for advanced prostate cancer. The Cochrane Library, 2006, , CD006250.	2.8	96
188	Efficacy of Clarithromycin-Naproxen-Oseltamivir Combination in the Treatment of Patients Hospitalized for Influenza A(H3N2) Infection. Chest, 2017, 151, 1069-1080.	0.8	95
189	Novel Betacoronavirus in Dromedaries of the Middle East, 2013. Emerging Infectious Diseases, 2014, 20, 560-572.	4.3	94
190	<i>MP1</i> Encodes an Abundant and Highly Antigenic Cell Wall Mannoprotein in the Pathogenic Fungus <i>Penicillium marneffei</i> Infection and Immunity, 1998, 66, 966-973.	2.2	94
191	Genetic relatedness of the novel human group C betacoronavirus to <i>Tylonycteris</i> bat coronavirus HKU4 and <i>Pipistrellus</i> bat coronavirus HKU5. Emerging Microbes and Infections, 2012, 1, 1-5.	6.5	93
192	Waning immune responses against SARS-CoV-2 variants of concern among vaccinees in Hong Kong. EBioMedicine, 2022, 77, 103904.	6.1	93
193	Streptococcus sinensis sp. nov., a Novel Species Isolated from a Patient with Infective Endocarditis. Journal of Clinical Microbiology, 2002, 40, 805-810.	3.9	92
194	Human Parainfluenza Virus 4 Outbreak and the Role of Diagnostic Tests. Journal of Clinical Microbiology, 2005, 43, 4515-4521.	3.9	92
195	Oseltamivir-Resistant Influenza A Pandemic (H1N1) 2009 Virus, Hong Kong, China. Emerging Infectious Diseases, 2009, 15, 1970-1972.	4.3	92
196	Origin, Genetic Diversity, and Evolutionary Dynamics of Novel Porcine Circovirus 3. Advanced Science, 2018, 5, 1800275.	11.2	92
197	Disseminated Penicilliosis, Recurrent Bacteremic Nontyphoidal Salmonellosis, and Burkholderiosis Associated with Acquired Immunodeficiency Due to Autoantibody against Gamma Interferon. Vaccine Journal, 2010, 17, 1132-1138.	3.1	90
198	Molecular Characterization of the 2011 Hong Kong Scarlet Fever Outbreak. Journal of Infectious Diseases, 2012, 206, 341-351.	4.0	89

#	Article	IF	Citations
199	Anaerobic, non-sporulating, Gram-positive bacilli bacteraemia characterized by 16S rRNA gene sequencing. Journal of Medical Microbiology, 2004, 53, 1247-1253.	1.8	88
200	A critical role of IL-17 in modulating the B-cell response during H5N1 influenza virus infection. Cellular and Molecular Immunology, 2011, 8, 462-468.	10.5	88
201	Selective Activation of Type II Interferon Signaling by Zika Virus NS5 Protein. Journal of Virology, 2017, 91, .	3.4	88
202	Epidemiology of Acute Myocarditis/Pericarditis in Hong Kong Adolescents Following Comirnaty Vaccination. Clinical Infectious Diseases, 2022, 75, 673-681.	5.8	88
203	Recent Transmission of a Novel Alphacoronavirus, Bat Coronavirus HKU10, from Leschenault's Rousettes to Pomona Leaf-Nosed Bats: First Evidence of Interspecies Transmission of Coronavirus between Bats of Different Suborders. Journal of Virology, 2012, 86, 11906-11918.	3.4	87
204	Robust SARS-CoV-2 infection in nasal turbinates after treatment with systemic neutralizing antibodies. Cell Host and Microbe, 2021, 29, 551-563.e5.	11.0	87
205	Granulicatella adiacens and Abiotrophia defectiva bacteraemia characterized by 16S rRNA gene sequencing. Journal of Medical Microbiology, 2003, 52, 137-140.	1.8	86
206	SARS Coronavirus Detection Methods. Emerging Infectious Diseases, 2005, 11, 1090-1092.	4.3	86
207	Introduction of an electronic monitoring system for monitoring compliance with Moments 1 and 4 of the WHO "My 5 Moments for Hand Hygiene" methodology. BMC Infectious Diseases, 2011, 11, 151.	2.9	86
208	A broad-spectrum virus- and host-targeting peptide against respiratory viruses including influenza virus and SARS-CoV-2. Nature Communications, 2020, 11, 4252.	12.8	86
209	Seroprevalence of SARS-CoV-2 in Hong Kong and in residents evacuated from Hubei province, China: a multicohort study. Lancet Microbe, The, 2020, 1, e111-e118.	7.3	86
210	Wild Type and Mutant 2009 Pandemic Influenza A (H1N1) Viruses Cause More Severe Disease and Higher Mortality in Pregnant BALB/c Mice. PLoS ONE, 2010, 5, e13757.	2.5	86
211	The management of coronavirus infections with particular reference to SARS. Journal of Antimicrobial Chemotherapy, 2008, 62, 437-441.	3.0	84
212	Coinfection by Severe Acute Respiratory Syndrome Coronavirus 2 and Influenza A(H1N1)pdm09 Virus Enhances the Severity of Pneumonia in Golden Syrian Hamsters. Clinical Infectious Diseases, 2021, 72, e978-e992.	5.8	84
213	Association of Laribacter hongkongensis in community-acquired gastroenteritis with travel and eating fish: a multicentre case-control study. Lancet, The, 2004, 363, 1941-1947.	13.7	83
214	Emergence of enterovirus 71 "double-recombinant―strains belonging to a novel genotype D originating from southern China: first evidence for combination of intratypic and intertypic recombination events in EV71. Archives of Virology, 2010, 155, 1413-1424.	2.1	83
215	Identification and characterization of bocaviruses in cats and dogs reveals a novel feline bocavirus and a novel genetic group of canine bocavirus. Journal of General Virology, 2012, 93, 1573-1582.	2.9	83
216	Olfactory Dysfunction in Coronavirus Disease 2019 Patients: Observational Cohort Study and Systematic Review. Open Forum Infectious Diseases, 2020, 7, ofaa199.	0.9	83

#	Article	IF	CITATIONS
217	High Titer and Avidity of Nonneutralizing Antibodies against Influenza Vaccine Antigen Are Associated with Severe Influenza. Vaccine Journal, 2012, 19, 1012-1018.	3.1	82
218	Potent and protective IGHV3-53/3-66 public antibodies and their shared escape mutant on the spike of SARS-CoV-2. Nature Communications, 2021, 12, 4210.	12.8	82
219	Evidence for <i>Elizabethkingia anophelis</i> Transmission from Mother to Infant, Hong Kong. Emerging Infectious Diseases, 2015, 21, 232-241.	4.3	81
220	Identification of Mycobacterium neoaurumIsolated from a Neutropenic Patient with Catheter-Related Bacteremia by 16S rRNA Sequencing. Journal of Clinical Microbiology, 2000, 38, 3515-3517.	3.9	81
221	Cytosine deamination and selection of CpG suppressed clones are the two major independent biological forces that shape codon usage bias in coronaviruses. Virology, 2007, 369, 431-442.	2.4	80
222	Receptor Usage of a Novel Bat Lineage C Betacoronavirus Reveals Evolution of Middle East Respiratory Syndrome-Related Coronavirus Spike Proteins for Human Dipeptidyl Peptidase 4 Binding. Journal of Infectious Diseases, 2018, 218, 197-207.	4.0	80
223	Complete Genome Analysis of Three Novel Picornaviruses from Diverse Bat Species. Journal of Virology, 2011, 85, 8819-8828.	3.4	79
224	A Functional Variation in CD55 Increases the Severity of 2009 Pandemic H1N1 Influenza A Virus Infection. Journal of Infectious Diseases, 2012, 206, 495-503.	4.0	79
225	Laboratory diagnosis of melioidosis: Past, present and future. Experimental Biology and Medicine, 2015, 240, 742-751.	2.4	79
226	False-Positive Results in a Recombinant Severe Acute Respiratory Syndrome-Associated Coronavirus (SARS-CoV) Nucleocapsid Enzyme-Linked Immunosorbent Assay Due to HCoV-OC43 and HCoV-229E Rectified by Western Blotting with Recombinant SARS-CoV Spike Polypeptide. Journal of Clinical Microbiology, 2004, 42, 5885-5888.	3.9	78
227	Differential susceptibility of different cell lines to swine-origin influenza A H1N1, seasonal human influenza A H1N1, and avian influenza A H5N1 viruses. Journal of Clinical Virology, 2009, 46, 325-330.	3.1	78
228	Parenteral Aminoglycoside Therapy. Drugs, 1994, 47, 902-913.	10.9	77
229	Rediscovery and genomic characterization of bovine astroviruses. Journal of General Virology, 2011, 92, 1888-1898.	2.9	77
230	Immunogenicity of Intradermal Trivalent Influenza Vaccine With Topical Imiquimod: A Double Blind Randomized Controlled Trial. Clinical Infectious Diseases, 2014, 59, 1246-1255.	5.8	77
231	Zika Virus Infection in Dexamethasone-immunosuppressed Mice Demonstrating Disseminated Infection with Multi-organ Involvement Including Orchitis Effectively Treated by Recombinant Type I Interferons. EBioMedicine, 2016, 14, 112-122.	6.1	77
232	Identification by 16S rRNA Gene Sequencing of Lactobacillus salivarius Bacteremic Cholecystitis. Journal of Clinical Microbiology, 2002, 40, 265-267.	3.9	76
233	Molecular epidemiology and household transmission of community-associated methicillin-resistant Staphylococcus aureus in Hong Kong. Diagnostic Microbiology and Infectious Disease, 2007, 57, 145-151.	1.8	76
234	Avian influenza A H5N1 virus: a continuous threat to humans. Emerging Microbes and Infections, 2012, 1, 1-12.	6.5	76

#	Article	IF	CITATIONS
235	Topical imiquimod before intradermal trivalent influenza vaccine for protection against heterologous non-vaccine and antigenically drifted viruses: a single-centre, double-blind, randomised, controlled phase 2b/3 trial. Lancet Infectious Diseases, The, 2016, 16, 209-218.	9.1	75
236	Detection of Specific Antibodies to an Antigenic Mannoprotein for Diagnosis of Penicillium marneffei Penicilliosis. Journal of Clinical Microbiology, 1998, 36, 3028-3031.	3.9	75
237	Relatively Alcohol-Resistant Mycobacteria Are Emerging Pathogens in Patients Receiving Acupuncture Treatment. Journal of Clinical Microbiology, 2002, 40, 1219-1224.	3.9	74
238	Priming with rAAV encoding RBD of SARS-CoV S protein and boosting with RBD-specific peptides for T cell epitopes elevated humoral and cellular immune responses against SARS-CoV infection. Vaccine, 2008, 26, 1644-1651.	3.8	74
239	Advantages of Using Matrix-Assisted Laser Desorption Ionization-Time of Flight Mass Spectrometry as a Rapid Diagnostic Tool for Identification of Yeasts and Mycobacteria in the Clinical Microbiological Laboratory. Journal of Clinical Microbiology, 2013, 51, 3981-3987.	3.9	74
240	A sensitive and specific antigen detection assay for Middle East respiratory syndrome coronavirus. Emerging Microbes and Infections, 2015, 4, 1-5.	6.5	74
241	Cross-species transmission and emergence of novel viruses from birds. Current Opinion in Virology, 2015, 10, 63-69.	5.4	74
242	Development of a Novel, Genome Subtraction-Derived, SARS-CoV-2-Specific COVID-19-nsp2 Real-Time RT-PCR Assay and Its Evaluation Using Clinical Specimens. International Journal of Molecular Sciences, 2020, 21, 2574.	4.1	74
243	Development and Evaluation of Novel Real-Time Reverse Transcription-PCR Assays with Locked Nucleic Acid Probes Targeting Leader Sequences of Human-Pathogenic Coronaviruses. Journal of Clinical Microbiology, 2015, 53, 2722-2726.	3.9	73
244	The biosynthetic pathway for a thousand-year-old natural food colorant and citrinin in Penicillium marneffei. Scientific Reports, 2014, 4, 6728.	3.3	73
245	The hemagglutinin structure of an avian H1N1 influenza A virus. Virology, 2009, 392, 73-81.	2.4	72
246	Detection of Cell Wall Mannoprotein Mp1p in Culture Supernatants of <i>Penicillium marneffei</i> and in Sera of Penicilliosis Patients. Journal of Clinical Microbiology, 1999, 37, 981-986.	3.9	72
247	Striking antibody evasion manifested by the Omicron variant of SARS-CoV-2. Nature, 0, , .	27.8	72
248	Relationship of Pretransplantation Polyoma BK Virus Serologic Findings and BK Viral Reactivation after Hematopoietic Stem Cell Transplantation. Clinical Infectious Diseases, 2007, 44, 830-837.	5.8	71
249	Dose sparing intradermal trivalent influenza (2010/2011) vaccination overcomes reduced immunogenicity of the 2009 H1N1 strain. Vaccine, 2012, 30, 6427-6435.	3.8	71
250	Misidentification of Aspergillus nomius and Aspergillus tamarii as Aspergillus flavus: Characterization by Internal Transcribed Spacer, β-Tubulin, and Calmodulin Gene Sequencing, Metabolic Fingerprinting, and Matrix-Assisted Laser Desorption Ionization–Time of Flight Mass Spectrometry. Journal of Clinical Microbiology, 2014, 52, 1153-1160.	3.9	71
251	Effect of Clinical and Virological Parameters on the Level of Neutralizing Antibody against Pandemic Influenza A Virus H1N1 2009. Clinical Infectious Diseases, 2010, 51, 274-279.	5.8	70
252	Middle East respiratory syndrome coronavirus infection: virus-host cell interactions and implications on pathogenesis. Virology Journal, 2015, 12, 218.	3.4	70

#	Article	IF	CITATIONS
253	<i>Streptococcus pyogenes</i> and re-emergence of scarlet fever as a public health problem. Emerging Microbes and Infections, 2012, 1, 1-10.	6.5	69
254	Hand-touch contact assessment of high-touch and mutual-touch surfaces among healthcare workers, patients, and visitors. Journal of Hospital Infection, 2015, 90, 220-225.	2.9	69
255	Identification of Arcobacter cryaerophilus isolated from a traffic accident victim with bacteremia by 16S ribosomal RNA gene sequencing. Diagnostic Microbiology and Infectious Disease, 2001, 40, 125-127.	1.8	68
256	Analysis of Melamine Cyanurate in Urine Using Matrix-Assisted Laser Desorption/Ionization Mass Spectrometry. Analytical Chemistry, 2009, 81, 3676-3682.	6.5	68
257	Clinical Spectrum of Exophiala Infections and a Novel Exophiala Species, Exophiala hongkongensis. Journal of Clinical Microbiology, 2013, 51, 260-267.	3.9	68
258	Carcinoembryonic Antigen-Related Cell Adhesion Molecule 5 Is an Important Surface Attachment Factor That Facilitates Entry of Middle East Respiratory Syndrome Coronavirus. Journal of Virology, 2016, 90, 9114-9127.	3.4	68
259	Nocardiosis in systemic lupus erythematosus. Seminars in Arthritis and Rheumatism, 1997, 26, 675-683.	3.4	67
260	Nasopharyngeal Carriage of Antimicrobial-Resistant Streptococcus pneumoniae among Young Children Attending 79 Kindergartens and Day Care Centers in Hong Kong. Antimicrobial Agents and Chemotherapy, 2001, 45, 2765-2770.	3.2	67
261	Clinical Role of ??-Lactam/??-Lactamase Inhibitor Combinations. Drugs, 2003, 63, 1511-1524.	10.9	67
262	Clinical isolates of Streptococcus iniae from Asia are more mucoid and \hat{l}^2 -hemolytic than those from North America. Diagnostic Microbiology and Infectious Disease, 2006, 54, 177-181.	1.8	67
263	Clinical and Virological Factors Associated with Viremia in Pandemic Influenza A/H1N1/2009 Virus Infection. PLoS ONE, 2011, 6, e22534.	2.5	67
264	Geographical difference of disease association in Streptococcus bovis bacteraemia. Journal of Medical Microbiology, 2003, 52, 903-908.	1.8	66
265	Viral Load Distribution in SARS Outbreak. Emerging Infectious Diseases, 2005, 11, 1882-1886.	4.3	66
266	SARS Coronavirus Detection Methods. Emerging Infectious Diseases, 2005, 11, 1108-1111.	4.3	66
267	Rapid Spread of Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) Omicron Subvariant BA.2 in a Single-Source Community Outbreak. Clinical Infectious Diseases, 2022, 75, e44-e49.	5.8	66
268	Catheter-Related Microbacterium Bacteremia Identified by 16S rRNA Gene Sequencing. Journal of Clinical Microbiology, 2002, 40, 2681-2685.	3.9	65
269	Natural Occurrence and Characterization of Two Internal Ribosome Entry Site Elements in a Novel Virus, Canine Picodicistrovirus, in the Picornavirus-Like Superfamily. Journal of Virology, 2012, 86, 2797-2808.	3.4	65
270	Metagenomic analysis of viromes of dromedary camel fecal samples reveals large number and high diversity of circoviruses and picobirnaviruses. Virology, 2014, 471-473, 117-125.	2.4	65

#	Article	IF	CITATIONS
271	Hospital Outbreak of Pulmonary and Cutaneous Zygomycosis due to Contaminated Linen Items From Substandard Laundry. Clinical Infectious Diseases, 2016, 62, 714-721.	5.8	65
272	"streptococcus milleri―endocarditis caused by S treptococcus anginosus. Diagnostic Microbiology and Infectious Disease, 2004, 48, 81-88.	1.8	64
273	Accelerated Evolutionary Rate May Be Responsible for the Emergence of Lineage-Specific Genes in Ascomycota. Journal of Molecular Evolution, 2006, 63, 1-11.	1.8	64
274	Systemic infection of avian influenza A virus H5N1 subtype in humans. Human Pathology, 2009, 40, 735-739.	2.0	64
275	Cellular response to influenza virus infection: a potential role for autophagy in CXCL10 and interferon-alpha induction. Cellular and Molecular Immunology, 2010, 7, 263-270.	10.5	64
276	Rhinovirus – From bench to bedside. Journal of the Formosan Medical Association, 2017, 116, 496-504.	1.7	64
277	Characterization of AFMP1 : a Novel Target for Serodiagnosis of Aspergillosis. Journal of Clinical Microbiology, 2001, 39, 3830-3837.	3.9	63
278	96 weeks combination of adefovir dipivoxil plus emtricitabine vs. adefovir dipivoxil monotherapy in the treatment of chronic hepatitis B. Journal of Hepatology, 2008, 48, 714-720.	3.7	63
279	Anti–IFN-γ autoantibodies are strongly associated with HLA-DR*15:02/16:02 and HLA-DQ*05:01/05:02 across Southeast Asia. Journal of Allergy and Clinical Immunology, 2016, 137, 945-948.e8.	2.9	63
280	Dissemination of the mcr-1 colistin resistance gene. Lancet Infectious Diseases, The, 2016, 16, 145-146.	9.1	63
281	Dual-functional peptide with defective interfering genes effectively protects mice against avian and seasonal influenza. Nature Communications, 2018, 9, 2358.	12.8	63
282	From SARS coronavirus to novel animal and human coronaviruses. Journal of Thoracic Disease, 2013, 5 Suppl 2, S103-8.	1.4	63
283	Synthetic Peptides outside the Spike Protein Heptad Repeat Regions as Potent Inhibitors of Sars-Associated Coronavirus. Antiviral Therapy, 2005, 10, 393-403.	1.0	63
284	Clinical Significance of Alimentary Tract Microbes in Bone Marrow Transplant Recipients. Diagnostic Microbiology and Infectious Disease, 1998, 30, 75-81.	1.8	62
285	Complete Genome Sequence of <i>Staphylococcus lugdunensis</i> Strain HKU09-01. Journal of Bacteriology, 2010, 192, 1471-1472.	2.2	62
286	Disseminated Infections with <i>Talaromyces marneffei </i> in Non-AIDS Patients Given Monoclonal Antibodies against CD20 and Kinase Inhibitors. Emerging Infectious Diseases, 2015, 21, 1101-1106.	4.3	62
287	Discovery and Sequence Analysis of Four Deltacoronaviruses from Birds in the Middle East Reveal Interspecies Jumping with Recombination as a Potential Mechanism for Avian-to-Avian and Avian-to-Mammalian Transmission. Journal of Virology, 2018, 92, .	3.4	62
288	Natural Transmission of Bat-like Severe Acute Respiratory Syndrome Coronavirus 2 Without Proline-Arginine-Arginine-Alanine Variants in Coronavirus Disease 2019 Patients. Clinical Infectious Diseases, 2021, 73, e437-e444.	5.8	62

#	Article	IF	CITATIONS
289	An orally available Mpro inhibitor is effective against wild-type SARS-CoV-2 and variants including Omicron. Nature Microbiology, 2022, 7, 716-725.	13.3	62
290	Accurate Diagnosis of COVID-19 by a Novel Immunogenic Secreted SARS-CoV-2 orf8 Protein. MBio, 2020, 11, .	4.1	61
291	Genomic and experimental evidence for a potential sexual cycle in the pathogenic thermal dimorphic fungusPenicillium marneffei. FEBS Letters, 2006, 580, 3409-3416.	2.8	60
292	Catabacter hongkongensis gen. nov., sp. nov., Isolated from Blood Cultures of Patients from Hong Kong and Canada. Journal of Clinical Microbiology, 2007, 45, 395-401.	3.9	60
293	Small Interfering RNA Targeting M2 Gene Induces Effective and Long Term Inhibition of Influenza A Virus Replication. PLoS ONE, 2009, 4, e5671.	2.5	60
294	Transmission of Omicron (B.1.1.529) - SARS-CoV-2 Variant of Concern in a designated quarantine hotel for travelers: a challenge of elimination strategy of COVID-19. The Lancet Regional Health - Western Pacific, 2022, 18, 100360.	2.9	60
295	Clinical Evaluation of the New High-Throughput Luminex NxTAG Respiratory Pathogen Panel Assay for Multiplex Respiratory Pathogen Detection. Journal of Clinical Microbiology, 2016, 54, 1820-1825.	3.9	59
296	Mycophenolic acid, an immunomodulator, has potent and broad-spectrum in vitro antiviral activity against pandemic, seasonal and avian influenza viruses affecting humans. Journal of General Virology, 2016, 97, 1807-1817.	2.9	59
297	Co-existence of multiple strains of two novel porcine bocaviruses in the same pig, a previously undescribed phenomenon in members of the family Parvoviridae, and evidence for inter- and intra-host genetic diversity and recombination. Journal of General Virology, 2011, 92, 2047-2059.	2.9	59
298	Unique immunogenicity of hepatitis B virus DNA vaccine presented by live-attenuated Salmonella typhimurium. Vaccine, 2001, 19, 2945-2954.	3.8	58
299	Identification by 16S ribosomal RNA gene sequencing of an Enterobacteriaceae species with ambiguous biochemical profile from a renal transplant recipient. Diagnostic Microbiology and Infectious Disease, 2001, 39, 85-93.	1.8	58
300	The Lower Serum Immunoglobulin G2 Level in Severe Cases than in Mild Cases of Pandemic H1N1 2009 Influenza Is Associated with Cytokine Dysregulation. Vaccine Journal, 2011, 18, 305-310.	3.1	58
301	A Novel Dirofilaria Species Causing Human and Canine Infections in Hong Kong. Journal of Clinical Microbiology, 2012, 50, 3534-3541.	3.9	58
302	Competing endogenous RNA network profiling reveals novel host dependency factors required for MERS-CoV propagation. Emerging Microbes and Infections, 2020, 9, 733-746.	6.5	58
303	Intravenous Injection of Coronavirus Disease 2019 (COVID-19) mRNA Vaccine Can Induce Acute Myopericarditis in Mouse Model. Clinical Infectious Diseases, 2022, 74, 1933-1950.	5.8	58
304	Detection of Antibodies Specific to an Antigenic Cell Wall Galactomannoprotein for Serodiagnosis of Aspergillus fumigatus Aspergillosis. Journal of Clinical Microbiology, 2002, 40, 2041-2045.	3.9	57
305	Identification and complete genome analysis of three novel paramyxoviruses, Tuhoko virus $1,2$ and 3 , in fruit bats from China. Virology, $2010,404,106-116$.	2.4	57
306	Molecular Characterization of Fluoroquinolone Resistance in <i>Mycobacterium tuberculosis</i> Functional Analysis of <i>gyrA</i> Mutation at Position 74. Antimicrobial Agents and Chemotherapy, 2011, 55, 608-614.	3.2	57

#	Article	IF	CITATIONS
307	Differences in Antibody Responses of Individuals with Natural Infection and Those Vaccinated against Pandemic H1N1 2009 Influenza. Vaccine Journal, 2011, 18, 867-873.	3.1	57
308	Transfer of scarlet fever-associated elements into the group A Streptococcus M1T1 clone. Scientific Reports, 2015, 5, 15877.	3.3	57
309	PD1-based DNA vaccine amplifies HIV-1 GAG-specific CD8+ T cells in mice. Journal of Clinical Investigation, 2013, 123, 2629-2642.	8.2	57
310	Omicron variant susceptibility to neutralizing antibodies induced in children by natural SARS-CoV-2 infection or COVID-19 vaccine. Emerging Microbes and Infections, 2022, 11, 543-547.	6.5	57
311	The mitochondrial genome of the thermal dimorphic fungusPenicillium marneffeiis more closely related to those of molds than yeasts. FEBS Letters, 2003, 555, 469-477.	2.8	56
312	The NS1 Protein of Influenza A Virus Interacts with Cellular Processing Bodies and Stress Granules through RNA-Associated Protein 55 (RAP55) during Virus Infection. Journal of Virology, 2012, 86, 12695-12707.	3.4	56
313	Discovery of the FDA-approved drugs bexarotene, cetilistat, diiodohydroxyquinoline, and abiraterone as potential COVID-19 treatments with a robust two-tier screening system. Pharmacological Research, 2020, 159, 104960.	7.1	56
314	Therapeutic efficacy of hepatitis B surface antigen–antibodies-recombinant DNA composite in HBsAg transgenic mice. Vaccine, 2001, 19, 4219-4225.	3.8	55
315	Concurrent comparison of epidemiology, clinical presentation and outcome between adult patients suffering from the pandemic influenza A (H1N1) 2009 virus and the seasonal influenza A virus infection. Postgraduate Medical Journal, 2010, 86, 515-521.	1.8	55
316	Identification of specific metabolites in culture supernatant of <i>Mycobacterium tuberculosis </i> busing metabolomics: exploration of potential biomarkers. Emerging Microbes and Infections, 2015, 4, 1-10.	6.5	55
317	Broad-Spectrum Host-Based Antivirals Targeting the Interferon and Lipogenesis Pathways as Potential Treatment Options for the Pandemic Coronavirus Disease 2019 (COVID-19). Viruses, 2020, 12, 628.	3.3	55
318	<i>Lasiodiplodia theobromae</i> Pneumonia in a Liver Transplant Recipient. Journal of Clinical Microbiology, 2008, 46, 380-384.	3.9	54
319	Comparative analysis of six genome sequences of three novel picornaviruses, turdiviruses 1, 2 and 3, in dead wild birds, and proposal of two novel genera, Orthoturdivirus and Paraturdivirus, in the family Picornaviridae. Journal of General Virology, 2010, 91, 2433-2448.	2.9	54
320	First Discovery of Two Polyketide Synthase Genes for Mitorubrinic Acid and Mitorubrinol Yellow Pigment Biosynthesis and Implications in Virulence of Penicillium marneffei. PLoS Neglected Tropical Diseases, 2012, 6, e1871.	3.0	54
321	Novel Partitivirus Enhances Virulence of and Causes Aberrant Gene Expression in Talaromyces marneffei. MBio, 2018, 9, .	4.1	54
322	Tsukamurella Conjunctivitis: a Novel Clinical Syndrome. Journal of Clinical Microbiology, 2003, 41, 3368-3371.	3.9	53
323	SARS coronavirus spike polypeptide DNA vaccine priming with recombinant spike polypeptide from Escherichia coli as booster induces high titer of neutralizing antibody against SARS coronavirus. Vaccine, 2005, 23, 4959-4968.	3.8	53
324	Reactive and Infective Dermatoses Associated with Adult-Onset Immunodeficiency due to Anti-Interferon-Gamma Autoantibody: Sweet's Syndrome and Beyond. Dermatology, 2013, 226, 157-166.	2.1	53

#	Article	IF	Citations
325	Broad-spectrum inhibition of common respiratory RNA viruses by a pyrimidine synthesis inhibitor with involvement of the host antiviral response. Journal of General Virology, 2017, 98, 946-954.	2.9	53
326	Antibiotics Modulate Vaccine-Induced Humoral Immune Response. Vaccine Journal, 1999, 6, 832-837.	2.6	52
327	Acupuncture Mycobacteriosis. New England Journal of Medicine, 2001, 345, 842-843.	27.0	52
328	Laribacter hongkongensis: a potential cause of infectious diarrhea. Diagnostic Microbiology and Infectious Disease, 2003, 47, 551-556.	1.8	52
329	A Patient with Asymptomatic Severe Acute Respiratory Syndrome (SARS) and Antigenemia from the 2003–2004 Community Outbreak of SARS in Guangzhou, China. Clinical Infectious Diseases, 2006, 43, e1-e5.	5.8	52
330	The Complete Genome and Proteome of Laribacter hongkongensis Reveal Potential Mechanisms for Adaptations to Different Temperatures and Habitats. PLoS Genetics, 2009, 5, e1000416.	3.5	52
331	A global call for talaromycosis to be recognised as a neglected tropical disease. The Lancet Global Health, 2021, 9, e1618-e1622.	6.3	52
332	A one step quantitative RT-PCR for detection of SARS coronavirus with an internal control for PCR inhibitors. Journal of Clinical Virology, 2004, 30, 214-217.	3.1	51
333	More and More Coronaviruses: Human Coronavirus HKU1. Viruses, 2009, 1, 57-71.	3.3	51
334	Clostridium difficile ribotype 027 arrives in Hong Kong. International Journal of Antimicrobial Agents, 2009, 34, 492-493.	2.5	51
335	Internal Transcribed Spacer Region Sequence Heterogeneity in <i>Rhizopus microsporus</i> implications for Molecular Diagnosis in Clinical Microbiology Laboratories. Journal of Clinical Microbiology, 2010, 48, 208-214.	3.9	51
336	Clostridium difficile isolates with increased sporulation: emergence of PCR ribotype 002 in Hong Kong. European Journal of Clinical Microbiology and Infectious Diseases, 2011, 30, 1371-81.	2.9	51
337	A Unique and Conserved Neutralization Epitope in H5N1 Influenza Viruses Identified by an Antibody against the A/Goose/Guangdong/1/96 Hemagglutinin. Journal of Virology, 2013, 87, 12619-12635.	3.4	51
338	Epidemiology of human parechovirus, Aichi virus and salivirus in fecal samples from hospitalized children with gastroenteritis in Hong Kong. Virology Journal, 2014, 11, 182.	3.4	51
339	An NS-segment exonic splicing enhancer regulates influenza A virus replication in mammalian cells. Nature Communications, 2017, 8, 14751.	12.8	51
340	Generation of DelNS1 Influenza Viruses: a Strategy for Optimizing Live Attenuated Influenza Vaccines. MBio, 2019, 10, .	4.1	51
341	Animal models in SARS-CoV-2 research. Nature Methods, 2022, 19, 392-394.	19.0	51
342	Bacteremia Due to <i>Clostridium hathewayi</i> in a Patient with Acute Appendicitis. Journal of Clinical Microbiology, 2004, 42, 5947-5949.	3.9	50

#	Article	IF	Citations
343	Ecoepidemiology of Laribacter hongkongensis , a Novel Bacterium Associated with Gastroenteritis. Journal of Clinical Microbiology, 2005, 43, 919-922.	3.9	50
344	Matrix-Assisted Laser Desorption Ionization–Time of Flight Mass Spectrometry for Rapid Identification of Burkholderia pseudomallei: Importance of Expanding Databases with Pathogens Endemic to Different Localities. Journal of Clinical Microbiology, 2012, 50, 3142-3143.	3.9	50
345	Serum Antibody Profile of a Patient With Coronavirus Disease 2019 Reinfection. Clinical Infectious Diseases, 2021, 72, e659-e662.	5.8	50
346	Avian influenza virus A H7N9 infects multiple mononuclear cell types in peripheral blood and induces dysregulated cytokine responses and apoptosis in infected monocytes. Journal of General Virology, 2017, 98, 922-934.	2.9	49
347	Complete Genome Sequence of a Novel Paramyxovirus, Tailam Virus, Discovered in Sikkim Rats. Journal of Virology, 2011, 85, 13473-13474.	3.4	48
348	A Novel Psittacine Adenovirus Identified During an Outbreak of Avian Chlamydiosis and Human Psittacosis: Zoonosis Associated with Virus-Bacterium Coinfection in Birds. PLoS Neglected Tropical Diseases, 2014, 8, e3318.	3.0	48
349	Unexpectedly Higher Morbidity and Mortality of Hospitalized Elderly Patients Associated with Rhinovirus Compared with Influenza Virus Respiratory Tract Infection. International Journal of Molecular Sciences, 2017, 18, 259.	4.1	48
350	Beneficial effect of combinational methylprednisolone and remdesivir in hamster model of SARS-CoV-2 infection. Emerging Microbes and Infections, 2021, 10, 291-304.	6.5	48
351	Clinical Characteristics and Transmission of COVID-19 in Children and Youths During 3 Waves of Outbreaks in Hong Kong. JAMA Network Open, 2021, 4, e218824.	5.9	48
352	Targeting highly pathogenic coronavirus-induced apoptosis reduces viral pathogenesis and disease severity. Science Advances, 2021, 7, .	10.3	48
353	Bacteremia due to Staphylococcus aureus with reduced susceptibility to vancomycin. Diagnostic Microbiology and Infectious Disease, 2000, 36, 261-268.	1.8	47
354	Characterization of SARS-CoV main protease and identification of biologically active small molecule inhibitors using a continuous fluorescence-based assay. FEBS Letters, 2004, 576, 325-330.	2.8	47
355	Guidelines for interpretation of 16S rRNA gene sequence-based results for identification of medically important aerobic Gram-positive bacteria. Journal of Medical Microbiology, 2009, 58, 1030-1036.	1.8	47
356	Proteome profiling of the dimorphic fungus <i>PenicilliumÂmarneffei </i> extracellular proteins and identification of glyceraldehyde-3-phosphate dehydrogenase as an important adhesion factor for conidial attachment. FEBS Journal, 2013, 280, 6613-6626.	4.7	47
357	Surfactant Protein B Gene Polymorphism Is Associated With Severe Influenza. Chest, 2014, 145, 1237-1243.	0.8	47
358	Toll-Like Receptor 7 Agonist Imiquimod in Combination with Influenza Vaccine Expedites and Augments Humoral Immune Responses against Influenza A(H1N1)pdm09 Virus Infection in BALB/c Mice. Vaccine Journal, 2014, 21, 570-579.	3.1	47
359	Isolation and Characterization of a Salmonella enterica Serotype Typhi Variant and Its Clinical and Public Health Implications. Journal of Clinical Microbiology, 2001, 39, 1190-1194.	3.9	46
360	Lipid metabolites as potential diagnostic and prognostic biomarkers for acute community acquired pneumonia. Diagnostic Microbiology and Infectious Disease, 2016, 85, 249-254.	1.8	46

#	Article	IF	CITATIONS
361	Immunization With a Novel Human Type 5 Adenovirus-Vectored Vaccine Expressing the Premembrane and Envelope Proteins of Zika Virus Provides Consistent and Sterilizing Protection in Multiple Immunocompetent and Immunocompromised Animal Models. Journal of Infectious Diseases, 2018, 218, 365-377.	4.0	46
362	Human coronavirus dependency on host heat shock protein 90 reveals an antiviral target. Emerging Microbes and Infections, 2020, 9, 2663-2672.	6.5	46
363	Cell-wall-deficient bacteria and culture-negative febrile episodes in bone-marrow-transplant recipients. Lancet, The, 2001, 357, 675-679.	13.7	45
364	groEL Encodes a Highly Antigenic Protein in Burkholderia pseudomallei. Vaccine Journal, 2001, 8, 832-836.	2.6	45
365	Single gene target bacterial identification. Diagnostic Microbiology and Infectious Disease, 2002, 44, 143-149.	1.8	45
366	Exploring the Penicillium marneffei genome. Archives of Microbiology, 2003, 179, 339-353.	2.2	45
367	Clinical and Molecular Epidemiology of Human Parainfluenza Virus 4 Infections in Hong Kong: Subtype 4B as Common as Subtype 4A. Journal of Clinical Microbiology, 2009, 47, 1549-1552.	3.9	45
368	Novel Pan-Genomic Analysis Approach in Target Selection for Multiplex PCR Identification and Detection of Burkholderia pseudomallei, Burkholderia thailandensis, and Burkholderia cepacia Complex Species: a Proof-of-Concept Study. Journal of Clinical Microbiology, 2011, 49, 814-821.	3.9	45
369	Analytical Sensitivity of Seven Point-of-Care Influenza Virus Detection Tests and Two Molecular Tests for Detection of Avian Origin H7N9 and Swine Origin H3N2 Variant Influenza A Viruses. Journal of Clinical Microbiology, 2013, 51, 3160-3161.	3.9	45
370	Evaluation of the commercially available LightMix® Modular E-gene kit using clinical and proficiency testing specimens for SARS-CoV-2 detection. Journal of Clinical Virology, 2020, 129, 104476.	3.1	45
371	Absence of nosocomial influenza and respiratory syncytial virus infection in the coronavirus disease 2019 (COVID-19) era: Implication of universal masking in hospitals. Infection Control and Hospital Epidemiology, 2021, 42, 218-221.	1.8	45
372	Hepatitis E Virus Species C Infection in Humans, Hong Kong. Clinical Infectious Diseases, 2022, 75, 288-296.	5.8	45
373	Contribution of low population immunity to the severe Omicron BA.2 outbreak in Hong Kong. Nature Communications, 2022, 13 , .	12.8	45
374	Diagnosis of pelvic actinomycosis by 16S ribosomal RNA gene sequencing and its clinical significance. Diagnostic Microbiology and Infectious Disease, 2002, 43, 113-118.	1.8	44
375	Automated Identification of Medically Important Bacteria by 16S rRNA Gene Sequencing Using a Novel Comprehensive Database, 16SpathDB. Journal of Clinical Microbiology, 2011, 49, 1799-1809.	3.9	44
376	Clonality Despite Sex: The Evolution of Host-Associated Sexual Neighborhoods in the Pathogenic Fungus Penicillium marneffei. PLoS Pathogens, 2012, 8, e1002851.	4.7	44
377	Identification of a Novel Feline Picornavirus from the Domestic Cat. Journal of Virology, 2012, 86, 395-405.	3.4	44
378	Complete Genome Sequences of Novel Canine Noroviruses in Hong Kong. Journal of Virology, 2012, 86, 9531-9532.	3.4	44

#	Article	IF	CITATIONS
379	NF90 Exerts Antiviral Activity through Regulation of PKR Phosphorylation and Stress Granules in Infected Cells. Journal of Immunology, 2014, 192, 3753-3764.	0.8	44
380	Metabolomic Profiling of Plasma from Patients with Tuberculosis by Use of Untargeted Mass Spectrometry Reveals Novel Biomarkers for Diagnosis. Journal of Clinical Microbiology, 2015, 53, 3750-3759.	3.9	44
381	Enterovirus D68 Infections Associated with Severe Respiratory Illness in Elderly Patients and Emergence of a Novel Clade in Hong Kong. Scientific Reports, 2016, 6, 25147.	3.3	44
382	Immunoassays Based on Penicillium marneffei Mp1p Derived from Pichia pastoris Expression System for Diagnosis of Penicilliosis. PLoS ONE, 2011, 6, e28796.	2.5	44
383	Identification of a Novel Bat Papillomavirus by Metagenomics. PLoS ONE, 2012, 7, e43986.	2.5	44
384	Actinomyces hongkongensis sp. nov. $\hat{a}\in$ A Novel Actinomyces species Isolated from a Patient with Pelvic Actinomycosis. Systematic and Applied Microbiology, 2003, 26, 518-522.	2.8	43
385	Leptotrichia hongkongensis sp. nov., a novel Leptotrichia species with the oral cavity as its natural reservoir. Journal of Zhejiang University: Science B, 2010, 11, 391-401.	2.8	43
386	Discovery and Genomic Characterization of a Novel Bat Sapovirus with Unusual Genomic Features and Phylogenetic Position. PLoS ONE, 2012, 7, e34987.	2.5	43
387	Functional variants regulating LGALS1 (Galectin 1) expression affect human susceptibility to influenza A(H7N9). Scientific Reports, 2015, 5, 8517.	3.3	43
388	Genetic diversity of Aspergillus species isolated from onychomycosis and Aspergillus hongkongensis sp. nov., with implications to antifungal susceptibility testing. Diagnostic Microbiology and Infectious Disease, 2016, 84, 125-134.	1.8	43
389	Activation of C-Type Lectin Receptor and (RIG)-I-Like Receptors Contributes to Proinflammatory Response in Middle East Respiratory Syndrome Coronavirus-Infected Macrophages. Journal of Infectious Diseases, 2020, 221, 647-659.	4.0	43
390	Cross-linking peptide and repurposed drugs inhibit both entry pathways of SARS-CoV-2. Nature Communications, 2021, 12, 1517.	12.8	43
391	Fatal Necrotizing Fasciitis due to Vibrio damsela. Scandinavian Journal of Infectious Diseases, 1993, 25, 659-661.	1.5	42
392	Enterococcus cecorum Empyema Thoracis Successfully Treated with Cefotaxime. Journal of Clinical Microbiology, 2004, 42, 919-922.	3.9	42
393	Characterization of Haemophilus segnis , an Important Cause of Bacteremia, by 16S rRNA Gene Sequencing. Journal of Clinical Microbiology, 2004, 42, 877-880.	3.9	42
394	Draft Genome Sequence of Penicillium marneffei Strain PM1. Eukaryotic Cell, 2011, 10, 1740-1741.	3 . 4	42
395	Genetic characterization of EV71 isolates from 2004 to 2010 reveals predominance and persistent circulation of the newly proposed genotype D and recent emergence of a distinct lineage of subgenotype C2 in Hong Kong. Virology Journal, 2013, 10, 222.	3.4	42
396	Transmission of methicillin-resistant staphylococcus aureus in the long term care facilities in Hong Kong. BMC Infectious Diseases, 2013, 13, 205.	2.9	42

#	Article	IF	Citations
397	Identification and characterization of a novel paramyxovirus, porcine parainfluenza virus 1, from deceased pigs. Journal of General Virology, 2013, 94, 2184-2190.	2.9	42
398	Hemagglutinin of influenza A virus binds specifically to cell surface nucleolin and plays a role in virus internalization. Virology, 2016, 494, 78-88.	2.4	42
399	Identification of a Novel Betacoronavirus (Merbecovirus) in Amur Hedgehogs from China. Viruses, 2019, 11, 980.	3.3	42
400	Impact of Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) Variant-Associated Receptor Binding Domain (RBD) Mutations on the Susceptibility to Serum Antibodies Elicited by Coronavirus Disease 2019 (COVID-19) Infection or Vaccination. Clinical Infectious Diseases, 2022, 74, 1623-1630.	5.8	42
401	Co-circulation of two SARS-CoV-2 variant strains within imported pet hamsters in Hong Kong. Emerging Microbes and Infections, 2022, 11, 689-698.	6.5	42
402	Detection of Cell Wall Galactomannoprotein Afmp1p in Culture Supernatants of Aspergillus fumigatus and in Sera of Aspergillosis Patients. Journal of Clinical Microbiology, 2002, 40, 4382-4387.	3.9	41
403	Current status and future directions for Laribacter hongkongensis, a novel bacterium associated with gastroenteritis and traveller's diarrhoea. Current Opinion in Infectious Diseases, 2005, 18, 413-419.	3.1	41
404	MBEToolbox: a Matlab toolbox for sequence data analysis in molecular biology and evolution. BMC Bioinformatics, 2005, 6, 64.	2.6	41
405	Recombinant adeno-associated virus expressing the receptor-binding domain of severe acute respiratory syndrome coronavirus S protein elicits neutralizing antibodies: Implication for developing SARS vaccines. Virology, 2006, 353, 6-16.	2.4	41
406	Polymorphisms of type I interferon receptor 1 promoter and their effects on chronic hepatitis B virus infection. Journal of Hepatology, 2007, 46, 198-205.	3.7	41
407	CoVDB: a comprehensive database for comparative analysis of coronavirus genes and genomes. Nucleic Acids Research, 2008, 36, D504-D511.	14.5	41
408	Antimicrobial resistance among uropathogens that cause acute uncomplicated cystitis in women in Hong Kong: a prospective multicenter study in 2006 to 2008. Diagnostic Microbiology and Infectious Disease, 2010, 66, 87-93.	1.8	41
409	Matrix-assisted laser desorption ionisation time-of-flight mass spectrometry for identification of clinically significant bacteria that are difficult to identify in clinical laboratories. Journal of Clinical Pathology, 2014, 67, 361-366.	2.0	41
410	Molecular diagnosis in clinical parasitology: When and why?. Experimental Biology and Medicine, 2014, 239, 1443-1460.	2.4	41
411	Japanese Encephalitis Virus Transmitted Via Blood Transfusion, Hong Kong, China. Emerging Infectious Diseases, 2018, 24, 49-57.	4.3	41
412	Differential immune activation profile of SARS-CoV-2 and SARS-CoV infection in human lung and intestinal cells: Implications for treatment with IFN- \hat{l}^2 and IFN inducer. Journal of Infection, 2020, 81, e1-e10.	3.3	41
413	SARS-CoV-2 Induces a More Robust Innate Immune Response and Replicates Less Efficiently Than SARS-CoV in the Human Intestines: An ExÂVivo Study With Implications on Pathogenesis of COVID-19. Cellular and Molecular Gastroenterology and Hepatology, 2021, 11, 771-781.	4.5	41
414	Analysis of a Viridans Group Strain Reveals a Case of Bacteremia Due to Lancefield Group G Alpha-Hemolytic Streptococcus dysgalactiae subsp. equisimilis in a Patient with Pyomyositis and Reactive Arthritis. Journal of Clinical Microbiology, 2003, 41, 613-618.	3.9	40

#	Article	IF	CITATIONS
415	AFLMP1 Encodes an Antigenic Cell Wall Protein in Aspergillus flavus. Journal of Clinical Microbiology, 2003, 41, 845-850.	3.9	40
416	Strategic measures for the control of surging antimicrobial resistance in Hong Kong and mainland of China. Emerging Microbes and Infections, 2015, 4, 1-13.	6.5	40
417	Pulmonary and extrapulmonary complications of human rhinovirus infection in critically ill patients. Journal of Clinical Virology, 2016, 77, 85-91.	3.1	40
418	Loss of orf3b in the circulating SARS-CoV-2 strains. Emerging Microbes and Infections, 2020, 9, 2685-2696.	6.5	40
419	Viruses harness $Yxx\tilde{A}^{-}$ motif to interact with host AP2M1 for replication: A vulnerable broad-spectrum antiviral target. Science Advances, 2020, 6, eaba7910.	10.3	40
420	Confirmation of the First Hong Kong Case of Human Infection by Novel Swine Origin Influenza A (H1N1) Virus Diagnosed Using Ultrarapid, Real-Time Reverse Transcriptase PCR. Journal of Clinical Microbiology, 2009, 47, 2344-2346.	3.9	39
421	A tricyclic pyrrolobenzodiazepine produced by Klebsiella oxytoca is associated with cytotoxicity in antibiotic-associated hemorrhagic colitis. Journal of Biological Chemistry, 2017, 292, 19503-19520.	3.4	39
422	Unique Clusters of Severe Acute Respiratory Syndrome Coronavirus 2 Causing a Large Coronavirus Disease 2019 Outbreak in Hong Kong. Clinical Infectious Diseases, 2021, 73, 137-142.	5.8	39
423	Human tryptophanyl-tRNA synthetase is an IFN-γ–inducible entry factor for Enterovirus. Journal of Clinical Investigation, 2018, 128, 5163-5177.	8.2	39
424	The study of cytokine dynamics at the operation site after mastectomy. Wound Repair and Regeneration, 2003, 11, 326-330.	3.0	38
425	Use of Cefoperazone MacConkey Agar for Selective Isolation of Laribacter hongkongensis. Journal of Clinical Microbiology, 2003, 41, 4839-4841.	3.9	38
426	Natural selection retains overrepresented out-of-frame stop codons against frameshift peptides in prokaryotes. BMC Genomics, 2010, 11, 491.	2.8	38
427	Comparative Evaluation of a Point-of-Care Immunochromatographic Test SNAP 4Dx with Molecular Detection Tests for Vector-Borne Canine Pathogens in Hong Kong. Vector-Borne and Zoonotic Diseases, 2011, 11, 1269-1277.	1.5	38
428	Signature Gene Expression Reveals Novel Clues to the Molecular Mechanisms of Dimorphic Transition in Penicillium marneffei. PLoS Genetics, 2014, 10, e1004662.	3.5	38
429	Human Intestinal Organoids Recapitulate Enteric Infections of Enterovirus and Coronavirus. Stem Cell Reports, 2021, 16, 493-504.	4.8	38
430	The impact of spike N501Y mutation on neutralizing activity and RBD binding of SARS-CoV-2 convalescent serum. EBioMedicine, 2021, 71, 103544.	6.1	38
431	Paired heavy- and light-chain signatures contribute to potent SARS-CoV-2 neutralization in public antibody responses. Cell Reports, 2021, 37, 109771.	6.4	38
432	First Report of Tsukamurella Keratitis: Association between T. tyrosinosolvens and T. pulmonis and Ophthalmologic Infections. Journal of Clinical Microbiology, 2009, 47, 1953-1956.	3.9	37

#	Article	IF	CITATIONS
433	Decolonization of gastrointestinal carriage of vancomycin-resistant Enterococcus faecium: case series and review of literature. BMC Infectious Diseases, 2014, 14, 514.	2.9	37
434	Corynebacterium kroppenstedtii Is an Emerging Cause of Mastitis Especially in Patients With Psychiatric Illness on Antipsychotic Medication. Open Forum Infectious Diseases, 2017, 4, ofx096.	0.9	37
435	Detection of SARS-CoV-2 in conjunctival secretions from patients without ocular symptoms. Infection, 2021, 49, 257-265.	4.7	37
436	IFCC interim guidelines on rapid point-of-care antigen testing for SARS-CoV-2 detection in asymptomatic and symptomatic individuals. Clinical Chemistry and Laboratory Medicine, 2021, 59, 1507-1515.	2.3	37
437	Diagnostic application of genotypic identification of mycobacteria. Journal of Medical Microbiology, 2006, 55, 529-536.	1.8	36
438	Logistical feasibility and potential benefits of a population-wide passive-immunotherapy program during an influenza pandemic. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 3269-3274.	7.1	36
439	NMR-based metabolomic urinalysis: A rapid screening test for urinary tract infection. Clinica Chimica Acta, 2014, 436, 217-223.	1.1	36
440	Use of MALDI Biotyper plus ClinProTools mass spectra analysis for correct identification of <i>Streptococcus pneumoniae</i> and <i>Streptococcus mitis</i> li>oralisDournal of Clinical Pathology, 2015, 68, 652-656.	2.0	36
441	Identification of nsp1 gene as the target of SARSâ€CoVâ€2 realâ€time RTâ€PCR using nanopore wholeâ€genome sequencing. Journal of Medical Virology, 2020, 92, 2725-2734.	5.0	36
442	Synthetic peptides outside the spike protein heptad repeat regions as potent inhibitors of SARS-associated coronavirus. Antiviral Therapy, 2005, 10, 393-403.	1.0	36
443	Molecular epidemiology of methicillin-resistant Staphylococcus aureus in residential care homes for the elderly in Hong Kong. Diagnostic Microbiology and Infectious Disease, 2008, 61, 135-142.	1.8	35
444	Identification of Major Histocompatibility Complex Class I C Molecule as an Attachment Factor That Facilitates Coronavirus HKU1 Spike-Mediated Infection. Journal of Virology, 2009, 83, 1026-1035.	3.4	35
445	Discovery and Genomic Characterization of a Novel Ovine Partetravirus and a New Genotype of Bovine Partetravirus. PLoS ONE, 2011, 6, e25619.	2.5	35
446	Recombinant Coxsackievirus A2 and Deaths of Children, Hong Kong, 2012. Emerging Infectious Diseases, 2013, 19, 1285-1288.	4.3	35
447	Avian Influenza A H7N9 Virus Induces Severe Pneumonia in Mice without Prior Adaptation and Responds to a Combination of Zanamivir and COX-2 Inhibitor. PLoS ONE, 2014, 9, e107966.	2.5	35
448	Ebola virus disease: a highly fatal infectious disease reemerging in West Africa. Microbes and Infection, 2015, 17, 84-97.	1.9	35
449	The celecoxib derivative kinase inhibitor AR-12 (OSU-03012) inhibits Zika virus via down-regulation of the PI3K/Akt pathway and protects Zika virus-infected A129 mice: A host-targeting treatment strategy. Antiviral Research, 2018, 160, 38-47.	4.1	35
450	Prophage exotoxins enhance colonization fitness in epidemic scarlet fever-causing Streptococcus pyogenes. Nature Communications, 2020, 11, 5018.	12.8	35

#	Article	IF	CITATIONS
451	DNA immunization using a secreted cell wall antigen Mp1p is protective against Penicillium marneffei infection. Vaccine, 2002, 20, 2878-2886.	3.8	34
452	Seasonal and tissue distribution of Laribacter hongkongensis, a novel bacterium associated with gastroenteritis, in retail freshwater fish in Hong Kong. International Journal of Food Microbiology, 2007, 113, 62-66.	4.7	34
453	Complete Genome Sequence of the Veterinary Pathogen <i>Staphylococcus pseudintermedius</i> Strain HKU10-03, Isolated in a Case of Canine Pyoderma. Journal of Bacteriology, 2011, 193, 1783-1784.	2.2	34
454	High Mortality Associated with Catabacter hongkongensis Bacteremia. Journal of Clinical Microbiology, 2012, 50, 2239-2243.	3.9	34
455	Unique reassortant of influenza A(H7N9) virus associated with severe disease emerging in Hong Kong. Journal of Infection, 2014, 69, 60-68.	3.3	34
456	Chickens host diverse picornaviruses originated from potential interspecies transmission with recombination. Journal of General Virology, 2014, 95, 1929-1944.	2.9	34
457	Improved detection of Zika virus <scp>RNA</scp> in human and animal specimens by a novel, highly sensitive and specific realâ€time RTâ€PCR assay targeting the 5′â€untranslated region of Zika virus. Tropical Medicine and International Health, 2017, 22, 594-603.	2.3	34
458	Early-Morning vs Spot Posterior Oropharyngeal Saliva for Diagnosis of SARS-CoV-2 Infection: Implication of Timing of Specimen Collection for Community-Wide Screening. Open Forum Infectious Diseases, 2020, 7, ofaa210.	0.9	34
459	Cloning and Characterization of a Chromosomal Class C \hat{l}^2 -Lactamase and Its Regulatory Gene in Laribacter hongkongensis. Antimicrobial Agents and Chemotherapy, 2005, 49, 1957-1964.	3.2	33
460	<i>In Silico</i> Analysis of ORF1ab in Coronavirus HKU1 Genome Reveals a Unique Putative Cleavage Site of Coronavirus HKU1 3Câ€Like Protease. Microbiology and Immunology, 2005, 49, 899-908.	1.4	33
461	Implementation of directly observed patient hand hygiene for hospitalized patients by hand hygiene ambassadors in Hong Kong. American Journal of Infection Control, 2016, 44, 621-624.	2.3	33
462	First Report of a Fatal Case Associated with EV-D68 Infection in Hong Kong and Emergence of an Interclade Recombinant in China Revealed by Genome Analysis. International Journal of Molecular Sciences, 2017, 18, 1065.	4.1	33
463	Replication of MERS and SARS coronaviruses in bat cells offers insights to their ancestral origins. Emerging Microbes and Infections, 2018, 7, 1-11.	6.5	33
464	Detection of katG Ser315Thr substitution in respiratory specimens from patients with isoniazid-resistant Mycobacterium tuberculosis using PCR-RFLP. Journal of Medical Microbiology, 2003, 52, 999-1003.	1.8	33
465	Age-associated SARS-CoV-2 breakthrough infection and changes in immune response in a mouse model. Emerging Microbes and Infections, 2022, 11, 368-383.	6.5	33
466	Acute Disseminated Encephalomyelitis After Para-influenza Infection Post Bone Marrow Transplantation. Leukemia and Lymphoma, 2002, 43, 455-457.	1.3	32
467	AFMP2 Encodes a Novel Immunogenic Protein of the Antigenic Mannoprotein Superfamily in Aspergillus fumigatus. Journal of Clinical Microbiology, 2004, 42, 2287-2291.	3.9	32
468	Life-Threatening Invasive Helcococcus kunzii Infections in Intravenous-Drug Users and ermA -Mediated Erythromycin Resistance. Journal of Clinical Microbiology, 2005, 43, 6205-6208.	3.9	32

#	Article	IF	Citations
469	Bacteremia Caused by Solobacterium moorei in a Patient with Acute Proctitis and Carcinoma of the Cervix. Journal of Clinical Microbiology, 2006, 44, 3031-3034.	3.9	32
470	Fatal co-infection with swine origin influenza virus A/H1N1 and community-acquired methicillin-resistant Staphylococcus aureus. Journal of Infection, 2009, 59, 366-370.	3.3	32
471	First Report of Spontaneous Intrapartum Atopobium vaginae Bacteremia. Journal of Clinical Microbiology, 2012, 50, 2525-2528.	3.9	32
472	Unraveling the Molecular Basis of Temperature-Dependent Genetic Regulation in Penicillium marneffei. Eukaryotic Cell, 2013, 12, 1214-1224.	3.4	32
473	Control of hospital endemicity of multiple-drug-resistant Acinetobacter baumannii ST457 with directly observed hand hygiene. European Journal of Clinical Microbiology and Infectious Diseases, 2015, 34, 713-718.	2.9	32
474	Antibody-Dependent Cell-Mediated Cytotoxicity Epitopes on the Hemagglutinin Head Region of Pandemic H1N1 Influenza Virus Play Detrimental Roles in H1N1-Infected Mice. Frontiers in Immunology, 2017, 8, 317.	4.8	32
475	Vaccine-Elicited CD8+ T Cells Cure Mesothelioma by Overcoming Tumor-Induced Immunosuppressive Environment. Cancer Research, 2014, 74, 6010-6021.	0.9	32
476	ACE2-like carboxypeptidase B38-CAP protects from SARS-CoV-2-induced lung injury. Nature Communications, 2021, 12, 6791.	12.8	32
477	Nasal prevention of SARS-CoV-2 infection by intranasal influenza-based boost vaccination in mouse models. EBioMedicine, 2022, 75, 103762.	6.1	32
478	A regulatory polymorphism in interferon- \hat{I}^3 receptor 1 promoter is associated with the susceptibility to chronic hepatitis B virus infection. Immunogenetics, 2009, 61, 423-430.	2.4	31
479	Viral lung infections. Current Opinion in Pulmonary Medicine, 2014, 20, 225-232.	2.6	31
480	Lipid mediators of inflammation as novel plasma biomarkers to identify patients with bacteremia. Journal of Infection, 2015, 70, 433-444.	3.3	31
481	Emergence of carbapenem-resistant hypervirulent Klebsiella pneumoniae. Lancet Infectious Diseases, The, 2018, 18, 24.	9.1	31
482	Donor-Derived Genotype 4 Hepatitis E Virus Infection, Hong Kong, China, 2018. Emerging Infectious Diseases, 2019, 25, 425-433.	4.3	31
483	Infection control challenges in setting up community isolation and treatment facilities for patients with coronavirus disease 2019 (COVID-19): Implementation of directly observed environmental disinfection. Infection Control and Hospital Epidemiology, 2021, 42, 1037-1045.	1.8	31
484	Peptide-based pan-CoV fusion inhibitors maintain high potency against SARS-CoV-2 Omicron variant. Cell Research, 2022, 32, 404-406.	12.0	31
485	An antibody class with a common CDRH3 motif broadly neutralizes sarbecoviruses. Science Translational Medicine, 2022, 14, eabn6859.	12.4	31
486	Adult Croup: A Rare but More Severe Condition. Respiration, 2000, 67, 684-688.	2.6	30

#	Article	IF	CITATIONS
487	Clinical and Molecular Epidemiology of Erythromycin-Resistant Beta-Hemolytic Lancefield Group G Streptococci Causing Bacteremia. Journal of Clinical Microbiology, 2003, 41, 5188-5191.	3.9	30
488	Differential selection and mutation between dsDNA and ssDNA phages shape the evolution of their genomic AT percentage., 2005, 6, 20.		30
489	Agar Block Smear Preparation: a Novel Method of Slide Preparation for Preservation of Native Fungal Structures for Microscopic Examination and Long-Term Storage. Journal of Clinical Microbiology, 2010, 48, 3053-3061.	3.9	30
490	First Report of Disseminated Mycobacterium Skin Infections in Two Liver Transplant Recipients and Rapid Diagnosis byhsp65Gene Sequencing. Journal of Clinical Microbiology, 2011, 49, 3733-3738.	3.9	30
491	Colistin-Resistant <i>Enterobacteriaceae</i> Carrying the <i>mcr-1</i> Gene among Patients in Hong Kong. Emerging Infectious Diseases, 2016, 22, 1667-1669.	4.3	30
492	Middle East Respiratory Syndrome Coronavirus ORF8b Accessory Protein Suppresses Type I IFN Expression by Impeding HSP70-Dependent Activation of IRF3 Kinase IKKÎμ. Journal of Immunology, 2020, 205, 1564-1579.	0.8	30
493	Vaccineâ€breakthrough infection by the SARSâ€CoVâ€2 omicron variant elicits broadly crossâ€reactive immune responses. Clinical and Translational Medicine, 2022, 12, e720.	4.0	30
494	Clarithromycin Attenuates Mastectomy-Induced Acute Inflammatory Response. Vaccine Journal, 2000, 7, 925-931.	2.6	29
495	Quantification of Adenovirus in the Lower Respiratory Tract of Patients without Clinical Adenovirus-Related Respiratory Disease. Clinical Infectious Diseases, 2005, 40, 1541-1544.	5.8	29
496	Fangia hongkongensis gen. nov., sp. nov., a novel gammaproteobacterium of the order Thiotrichales isolated from coastal seawater of Hong Kong. International Journal of Systematic and Evolutionary Microbiology, 2007, 57, 2665-2669.	1.7	29
497	A Systematic Approach to Novel Virus Discovery in Emerging Infectious Disease Outbreaks. Journal of Molecular Diagnostics, 2015, 17, 230-241.	2.8	29
498	<i>Coronavirus HKU15</i> in respiratory tract of pigs and first discovery of coronavirus quasispecies in 5 ′-untranslated region. Emerging Microbes and Infections, 2017, 6, 1-7.	6.5	29
499	Multimodal investigation of rat hepatitis E virus antigenicity: Implications for infection, diagnostics, and vaccine efficacy. Journal of Hepatology, 2021, 74, 1315-1324.	3.7	29
500	Mp1p Is a Virulence Factor in Talaromyces (Penicillium) marneffei. PLoS Neglected Tropical Diseases, 2016, 10, e0004907.	3.0	29
501	SARS-CoV-2 exploits host DGAT and ADRP for efficient replication. Cell Discovery, 2021, 7, 100.	6.7	29
502	Bacillus Calmette-Guérin–induced trained immunity protects against SARS-CoV-2 challenge in K18-hACE2 mice. JCI Insight, 2022, 7, .	5.0	29
503	<i>MP1</i> Homologue-Based Multilocus Sequence System for Typing the Pathogenic Fungus <i>Penicillium marneffei</i> : a Novel Approach Using Lineage-Specific Genes. Journal of Clinical Microbiology, 2007, 45, 3647-3654.	3.9	28
504	Fine antigenic variation within H5N1 influenza virus hemagglutinin's antigenic sites defined by yeast cell surface display. European Journal of Immunology, 2009, 39, 3498-3510.	2.9	28

#	Article	IF	CITATIONS
505	Functional dissection of an IFN- $\hat{l}\pm\hat{l}^2$ receptor 1 promoter variant that confers higher risk to chronic hepatitis B virus infection. Journal of Hepatology, 2009, 51, 322-332.	3.7	28
506	Epidemiology and clinical features of Shewanella infection over an eight-year period. Scandinavian Journal of Infectious Diseases, 2010, 42, 757-762.	1.5	28
507	Complete Genome Sequence of a Novel Picornavirus, Canine Picornavirus, Discovered in Dogs. Journal of Virology, 2012, 86, 3402-3403.	3.4	28
508	Detection of human rhinovirus C in fecal samples of children with gastroenteritis. Journal of Clinical Virology, 2012, 53, 290-296.	3.1	28
509	Use of Nasopharyngeal Aspirate for Diagnosis of Pneumocystis Pneumonia. Journal of Clinical Microbiology, 2013, 51, 1570-1574.	3.9	28
510	Identification of Novel Fusion Inhibitors of Influenza A Virus by Chemical Genetics. Journal of Virology, 2016, 90, 2690-2701.	3.4	28
511	A bipotential organoid model of respiratory epithelium recapitulates high infectivity of SARS-CoV-2 Omicron variant. Cell Discovery, 2022, 8, .	6.7	28
512	Alkanindiges hongkongensis sp. nov. A novel Alkanindiges species isolated from a patient with parotid abscess. Systematic and Applied Microbiology, 2005, 28, 316-322.	2.8	27
513	Isolation of Laribacter hongkongensis, a novel bacterium associated with gastroenteritis, from Chinese tiger frog. International Journal of Food Microbiology, 2009, 129, 78-82.	4.7	27
514	High Incidence of Severe Influenza among Individuals over 50 Years of Age. Vaccine Journal, 2011, 18, 1918-1924.	3.1	27
515	Rapid Identification and Validation of Specific Molecular Targets for Detection of Escherichia coli O104:H4 Outbreak Strain by Use of High-Throughput Sequencing Data from Nine Genomes. Journal of Clinical Microbiology, 2011, 49, 3714-3716.	3.9	27
516	Monitoring the fitness of antiviral-resistant influenza strains during an epidemic: a mathematical modelling study. Lancet Infectious Diseases, The, 2017, 17, 339-347.	9.1	27
517	Rhinovirus respiratory tract infection in hospitalized adult patients is associated with T H 2 response irrespective of asthma. Journal of Infection, 2018, 76, 465-474.	3.3	27
518	Performance of a Surrogate SARS-CoV-2-Neutralizing Antibody Assay in Natural Infection and Vaccination Samples. Diagnostics, 2021, 11, 1757.	2.6	27
519	Intranasal administration of a single dose of a candidate live attenuated vaccine derived from an NSP16-deficient SARS-CoV-2 strain confers sterilizing immunity in animals. , 2022, 19, 588-601.		27
520	Regular virologic surveillance showed very frequent cytomegalovirus reactivation in patients treated with alemtuzumab. American Journal of Hematology, 2007, 82, 108-111.	4.1	26
521	Well-Characterized Monoclonal Antibodies against Cell Wall Antigen of Aspergillus Species Improve Immunoassay Specificity and Sensitivity. Vaccine Journal, 2008, 15, 194-202.	3.1	26
522	Complete Genome Sequence of a Novel Picobirnavirus, Otarine Picobirnavirus, Discovered in California Sea Lions. Journal of Virology, 2012, 86, 6377-6378.	3.4	26

#	Article	IF	CITATIONS
523	Recombinant ESAT-6-Like Proteins Provoke Protective Immune Responses against Invasive Staphylococcus aureus Disease in a Murine Model. Infection and Immunity, 2015, 83, 339-345.	2.2	26
524	Talaromyces marneffei Mp1p Is a Virulence Factor that Binds and Sequesters a Key Proinflammatory Lipid to Dampen Host Innate Immune Response. Cell Chemical Biology, 2017, 24, 182-194.	5.2	26
525	Control of multidrug-resistant Acinetobacter baumannii in Hong Kong: Role of environmental surveillance in communal areas after a hospital outbreak. American Journal of Infection Control, 2018, 46, 60-66.	2.3	26
526	Characterization of an attenuated SARS-CoV-2 variant with a deletion at the $$1/$S2$ junction of the spike protein. Nature Communications, 2021, 12, 2790.	12.8	26
527	Phylogenomic analysis of COVID-19 summer and winter outbreaks in Hong Kong: An observational study. The Lancet Regional Health - Western Pacific, 2021, 10, 100130.	2.9	26
528	Thermo-tolerant Campylobacter fetus bacteraemia identified by 16S ribosomal RNA gene sequencing: an emerging pathogen in immunocompromised patients. Journal of Medical Microbiology, 2002, 51, 740-746.	1.8	26
529	Use of Recombinant Mitogillin for Serodiagnosis of Aspergillus fumigatus -Associated Diseases. Journal of Clinical Microbiology, 2001, 39, 4598-4600.	3.9	25
530	Seronegative Bacteremic Melioidosis Caused by Burkholderia pseudomallei with Ambiguous Biochemical Profile: Clinical Importance of Accurate Identification by 16S rRNA Gene and groEL Gene Sequencing. Journal of Clinical Microbiology, 2003, 41, 3973-3977.	3.9	25
531	Distribution and molecular characterization of tetracycline resistance in Laribacter hongkongensis. Journal of Antimicrobial Chemotherapy, 2008, 61, 488-497.	3.0	25
532	Detection of human novel influenza A (H1N1) viruses using multi-fluorescent real-time RT-PCR. Virus Research, 2010, 147, 85-90.	2.2	25
533	Complete Genome Sequence of a Coxsackievirus A22 Strain in Hong Kong Reveals a Natural Intratypic Recombination Event. Journal of Virology, 2011, 85, 12098-12099.	3.4	25
534	A significant number of reported <i>Absidia corymbifera</i> (<i>Lichtheimia corymbifera</i>) infections are caused by <i>Lichtheimia ramosa</i> (syn. <i>Lichtheimia hongkongensis</i>): an emerging cause of mucormycosis. Emerging Microbes and Infections, 2012, 1, 1-8.	6.5	25
535	Proactive infection control measures to prevent nosocomial transmission of vancomycin-resistant enterococci in HongÂKong. Journal of the Formosan Medical Association, 2014, 113, 734-741.	1.7	25
536	Host genes and influenza pathogenesis in humans: an emerging paradigm. Current Opinion in Virology, 2015, 14, 7-15.	5.4	25
537	High Diversity of Genogroup I Picobirnaviruses in Mammals. Frontiers in Microbiology, 2016, 7, 1886.	3.5	25
538	A six-year descriptive epidemiological study of human coronavirus infections in hospitalized patients in Hong Kong. Virologica Sinica, 2016, 31, 41-48.	3.0	25
539	Identification and genomic characterization of a novel rat bocavirus from brown rats in China. Infection, Genetics and Evolution, 2017, 47, 68-76.	2.3	25
540	Development and evaluation of a conventional RTâ€PCR for differentiating emerging influenza B/Victoria lineage viruses with hemagglutinin amino acid deletion from B/Yamagata lineage viruses. Journal of Medical Virology, 2020, 92, 382-385.	5.0	25

#	Article	IF	CITATIONS
541	A monoclonal antibody that neutralizes SARS-CoV-2 variants, SARS-CoV, and other sarbecoviruses. Emerging Microbes and Infections, 2022, 11, 147-157.	6.5	25
542	Streptococcus sinensis may react with Lancefield group F antiserum. Journal of Medical Microbiology, 2004, 53, 1083-1088.	1.8	24
543	Lichtheimia hongkongensis sp. nov., a novel Lichtheimia spp. associated with rhinocerebral, gastrointestinal, and cutaneous mucormycosis. Diagnostic Microbiology and Infectious Disease, 2010, 66, 274-284.	1.8	24
544	Epidemiology of Klebsiella oxytoca-Associated Diarrhea Detected by Simmons Citrate Agar Supplemented with Inositol, Tryptophan, and Bile Salts. Journal of Clinical Microbiology, 2012, 50, 1571-1579.	3.9	24
545	Streptococcus hongkongensis sp. nov., isolated from a patient with an infected puncture wound and from a marine flatfish. International Journal of Systematic and Evolutionary Microbiology, 2013, 63, 2570-2576.	1.7	24
546	Rapid Generation of Human-Like Neutralizing Monoclonal Antibodies in Urgent Preparedness for Influenza Pandemics and Virulent Infectious Diseases. PLoS ONE, 2013, 8, e66276.	2.5	24
547	A phylogenetically distinct Middle East respiratory syndrome coronavirus detected in a dromedary calf from a closed dairy herd in Dubai with rising seroprevalence with age. Emerging Microbes and Infections, 2015, 4, 1-5.	6.5	24
548	Polyphyletic origin of MERS coronaviruses and isolation of a novel clade A strain from dromedary camels in the United Arab Emirates. Emerging Microbes and Infections, 2016, 5, 1-9.	6.5	24
549	A Novel Risk Factor Associated With Colonization by Carbapenemase-Producing Enterobacteriaceae: Use of Proton Pump Inhibitors in Addition to Antimicrobial Treatment. Infection Control and Hospital Epidemiology, 2016, 37, 1418-1425.	1.8	24
550	Seasonal Outbreak of Bacillus Bacteremia Associated With Contaminated Linen in Hong Kong. Clinical Infectious Diseases, 2017, 64, S91-S97.	5.8	24
551	Rapid detection of MERS coronavirus-like viruses in bats: potential for tracking MERS coronavirus transmission and animal origin. Emerging Microbes and Infections, 2018, 7, 1-7.	6.5	24
552	Immunogenicity, safety and tolerability of intradermal influenza vaccines. Human Vaccines and Immunotherapeutics, 2018, 14, 565-570.	3.3	24
553	SMRT sequencing revealed the diversity and characteristics of defective interfering RNAs in influenza A (H7N9) virus infection. Emerging Microbes and Infections, 2019, 8, 662-674.	6.5	24
554	Absence of nosocomial transmission of coronavirus disease 2019 (COVID-19) due to SARS-CoV-2 in the prepandemic phase in Hong Kong. American Journal of Infection Control, 2020, 48, 890-896.	2.3	24
555	A pan-sarbecovirus vaccine induces highly potent and durable neutralizing antibody responses in non-human primates against SARS-CoV-2 Omicron variant. Cell Research, 2022, 32, 495-497.	12.0	24
556	Polymicrobial Outbreak of Intermittent Peritoneal Dialysis Peritonitis during External Wall Renovation at a Dialysis Center. Peritoneal Dialysis International, 2001, 21, 296-301.	2.3	23
557	The fight over flu. Nature, 2012, 481, 257-259.	27.8	23
558	Surveillance and Genome Analysis of Human Bocavirus in Patients with Respiratory Infection in Guangzhou, China. PLoS ONE, 2012, 7, e44876.	2.5	23

#	Article	IF	Citations
559	Evaluation of the molecular Xpert Xpress Flu/RSV assay vs. Alere i Influenza A & B assay for rapid detection of influenza viruses. Diagnostic Microbiology and Infectious Disease, 2018, 90, 177-180.	1.8	23
560	Whole-genome sequencing data-based modeling for the investigation of an outbreak of community-associated methicillin-resistant Staphylococcus aureus in a neonatal intensive care unit in Hong Kong. European Journal of Clinical Microbiology and Infectious Diseases, 2019, 38, 563-573.	2.9	23
561	A novel astrovirus from dromedaries in the Middle East. Journal of General Virology, 2015, 96, 2697-2707.	2.9	23
562	Targeting papain-like protease for broad-spectrum coronavirus inhibition. Protein and Cell, 2022, 13, 940-953.	11.0	23
563	Boosting of serum neutralizing activity against the Omicron variant among recovered COVID-19 patients by BNT162b2 and CoronaVac vaccines. EBioMedicine, 2022, 79, 103986.	6.1	23
564	A crucial role of macrophages in the immune responses to oral DNA vaccination against hepatitis B virus in a murine model. Vaccine, 2001, 20, 140-147.	3.8	22
565	Evolution of Antibiotic Resistance Mechanisms and their Relevance to Dialysis-Related Infections. Peritoneal Dialysis International, 2007, 27, 272-280.	2.3	22
566	Increased Genetic Diversity of HIV-1 Circulating in Hong Kong. PLoS ONE, 2010, 5, e12198.	2.5	22
567	Acupuncture transmitted infections. BMJ: British Medical Journal, 2010, 340, c1268-c1268.	2.3	22
568	Safety and immunogenicity of two different doses of a Vero cell-derived, whole virus clade 2 H5N1 (A/Indonesia/05/2005) influenza vaccine. Vaccine, 2012, 30, 329-335.	3.8	22
569	Complete Genome Sequence of a Novel Feline Astrovirus from a Domestic Cat in Hong Kong. Genome Announcements, 2013, 1, .	0.8	22
570	Isolation and Characterization of Dromedary Camel Coronavirus UAE-HKU23 from Dromedaries of the Middle East: Minimal Serological Cross-Reactivity between MERS Coronavirus and Dromedary Camel Coronavirus UAE-HKU23. International Journal of Molecular Sciences, 2016, 17, 691.	4.1	22
571	The persistent prevalence and evolution of cross-family recombinant coronavirus GCCDC1 among a bat population: a two-year follow-up. Science China Life Sciences, 2017, 60, 1357-1363.	4.9	22
572	Novel Picobirnaviruses in Respiratory and Alimentary Tracts of Cattle and Monkeys with Large Intraand Inter-Host Diversity. Viruses, 2019, 11, 574.	3.3	22
573	Identification of a Novel Ichthyic Parvovirus in Marine Species in Hainan Island, China. Frontiers in Microbiology, 2019, 10, 2815.	3.5	22
574	Development and Evaluation of Novel and Highly Sensitive Single-Tube Nested Real-Time RT-PCR Assays for SARS-CoV-2 Detection. International Journal of Molecular Sciences, 2020, 21, 5674.	4.1	22
575	Dynamic PB2-E627K substitution of influenza H7N9 virus indicates the in vivo genetic tuning and rapid host adaptation. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 23807-23814.	7.1	22
576	Host-derived lipids orchestrate pulmonary $\hat{I}^3\hat{I}$ Cell response to provide early protection against influenza virus infection. Nature Communications, 2021, 12, 1914.	12.8	22

#	Article	IF	Citations
577	SARS-CoV-2 infection induces inflammatory bone loss in golden Syrian hamsters. Nature Communications, 2022, 13, 2539.	12.8	22
578	A broadly neutralizing antibody protects Syrian hamsters against SARS-CoV-2 Omicron challenge. Nature Communications, 2022, 13, .	12.8	22
579	Candida tropicalis and Penicillium marneffei Mixed Fungaemia in a Patient with Waldenstr¶m's Macroglobulinaemia. European Journal of Clinical Microbiology and Infectious Diseases, 2001, 20, 132-135.	2.9	21
580	A 7‥EARâ€OLD BOY DYING OF ACUTE ENCEPHALOPATHY. Brain Pathology, 2010, 20, 261-264.	4.1	21
581	The 2008–2009 H1N1 influenza virus exhibits reduced susceptibility to antibody inhibition: Implications for the prevalence of oseltamivir resistant variant viruses. Antiviral Research, 2012, 93, 144-153.	4.1	21
582	Comparison of Pyrosequencing, Sanger Sequencing, and Melting Curve Analysis for Detection of Low-Frequency Macrolide-Resistant Mycoplasma pneumoniae Quasispecies in Respiratory Specimens. Journal of Clinical Microbiology, 2013, 51, 2592-2598.	3.9	21
583	Co-stimulation With TLR7 Agonist Imiquimod and Inactivated Influenza Virus Particles Promotes Mouse B Cell Activation, Differentiation, and Accelerated Antigen Specific Antibody Production. Frontiers in Immunology, 2018, 9, 2370.	4.8	21
584	H7N9 influenza A virus activation of necroptosis in human monocytes links innate and adaptive immune responses. Cell Death and Disease, 2019, 10, 442.	6.3	21
585	Clinical Performance of the Luminex NxTAG CoV Extended Panel for SARS-CoV-2 Detection in Nasopharyngeal Specimens from COVID-19 Patients in Hong Kong. Journal of Clinical Microbiology, 2020, 58, .	3.9	21
586	Virus subtype-specific suppression of MAVS aggregation and activation by PB1-F2 protein of influenza A (H7N9) virus. PLoS Pathogens, 2020, 16 , e 1008611 .	4.7	21
587	High prevalence of four novel astrovirus genotype species identified from rodents in China. Journal of General Virology, 2017, 98, 1004-1015.	2.9	21
588	The Management of the 2009 pandemic Influenza A H1N1 virus infection. Journal of Thoracic Disease, 2012, 4, 4-6.	1.4	21
589	A Case Report of Anaerobiospirillum Causing Septicemia. Journal of Infectious Diseases, 1989, 159, 153-154.	4.0	20
590	Association of Presence of Aspergillus Antibodies with Hemoptysis in Patients with Old Tuberculosis or Bronchiectasis but No Radiologically Visible Mycetoma. Journal of Clinical Microbiology, 2004, 42, 665-669.	3.9	20
591	Direct detection of Mycobacterium tuberculosis in clinical specimens using single-tube biotinylated nested polymerase chain reaction-enzyme linked immunoassay (PCR-ELISA). Diagnostic Microbiology and Infectious Disease, 2004, 48, 271-275.	1.8	20
592	The use of high-resolution melting analysis for rapid spa typing on methicillin-resistant Staphylococcus aureus clinical isolates. Journal of Microbiological Methods, 2013, 92, 99-102.	1.6	20
593	Characterization of a Tsukamurella Pseudo-Outbreak by Phenotypic Tests, 16S rRNA Sequencing, Pulsed-Field Gel Electrophoresis, and Metabolic Footprinting. Journal of Clinical Microbiology, 2013, 51, 334-338.	3.9	20
594	Recombinant influenza A virus hemagglutinin HA2 subunit protects mice against influenza A(H7N9) virus infection. Archives of Virology, 2015, 160, 777-786.	2.1	20

#	Article	IF	CITATIONS
595	A nurse-delivered brief health education intervention to improve pneumococcal vaccination rate among older patients with chronic diseases: A cluster randomized controlled trial. International Journal of Nursing Studies, 2015, 52, 317-324.	5.6	20
596	Screening of an FDA-Approved Drug Library with a Two-Tier System Identifies an Entry Inhibitor of Severe Fever with Thrombocytopenia Syndrome Virus. Viruses, 2019, 11, 385.	3.3	20
597	Identification and characterization of $\langle scp \rangle GLDC \langle scp \rangle$ as host susceptibility gene to severe influenza. EMBO Molecular Medicine, 2019, 11, .	6.9	20
598	Intra-host non-synonymous diversity at a neutralizing antibody epitope of SARS-CoV-2 spike protein N-terminal domain. Clinical Microbiology and Infection, 2021, 27, 1350.e1-1350.e5.	6.0	20
599	A Double-blind, Randomized Phase 2 Controlled Trial of Intradermal Hepatitis B Vaccination With a Topical Toll-like Receptor 7 Agonist Imiquimod, in Patients on Dialysis. Clinical Infectious Diseases, 2021, 73, e304-e311.	5.8	20
600	Isolation of MERS-related coronavirus from lesser bamboo bats that uses DPP4 and infects human-DPP4-transgenic mice. Nature Communications, 2021, 12, 216.	12.8	20
601	Low dose inocula of SARS-CoV-2 Alpha variant transmits more efficiently than earlier variants in hamsters. Communications Biology, 2021, 4, 1102.	4.4	20
602	Bats host diverse parvoviruses as possible origin of mammalian dependoparvoviruses and source for bat–swine interspecies transmission. Journal of General Virology, 2017, 98, 3046-3059.	2.9	20
603	Probable Animal-to-Human Transmission of Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) Delta Variant AY.127 Causing a Pet Shop-Related Coronavirus Disease 2019 (COVID-19) Outbreak in Hong Kong. Clinical Infectious Diseases, 2022, 75, e76-e81.	5.8	20
604	Early diagnosis of Exophiala CAPD peritonitis by 18S ribosomal RNA gene sequencing and its clinical significance. Diagnostic Microbiology and Infectious Disease, 2003, 46, 95-102.	1.8	19
605	Typhoid Fever Associated with Acute Appendicitis Caused by an H1-j Strain of Salmonella enteric a Serotype Typhi. Journal of Clinical Microbiology, 2005, 43, 1470-1472.	3.9	19
606	A novel approach for screening immunogenic proteins inPenicillium marneffeiusing the ΔAFMP1ΔAFMP2deletion mutant ofAspergillus fumigatus. FEMS Microbiology Letters, 2006, 262, 138-147	7.1.8	19
607	Detection of Babesia hongkongensis sp. nov. in a Free-Roaming Felis catus Cat in Hong Kong. Journal of Clinical Microbiology, 2012, 50, 2799-2803.	3.9	19
608	Phaeoacremonium parasiticum invasive infections and airway colonization characterized by agar block smear and ITS and \hat{l}^2 -tubulin gene sequencing. Diagnostic Microbiology and Infectious Disease, 2012, 74, 190-197.	1.8	19
609	Suboptimal Humoral Immune Response against Influenza A(H7N9) Virus Is Related to Its Internal Genes. Vaccine Journal, 2015, 22, 1235-1243.	3.1	19
610	An A14U Substitution in the $3\hat{a} \in \mathbb{Z}^2$ Noncoding Region of the M Segment of Viral RNA Supports Replication of Influenza Virus with an NS1 Deletion by Modulating Alternative Splicing of M Segment mRNAs. Journal of Virology, 2015, 89, 10273-10285.	3.4	19
611	Targeting SUMO Modification of the Non-Structural Protein 5 of Zika Virus as a Host-Targeting Antiviral Strategy. International Journal of Molecular Sciences, 2019, 20, 392.	4.1	19
612	Improved Detection of Antibodies against SARS-CoV-2 by Microsphere-Based Antibody Assay. International Journal of Molecular Sciences, 2020, 21, 6595.	4.1	19

#	Article	IF	CITATIONS
613	Nanopore Sequencing Reveals Novel Targets for Detection and Surveillance of Human and Avian Influenza A Viruses. Journal of Clinical Microbiology, 2020, 58, .	3.9	19
614	Orally administered bismuth drug together with <i>N</i> -acetyl cysteine as a broad-spectrum anti-coronavirus cocktail therapy. Chemical Science, 2022, 13, 2238-2248.	7.4	19
615	Biotyping of Penicillium marneffei Reveals Concentration-Dependent Growth Inhibition by Galactose. Journal of Clinical Microbiology, 2001, 39, 1416-1421.	3.9	18
616	Potential Benefit of Plasma Exchange in Treatment of Severe Icteric Leptospirosis Complicated by Acute Renal Failure. Vaccine Journal, 2002, 9, 482-484.	3.1	18
617	Unusual Laboratory Findings in a Case of Norwegian Scabies Provided a Clue to Diagnosis. Journal of Clinical Microbiology, 2005, 43, 2542-2544.	3.9	18
618	In silico analysis of 16S ribosomal RNA gene sequencing-based methods for identification of medically important anaerobic bacteria. Journal of Clinical Pathology, 2006, 60, 576-579.	2.0	18
619	Development of a multi-locus sequence typing scheme for Laribacter hongkongensis, a novel bacterium associated with freshwater fish-borne gastroenteritis and traveler's diarrhea. BMC Microbiology, 2009, 9, 21.	3.3	18
620	Development of monoclonal antibody-based galactomannoprotein antigen-capture ELISAs to detect Aspergillus fumigatus infection in the invasive aspergillosis rabbit models. European Journal of Clinical Microbiology and Infectious Diseases, 2012, 31, 2943-2950.	2.9	18
621	Novel Variant of Beilong Paramyxovirus in Rats, China. Emerging Infectious Diseases, 2012, 18, 1022-1024.	4.3	18
622	Re-Annotation of Protein-Coding Genes in 10 Complete Genomes of Neisseriaceae Family by Combining Similarity-Based and Composition-Based Methods. DNA Research, 2013, 20, 273-286.	3.4	18
623	Successful control of emerging vancomycin-resistant enterococci by territory-wide implementation of directly observed hand hygiene in patients in Hong Kong. American Journal of Infection Control, 2016, 44, 1168-1171.	2.3	18
624	Identification and interspecies transmission of a novel bocaparvovirus among different bat species in China. Journal of General Virology, 2016, 97, 3345-3358.	2.9	18
625	<i>Anaerospora hongkongensis</i> Gen. Nov. Sp. Nov., a Novel Genus and Species with Ribosomal DNA Operon Heterogeneity Isolated from an Intravenous Drug Abuser with Pseudobacteremia. Microbiology and Immunology, 2005, 49, 31-39.	1.4	17
626	First Report of <i>Gordonibacter pamelaeae</i> Bacteremia. Journal of Clinical Microbiology, 2010, 48, 319-322.	3.9	17
627	Automated Pangenomic Analysis in Target Selection for PCR Detection and Identification of Bacteria by Use of ssGeneFinder Webserver and Its Application to Salmonella enterica Serovar Typhi. Journal of Clinical Microbiology, 2012, 50, 1905-1911.	3.9	17
628	A novel dromedary camel enterovirus in the family Picornaviridae from dromedaries in the Middle East. Journal of General Virology, 2015, 96, 1723-1731.	2.9	17
629	Effect of proactive infection control measures on benchmarked rate of hospital outbreaks: An analysis of public hospitals in Hong Kong over 5Âyears. American Journal of Infection Control, 2015, 43, 965-970.	2.3	17
630	Human H7N9 virus induces a more pronounced pro-inflammatory cytokine but an attenuated interferon response in human bronchial epithelial cells when compared with an epidemiologically-linked chicken H7N9 virus. Virology Journal, 2016, 13, 42.	3.4	17

#	Article	IF	Citations
631	Prostaglandin E2-Mediated Impairment of Innate Immune Response to A(H1N1)pdm09 Infection in Diet-Induced Obese Mice Could Be Restored by Paracetamol. Journal of Infectious Diseases, 2019, 219, 795-807.	4.0	17
632	Mammalian cells use the autophagy process to restrict avian influenza virus replication. Cell Reports, 2021, 35, 109213.	6.4	17
633	Low Environmental Temperature Exacerbates Severe Acute Respiratory Syndrome Coronavirus 2 Infection in Golden Syrian Hamsters. Clinical Infectious Diseases, 2022, 75, e1101-e1111.	5.8	17
634	Proactive infection control measures to prevent nosocomial transmission of carbapenem-resistant Enterobacteriaceae in a non-endemic area. Chinese Medical Journal, 2013, 126, 4504-9.	2.3	17
635	Direct Detection of Mycobacterium tuberculosis in Respiratory Specimens Using an Automated DNA Amplification Assay and a Single Tube Nested Polymerase Chain Reaction (PCR). Clinical Chemistry and Laboratory Medicine, 1998, 36, 597-9.	2.3	16
636	Leukocytoclastic vasculitis complicating Klebsiella pneumoniae bacteremia. Diagnostic Microbiology and Infectious Disease, 2000, 37, 275-277.	1.8	16
637	Treatment of multidrug-resistant and extensively drug-resistant tuberculosis: current status and future prospects. Expert Review of Clinical Pharmacology, 2009, 2, 405-421.	3.1	16
638	Matrix-assisted laser desorption ionisation–time of flight mass spectrometry for rapid identification of Laribacter hongkongensis. Journal of Clinical Pathology, 2013, 66, 1081-1083.	2.0	16
639	Fatal Empyema Thoracis Caused by Schizophyllum commune with Cross-Reactive Cryptococcal Antigenemia. Journal of Clinical Microbiology, 2014, 52, 683-687.	3.9	16
640	Molecular identification of cestodes and nematodes by cox1 gene real-time PCR and sequencing. Diagnostic Microbiology and Infectious Disease, 2017, 89, 185-190.	1.8	16
641	Absence of Vaccine-enhanced Disease With Unexpected Positive Protection Against severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) by Inactivated Vaccine Given Within 3 Days of Virus Challenge in Syrian Hamster Model. Clinical Infectious Diseases, 2021, 73, e719-e734.	5.8	16
642	Fusion-inhibition peptide broadly inhibits influenza virus and SARS-CoV-2, including Delta and Omicron variants. Emerging Microbes and Infections, 2022, 11, 926-937.	6.5	16
643	A single dose of oral DNA immunization delivered by attenuated Salmonella typhimurium down-regulates transgene expression in HBsAg transgenic mice. European Journal of Immunology, 2002, 32, 3294-3304.	2.9	15
644	Molecular Characterization of a Strain of Group A Streptococcus Isolated from a Patient with a Psoas Abscess. Journal of Clinical Microbiology, 2003, 41, 4888-4891.	3.9	15
645	Two Cases of Continuous Ambulatory Peritoneal Dialysis-Associated Peritonitis Due to Plesiomonas shigelloides. Journal of Clinical Microbiology, 2004, 42, 933-935.	3.9	15
646	Lymphocyte surge as a marker for immunorestitution disease due to Pneumocystis jiroveci pneumonia in HIV-negative immunosuppressed hosts. European Journal of Clinical Microbiology and Infectious Diseases, 2004, 23, 512-514.	2.9	15
647	Clinical, phenotypic, and genotypic evidence for Streptococcus sinensis as the common ancestor of anginosus and mitis groups of streptococci. Medical Hypotheses, 2006, 66, 345-351.	1.5	15
648	Structural basis and sequence co-evolution analysis of the hemagglutinin protein of pandemic influenza A/H1N1 (2009) virus. Experimental Biology and Medicine, 2011, 236, 915-925.	2.4	15

#	Article	IF	CITATIONS
649	Direct detection of isoniazid-resistant Mycobacterium tuberculosis in respiratory specimens by multiplex allele-specific polymerase chain reaction. Diagnostic Microbiology and Infectious Disease, 2011, 69, 51-58.	1.8	15
650	Complete Genome Sequences of Novel Rat Noroviruses in Hong Kong. Journal of Virology, 2012, 86, 12435-12436.	3.4	15
651	Metabolomic Profiling of Plasma from Melioidosis Patients Using UHPLC-QTOF MS Reveals Novel Biomarkers for Diagnosis. International Journal of Molecular Sciences, 2016, 17, 307.	4.1	15
652	The role of nuclear NS1 protein in highly pathogenic H5N1 influenza viruses. Microbes and Infection, 2017, 19, 587-596.	1.9	15
653	Low population serum microneutralization antibody titer against the predominating influenza A(H3N2) N121K virus during the severe influenza summer peak of Hong Kong in 2017. Emerging Microbes and Infections, 2018, 7, 1-9.	6.5	15
654	Live attenuated Salmonella typhimurium vaccines delivering SaEsxA and SaEsxB via type III secretion system confer protection against Staphylococcus aureus infection. BMC Infectious Diseases, 2018, 18, 195.	2.9	15
655	Novel Bat Alphacoronaviruses in Southern China Support Chinese Horseshoe Bats as an Important Reservoir for Potential Novel Coronaviruses. Viruses, 2019, 11, 423.	3.3	15
656	Comparative Transcriptomic Analysis of Rhinovirus and Influenza Virus Infection. Frontiers in Microbiology, 2020, 11, 1580.	3.5	15
657	Middle East Respiratory Syndrome Coronavirus Antibodies in Bactrian and Hybrid Camels from Dubai. MSphere, 2020, 5, .	2.9	15
658	Polymorphisms of CCR5 gene in a southern Chinese population and their effects on disease progression in HIV infections. Aids, 2002, 16, 2480-2482.	2,2	15
659	Early Treatment of High-Risk Hospitalized Coronavirus Disease 2019 (COVID-19) Patients With a Combination of Interferon Beta-1b and Remdesivir: A Phase 2 Open-label Randomized Controlled Trial. Clinical Infectious Diseases, 2023, 76, e216-e226.	5.8	15
660	Construction of an inducible expression shuttle vector for Laribacter hongkongensis, a novel bacterium associated with gastroenteritis. FEMS Microbiology Letters, 2005, 252, 57-65.	1.8	14
661	First report of methicillin-resistant Staphylococcus aureus septic arthritis complicating acupuncture: simple procedure resulting in most devastating outcome. Diagnostic Microbiology and Infectious Disease, 2009, 63, 92-95.	1.8	14
662	Lactobacillus rhamnosus hepatic abscess associated with Mirizzi syndrome: a case report and review of the literature. Diagnostic Microbiology and Infectious Disease, 2010, 66, 94-97.	1.8	14
663	Analysis of multilocus sequence typing schemes for 35 different bacteria revealed that gene loci of 10 bacteria could be replaced to improve cost-effectiveness. Diagnostic Microbiology and Infectious Disease, 2011, 70, 316-323.	1.8	14
664	Molecular Characterization of a Catalase-Negative Staphylococcus aureus subsp.aureusStrain Collected from a Patient with Mitral Valve Endocarditis and Pericarditis Revealed a Novel Nonsense Mutation in thekatAGene. Journal of Clinical Microbiology, 2011, 49, 3398-3402.	3.9	14
665	Minimal Intervention for Controlling Nosocomial Transmission of Methicillin-Resistant Staphylococcus aureus in Resource Limited Setting with High Endemicity. PLoS ONE, 2014, 9, e100493.	2.5	14
666	A 10-year study reveals clinical and laboratory evidence for the â€~semi-invasive' properties of chronic pulmonary aspergillosis. Emerging Microbes and Infections, 2016, 5, 1-7.	6.5	14

#	Article	IF	CITATIONS
667	The challenge of patient empowerment in hand hygiene promotion in health care facilities in Hong Kong. American Journal of Infection Control, 2017, 45, 562-565.	2.3	14
668	Measuring nonâ€polyaminated lipocalinâ€2 for cardiometabolic risk assessment. ESC Heart Failure, 2017, 4, 563-575.	3.1	14
669	Novel selective medium for the isolation of <i>corynebacterium kroppenstedtii </i> from heavily colonised clinical specimens. Journal of Clinical Pathology, 2018, 71, 781-786.	2.0	14
670	Discovery of a Novel Specific Inhibitor Targeting Influenza A Virus Nucleoprotein with Pleiotropic Inhibitory Effects on Various Steps of the Viral Life Cycle. Journal of Virology, 2021, 95, .	3.4	14
671	Clonorchiasis in Bone Marrow Transplant Recipients. Clinical Infectious Diseases, 1998, 27, 382-384.	5.8	13
672	Bacteremia in a patient with colonic carcinoma caused by a novel Sedimentibacter species: Sedimentibacter hongkongensis sp. nov. Diagnostic Microbiology and Infectious Disease, 2004, 50, 81-87.	1.8	13
673	The severe acute respiratory syndrome (SARS). Journal of NeuroVirology, 2005, 11, 455-468.	2.1	13
674	Surgical site abscess caused by Lactobacillus fermentum identified by 16S ribosomal RNA gene sequencing. Diagnostic Microbiology and Infectious Disease, 2007, 58, 251-254.	1.8	13
675	Virulence determinants, drug resistance and mobile genetic elements of Laribacter hongkongensis: a genome-wide analysis. Cell and Bioscience, 2011, 1, 17.	4.8	13
676	Environmental adaptability and stress tolerance of Laribacter hongkongensis: a genome-wide analysis. Cell and Bioscience, $2011, 1, 22$.	4.8	13
677	Structural and functional insight into the mechanism of an alkaline exonuclease from Laribacter hongkongensis. Nucleic Acids Research, 2011, 39, 9803-9819.	14.5	13
678	Anti-ganglioside antibodies were not detected in human subjects infected with or vaccinated against 2009 pandemic influenza A (H1N1) virus. Vaccine, 2012, 30, 2605-2610.	3.8	13
679	Infections associated with body modification. Journal of the Formosan Medical Association, 2012, 111, 667-681.	1.7	13
680	Mycobacterium abscessus Bacteremia After Receipt of Intravenous Infusate of Cytokine-Induced Killer Cell Therapy for Body Beautification and Health Boosting. Clinical Infectious Diseases, 2013, 57, 981-991.	5.8	13
681	Structure of the S1 subunit C-terminal domain from bat-derived coronavirus HKU5 spike protein. Virology, 2017, 507, 101-109.	2.4	13
682	Gut-homing \hat{l} "42PD1+ \hat{V} "2 T cells promote innate mucosal damage via TLR4 during acute HIV type 1 infection. Nature Microbiology, 2017, 2, 1389-1402.	13.3	13
683	Evaluation of NxTAG Respiratory Pathogen Panel and Comparison with xTAG Respiratory Viral Panel Fast v2 and Film Array Respiratory Panel for Detecting Respiratory Pathogens in Nasopharyngeal Aspirates and Swine/Avian-Origin Influenza A Subtypes in Culture Isolates. Advances in Virology, 2017, 2017, 1-8.	1.1	13
684	Large-scale sequence analysis reveals novel human-adaptive markers in PB2 segment of seasonal influenza A viruses. Emerging Microbes and Infections, 2018, 7, 1-12.	6.5	13

#	Article	IF	CITATIONS
685	Molecular epidemiology of coxsackievirus A6 circulating in Hong Kong reveals common neurological manifestations and emergence of novel recombinant groups. Journal of Clinical Virology, 2018, 108, 43-49.	3.1	13
686	Nosocomial transmission of chickenpox and varicella zoster virus seroprevalence rate amongst healthcare workers in a teaching hospital in China. BMC Infectious Diseases, 2019, 19, 582.	2.9	13
687	<i>Talaromyces marneffei</i> Mp1 Protein, a Novel Virulence Factor, Carries Two Arachidonic Acid-Binding Domains To Suppress Inflammatory Responses in Hosts. Infection and Immunity, 2019, 87, .	2.2	13
688	Molecular Evolution of Human Coronavirus 229E in Hong Kong and a Fatal COVID-19 Case Involving Coinfection with a Novel Human Coronavirus 229E Genogroup. MSphere, 2021, 6, .	2.9	13
689	A new class of \hat{l}_{\pm} -ketoamide derivatives with potent anticancer and anti-SARS-CoV-2 activities. European Journal of Medicinal Chemistry, 2021, 215, 113267.	5.5	13
690	Haemophilus segnis polymicrobial and monomicrobial bacteraemia identified by 16S ribosomal RNA gene sequencing. Journal of Medical Microbiology, 2002, 51, 635-640.	1.8	13
691	Two novel dromedary camel bocaparvoviruses from dromedaries in the Middle East with unique genomic features. Journal of General Virology, 2017, 98, 1349-1359.	2.9	13
692	Capnocytophaga sputigena primary iliopsoas abscess. Journal of Medical Microbiology, 2010, 59, 1368-1370.	1.8	13
693	Reduction of Platelet Transfusion– Associated Sepsis by Short–Term Bacterial Culture. Vox Sanguinis, 1999, 77, 1-5.	1.5	13
694	Assessment of SARS-CoV-2 Immunity in Convalescent Children and Adolescents. Frontiers in Immunology, 2021, 12, 797919.	4.8	13
695	A Palmitic Acid-Conjugated, Peptide-Based pan-CoV Fusion Inhibitor Potently Inhibits Infection of SARS-CoV-2 Omicron and Other Variants of Concern. Viruses, 2022, 14, 549.	3.3	13
696	Explosive outbreak of SARS-CoV-2 Omicron variant is associated with vertical transmission in high-rise residential buildings in Hong Kong. Building and Environment, 2022, 221, 109323.	6.9	13
697	Enhancement by Ampicillin of Antibody Responses Induced by a Protein Antigen and a DNA Vaccine Carried by Live-Attenuated Salmonella enterica Serovar Typhi. Vaccine Journal, 2000, 7, 596-599.	2.6	12
698	Cryptococcal infection associated with fludarabine therapy. American Journal of Medicine, 2000, 108, 523-524.	1.5	12
699	Antigenicity and transmissibility of a novel clade 2.3.2.1 avian influenza H5N1 virus. Journal of General Virology, 2013, 94, 2616-2626.	2.9	12
700	Fatal anti-aquaporin-4 seropositive neuromyelitis optica spectrum disorder in tuberculosis. BMC Infectious Diseases, 2014, 14, 470.	2.9	12
701	Metabolomic profiling of Burkholderia pseudomallei using UHPLC-ESI-Q-TOF-MS reveals specific biomarkers including 4-methyl-5-thiazoleethanol and unique thiamine degradation pathway. Cell and Bioscience, 2015, 5, 26.	4.8	12
702	The Therapeutic Effect of Pamidronate on Lethal Avian Influenza A H7N9 Virus Infected Humanized Mice. PLoS ONE, 2015, 10, e0135999.	2.5	12

#	Article	IF	CITATIONS
703	A DNA vaccine targeting TcdA and TcdB induces protective immunity against Clostridium difficile. BMC Infectious Diseases, 2016, 16, 596.	2.9	12
704	Molecular epidemiology of canine picornavirus in Hong Kong and Dubai and proposal of a novel genus in Picornaviridae. Infection, Genetics and Evolution, 2016, 41, 191-200.	2.3	12
705	Comparative genome and evolutionary analysis of naturally occurring Beilong virus in brown and black rats. Infection, Genetics and Evolution, 2016, 45, 311-319.	2.3	12
706	Novel residues in the PA protein of avian influenza H7N7 virus affect virulence in mammalian hosts. Virology, 2016, 498, 1-8.	2.4	12
707	Ongoing transmission of avian influenza A viruses in Hong Kong despite very comprehensive poultry control measures: A prospective seroepidemiology study. Journal of Infection, 2016, 72, 207-213.	3.3	12
708	Immunotherapy Targeting Adenosine Synthase A Decreases Severity of Staphylococcus aureus Infection in Mouse Model. Journal of Infectious Diseases, 2017, 216, 245-253.	4.0	12
709	Comparative evaluation of a laboratory-developed real-time PCR assay and RealStar \hat{A}^{\otimes} Adenovirus PCR Kit for quantitative detection of human adenovirus. Virology Journal, 2018, 15, 149.	3.4	12
710	Epidemiological and Clinical Characteristics of Human Hepegivirus 1 Infection in Patients With Hepatitis C. Open Forum Infectious Diseases, 2019, 6, ofz329.	0.9	12
711	<i>In silico</i> structure-based discovery of a SARS-CoV-2 main protease inhibitor. International Journal of Biological Sciences, 2021, 17, 1555-1564.	6.4	12
712	Inhaled Dry Powder Formulation of Tamibarotene, a Broadâ€Spectrum Antiviral against Respiratory Viruses Including SARSâ€CoVâ€2 and Influenza Virus. Advanced Therapeutics, 2021, 4, 2100059.	3.2	12
713	Cloning and characterisation of malE in Burkholderia pseudomallei. Journal of Medical Microbiology, 2001, 50, 330-338.	1.8	12
714	Extensive contact tracing and screening to control the spread of vancomycin-resistant Enterococcus faecium ST414 in Hong Kong. Chinese Medical Journal, 2012, 125, 3450-7.	2.3	12
715	A nasal omicron vaccine booster elicits potent neutralizing antibody response against emerging SARS-CoV-2 variants. Emerging Microbes and Infections, 2022, 11, 964-967.	6.5	12
716	Pathogenicity of SARSâ€CoVâ€⊋ Omicron. Clinical and Translational Medicine, 2022, 12, e880.	4.0	12
717	Decreased Antibiotic Consumption Coincided with Reduction in Bacteremia Caused by Bacterial Species with Respiratory Transmission Potential during the COVID-19 Pandemic. Antibiotics, 2022, 11, 746.	3.7	12
718	Phenotypic and Molecular Characterization of Erythromycin Resistance in Four Isolates of Streptococcus -Like Gram-Positive Cocci Causing Bacteremia. Journal of Clinical Microbiology, 2004, 42, 3303-3305.	3.9	11
719	Lack of Evidence that DNA in Antibiotic Preparations Is a Source of Antibiotic Resistance Genes in Bacteria from Animal or Human Sources. Antimicrobial Agents and Chemotherapy, 2004, 48, 3141-3146.	3.2	11
720	Misidentification of a Mucoid Strain of Salmonella enterica Serotype Choleraesuis as Hafnia alvei by the Vitek GNI+ Card System. Journal of Clinical Microbiology, 2006, 44, 4605-4608.	3.9	11

#	Article	IF	CITATIONS
721	Antiviral therapy for respiratory tract infections. Respirology, 2008, 13, 950-971.	2.3	11
722	In silico analysis of 16S rRNA gene sequencing based methods for identification of medically important aerobic Gram-negative bacteria. Journal of Medical Microbiology, 2011, 60, 1281-1286.	1.8	11
723	A method to generate recombinant Salmonella typhi Ty21a strains expressing multiple heterologous genes using an improved recombineering strategy. Applied Microbiology and Biotechnology, 2011, 91, 177-188.	3.6	11
724	First Report of Brain Abscess Caused by a Satelliting Phenotypic Variant of Helcococcus kunzii. Journal of Clinical Microbiology, 2014, 52, 370-373.	3.9	11
725	Fatal Systemic Necrotizing Infections Associated with a Novel Paramyxovirus, Anaconda Paramyxovirus, in Green Anaconda Juveniles. Journal of Clinical Microbiology, 2014, 52, 3614-3623.	3.9	11
726	Role of Hand Hygiene Ambassador and Implementation of Directly Observed Hand Hygiene Among Residents in Residential Care Homes for the Elderly in Hong Kong. Infection Control and Hospital Epidemiology, 2018, 39, 571-577.	1.8	11
727	Measles outbreak from Hong Kong International Airport to the hospital due to secondary vaccine failure in healthcare workers. Infection Control and Hospital Epidemiology, 2019, 40, 1407-1415.	1.8	11
728	A novel linker-immunodominant site (LIS) vaccine targeting the SARS-CoV-2 spike protein protects against severe COVID-19 in Syrian hamsters. Emerging Microbes and Infections, 2021, 10, 874-884.	6.5	11
729	Mining of linear B cell epitopes of SARS-CoV-2 ORF8 protein from COVID-19 patients. Emerging Microbes and Infections, 2021, 10, 1016-1023.	6.5	11
730	Neurosensory Rehabilitation and Olfactory Network Recovery in Covid-19-Related Olfactory Dysfunction. Brain Sciences, 2021, 11, 686.	2.3	11
731	Adenosine synthase A contributes to recurrent Staphylococcus aureus infection by dampening protective immunity. EBioMedicine, 2021, 70, 103505.	6.1	11
732	Hemophagocytosis in the peripheral blood due to tuberculosis mycobacteremia. American Journal of Medicine, 2005, 118, 1298-1299.	1.5	10
733	False-negative cerebrospinal fluid cryptococcal antigen test due to small-colony variants of Cryptococcus neoformans meningitis in a patient with cystopleural shunt. Scandinavian Journal of Infectious Diseases, 2006, 38, 1110-1114.	1.5	10
734	Susceptibility patterns of clinical and fish isolates of Laribacter hongkongensis: comparison of the Etest, disc diffusion and broth microdilution methods. Journal of Antimicrobial Chemotherapy, 2009, 63, 704-708.	3.0	10
735	Resequencing microarray for detection of human adenoviruses in patients with conjunctivitis. Journal of Clinical Virology, 2010, 47, 282-285.	3.1	10
736	General metabolism of Laribacter hongkongensis: a genome-wide analysis. Cell and Bioscience, 2011, 1, 16.	4.8	10
737	Hyperimmune IV Immunoglobulin Treatment of 2009 Influenza A(H1N1): Response. Chest, 2013, 144, 712-713.	0.8	10
738	A Replicating Modified Vaccinia Tiantan Strain Expressing an Avian-Derived Influenza H5N1 Hemagglutinin Induce Broadly Neutralizing Antibodies and Cross-Clade Protective Immunity in Mice. PLoS ONE, 2013, 8, e83274.	2.5	10

#	Article	IF	Citations
739	Rapid reduction of viruria and stabilization of allograft function by fusidic acid in a renal transplant recipient with JC virus-associated nephropathy. Infection, 2015, 43, 577-581.	4.7	10
740	First Report of Human Infection by Agromycesmediolanus, a Gram-Positive Organism Found in Soil. Journal of Clinical Microbiology, 2015, 53, 3377-3379.	3.9	10
741	False-positive SARS-CoV-2 serology in 3 children with Kawasaki disease. Diagnostic Microbiology and Infectious Disease, 2020, 98, 115141.	1.8	10
742	Polyclonal <i>Burkholderia cepacia</i> Complex Outbreak in Peritoneal Dialysis Patients Caused by Contaminated Aqueous Chlorhexidine. Emerging Infectious Diseases, 2020, 26, 1987-1997.	4.3	10
743	Interferon-gamma inhibits influenza A virus cellular attachment by reducing sialic acid cluster size. IScience, 2022, 25, 104037.	4.1	10
744	Functional Analysis of Naturally Occurring Mutations in the Open Reading Frame of CCR5 in HIV-Infected Chinese Patients and Healthy Controls. Journal of Acquired Immune Deficiency Syndromes (1999), 2005, 38, 509-517.	2.1	9
745	Crystal Structure of the Mp1p Ligand Binding Domain 2 Reveals Its Function as a Fatty Acid-binding Protein. Journal of Biological Chemistry, 2010, 285, 9211-9220.	3.4	9
746	Comparative immunological evaluation of recombinant Salmonella Typhimurium strains expressing model antigens as live oral vaccines. BMC Immunology, 2012, 13, 54.	2.2	9
747	Identification of Novel Rosavirus Species That Infects Diverse Rodent Species and Causes Multisystemic Dissemination in Mouse Model. PLoS Pathogens, 2016, 12, e1005911.	4.7	9
748	The importance of matrix-assisted laser desorption ionization–time of flight mass spectrometry for correct identification of Clostridium difficile isolated from chromID C. difficile chromogenic agar. Journal of Microbiology, Immunology and Infection, 2017, 50, 723-726.	3.1	9
749	Rapid Differentiation of Haemophilus influenzae and Haemophilus haemolyticus by Use of Matrix-Assisted Laser Desorption Ionization–Time of Flight Mass Spectrometry with ClinProTools Mass Spectrum Analysis. Journal of Clinical Microbiology, 2017, 55, 2679-2685.	3.9	9
750	The Comeback of Scarlet Fever. EBioMedicine, 2018, 28, 7-8.	6.1	9
751	Triple combination of FDA-approved drugs including flufenamic acid, clarithromycin and zanamivir improves survival of severe influenza in mice. Archives of Virology, 2018, 163, 2349-2358.	2.1	9
752	Targeting the Inositol-Requiring Enzyme-1 Pathway Efficiently Reverts Zika Virus-Induced Neurogenesis and Spermatogenesis Marker Perturbations. ACS Infectious Diseases, 2020, 6, 1745-1758.	3.8	9
753	Human rhinovirus C: a newly discovered human rhinovirus species. Emerging Health Threats Journal, 2010, 3, 7106.	3.0	9
754	Gastrointestinal colonization of meticillin-resistant Staphylococcus aureus: an unrecognized burden upon hospital infection control. Journal of Hospital Infection, 2022, 121, 65-74.	2.9	9
755	Correlation of Immunogenicity and Reactogenicity of BNT162b2 and CoronaVac SARS-CoV-2 Vaccines. MSphere, 2022, 7, e0091521.	2.9	9
756	Multi-resistant Streptococcus pneumoniae in Hong Kong. Journal of Antimicrobial Chemotherapy, 1990, 25, 721-723.	3.0	8

#	Article	IF	CITATIONS
757	Genomic evidence for antibiotic resistance genes of actinomycetes as origins of antibiotic resistance genes in pathogenic bacteria simply because actinomycetes are more ancestral than pathogenic bacteria. Medical Hypotheses, 2006, 67, 1297-1304.	1.5	8
758	Caecal-caecal intussusception caused by Trichuris trichiura in a young healthy adult. Scandinavian Journal of Infectious Diseases, 2006, 38, 813-815.	1.5	8
759	Ascaris-induced eosinophilic pneumonitis in an HIV-infected patient. Journal of Clinical Pathology, 2006, 60, 202-203.	2.0	8
760	Plasmid profile and construction of a small shuttle vector in Laribacter hongkongensis. Biotechnology Letters, 2007, 29, 1575-1582.	2.2	8
761	Management of an incident of failed sterilization of surgical instruments in a dental clinic in Hong Kong. Journal of the Formosan Medical Association, 2013, 112, 666-675.	1.7	8
762	Viral genome and antiviral drug sensitivity analysis of two patients from a family cluster caused by the influenza A(H7N9) virus in Zhejiang, China, 2013. International Journal of Infectious Diseases, 2014, 29, 254-258.	3.3	8
763	Containment of Clostridium difficile infection without reduction in antimicrobial use in Hong Kong. European Journal of Clinical Microbiology and Infectious Diseases, 2015, 34, 1381-1386.	2.9	8
764	Comparative evaluation of a laboratory developed real-time PCR assay and the RealStar \hat{A}^{\otimes} HHV-6 PCR Kit for quantitative detection of human herpesvirus 6. Journal of Virological Methods, 2017, 246, 112-116.	2.1	8
765	Case of "relapsing―COVID â€19 in a kidney transplant recipient. Nephrology, 2020, 25, 933-936.	1.6	8
766	Two novel noroviruses and a novel norovirus genogroup in California sea lions. Journal of General Virology, 2018, 99, 777-782.	2.9	8
767	hnRNP C modulates MERS-CoV and SARS-CoV-2 replication by governing the expression of a subset of circRNAs and cognitive mRNAs. Emerging Microbes and Infections, 2022, 11, 519-531.	6.5	8
768	Computation of Antigenicity Predicts SARS-CoV-2 Vaccine Breakthrough Variants. Frontiers in Immunology, 2022, 13, 861050.	4.8	8
769	Acute pyelonephritis caused by Mycoplasma hominis. Pathology, 1995, 27, 61-63.	0.6	7
770	Pneumococcal native aortic valve endocarditis with mycotic abdominal aortic aneurysm, paraspinal and iliopsoas abscesses and pneumonia revealing a multiple myeloma. Journal of Medical Microbiology, 2011, 60, 851-855.	1.8	7
771	Production of 2-Aminophenoxazin-3-one by Staphylococcus aureus Causes False-Positive Results in β-Galactosidase Assays. Journal of Clinical Microbiology, 2012, 50, 3780-3782.	3.9	7
772	In memory of Patrick Manson, founding father of tropical medicine and the discovery of vector-borne infections. Emerging Microbes and Infections, 2012, 1, 1-7.	6.5	7
773	Clinical significance of Pneumocystis jiroveci in patients with active tuberculosis. Diagnostic Microbiology and Infectious Disease, 2013, 75, 260-265.	1.8	7
774	PExFInS: An Integrative Post-GWAS Explorer for Functional Indels and SNPs. Scientific Reports, 2015, 5, 17302.	3.3	7

#	Article	IF	CITATIONS
775	Metabolomics Analysis Reveals Specific Novel Tetrapeptide and Potential Anti-Inflammatory Metabolites in Pathogenic Aspergillus species. International Journal of Molecular Sciences, 2015, 16, 13850-13867.	4.1	7
776	The Association Between Methicillin Resistant Staphylococcus aureus Colonization and Mortality in Chinese Nursing Home Older Adults: A 2-Year Prospective Cohort. Journal of the American Medical Directors Association, 2015, 16, 796-797.	2 . 5	7
777	First detection and complete genome sequence of a phylogenetically distinct human polyomavirus 6 highly prevalent in human bile samples. Journal of Infection, 2017, 74, 50-59.	3.3	7
778	Broad and Effective Protection against Staphylococcus aureus Is Elicited by a Multivalent Vaccine Formulated with Novel Antigens. MSphere, 2019, 4, .	2.9	7
779	Inhibition of HBV replication and gene expression in vitro and in vivo with a single AAV vector delivering two shRNA molecules. BMB Reports, 2009, 42, 59-64.	2.4	7
780	Identification and Evaluation of Recombinant Outer Membrane Proteins as Vaccine Candidates Against Klebsiella pneumoniae. Frontiers in Immunology, 2021, 12, 730116.	4.8	7
781	An unprecedented outbreak investigation for nosocomial and community-acquired legionellosis in Hong Kong. Chinese Medical Journal, 2012, 125, 4283-90.	2.3	7
782	Adult-onset immunodeficiency due to anti-interferon-gamma autoantibody in mainland Chinese. Chinese Medical Journal, 2014, 127, 1189-90.	2.3	7
783	Air dispersal of meticillin-resistant Staphylococcus aureus in residential care homes for the elderly: implications for transmission during the COVID-19 pandemic. Journal of Hospital Infection, 2022, 123, 52-60.	2.9	7
784	A trifunctional peptide broadly inhibits SARS-CoV-2 Delta and Omicron variants in hamsters. Cell Discovery, 2022, 8, .	6.7	7
785	Nasopharyngeal Detection of Severe Acute Respiratory Syndrome-Associated Coronavirus RNA in Health-Care Workers. Chest, 2006, 129, 12-13.	0.8	6
786	Asymptomatic Penicillium marneffei fungemia in an HIV-infected patient. International Journal of Infectious Diseases, 2007, 11 , 280-281.	3.3	6
787	Human metapneumovirus infection in an immunocompetent adult presenting as mononucleosis-like illness. Journal of Infection, 2008, 56, 389-392.	3.3	6
788	Detection of Asymptomatic Antigenemia in Pigs Infected by Porcine Reproductive and Respiratory Syndrome Virus (PRRSV) by a Novel Capture Immunoassay with Monoclonal Antibodies against the Nucleocapsid Protein of PRRSV. Vaccine Journal, 2009, 16, 1822-1828.	3.1	6
789	BK Virus Nephropathy Due to KOM-3 Strain. American Journal of Kidney Diseases, 2009, 54, 122-126.	1.9	6
790	Transport genes and chemotaxis in Laribacter hongkongensis: a genome-wide analysis. Cell and Bioscience, 2011, 1, 28.	4.8	6
791	Emerging infectious diseases as a global health threat. Experimental Biology and Medicine, 2011, 236, 897-898.	2.4	6
792	Assessment of Antigen and Molecular Tests with Serial Specimens from a Patient with Influenza A(H7N9) Infection. Journal of Clinical Microbiology, 2014, 52, 2272-2274.	3.9	6

#	Article	IF	CITATIONS
793	Characterization of the antigenicity of Cpl1, a surface protein of <i>Cryptococcus neoformans </i> neoformans). Mycologia, 2015, 107, 39-45.	1.9	6
794	Application of a dual target PCR-high resolution melting (HRM) method for rapid nontuberculous mycobacteria identification. Journal of Microbiological Methods, 2017, 132, 1-3.	1.6	6
795	Impact of inter-genotypic recombination and probe cross-reactivity on the performance of the Abbott RealTime HCV Genotype II assay for hepatitis C genotyping. Diagnostic Microbiology and Infectious Disease, 2018, 91, 34-37.	1.8	6
796	Integrated analysis of mRNA-seq and miRNA-seq for host susceptibilities to influenza A (H7N9) infection in inbred mouse lines. Functional and Integrative Genomics, 2018, 18, 411-424.	3.5	6
797	Assessment of population susceptibility to upcoming seasonal influenza epidemic strain using interepidemic emerging influenza virus strains. Epidemiology and Infection, 2019, 147, e279.	2.1	6
798	Candida Tropicalis renal microabscesses in a child with leukemia confirmed using nucleic acid amplification and recovery after prolonged antifungal and corticosteroid treatment. International Journal of Infectious Diseases, 2019, 81, 110-113.	3.3	6
799	Persistent lentivirus infection induces early myeloid suppressor cells expansion to subvert protective memory CD8 T cell response✰,✰✰. EBioMedicine, 2020, 60, 103008.	6.1	6
800	Repurposing of Miltefosine as an Adjuvant for Influenza Vaccine. Vaccines, 2020, 8, 754.	4.4	6
801	Intradermal vaccination of live attenuated influenza vaccine protects mice against homologous and heterologous influenza challenges. Npj Vaccines, 2021, 6, 95.	6.0	6
802	False Coronavirus Disease 2019 Cases due to Contamination by Inactivated Virus Vaccine. Clinical Infectious Diseases, 2022, 74, 1485-1488.	5.8	6
803	Correlation between Commercial Anti-RBD IgG Titer and Neutralization Titer against SARS-CoV-2 Beta Variant. Diagnostics, 2021, 11, 2216.	2.6	6
804	Verocytotoxinâ€producing <i>Escherichia coli</i> infection: The Hong Kong experience. Journal of Gastroenterology and Hepatology (Australia), 1998, 13, S289-S293.	2.8	5
805	Continuous Ambulatory Peritoneal Dialysis-Related Peritonitis Associated with Lancefield Group G Beta-Hemolytic Streptococcus: Report of Two Cases Requiring Tenckhoff Catheter Removal. Journal of Clinical Microbiology, 2004, 42, 4399-4402.	3.9	5
806	Resequencing microarray for detection of human adenoviruses in patients with community-acquired gastroenteritis: a proof-of-concept study. Journal of Medical Microbiology, 2010, 59, 1387-1390.	1.8	5
807	A new ASPECT for complicated urinary tract infections. Lancet, The, 2015, 385, 1920-1922.	13.7	5
808	Prevention of nosocomial transmission of influenza A (H7N9) in Hong Kong. Journal of Hospital Infection, 2015, 90, 355-356.	2.9	5
809	Development and in-use evaluation of a novel Luminex MicroPlex microsphere-based (TRIOL) assay for simultaneous identification of <i>Mycobacterium tuberculosis</i> and detection of first-line and second-line anti-tuberculous drug resistance in China. Journal of Clinical Pathology, 2017, 70, 342-349.	2.0	5
810	Mp1p homologues as virulence factors in Aspergillus fumigatus. Medical Mycology, 2018, 56, 350-360.	0.7	5

#	Article	IF	Citations
811	Emergence of Cytomegalovirus Mononucleosis Syndrome Among Young Adults in Hong Kong Linked to Falling Seroprevalence: Results of a 14-Year Seroepidemiological Study. Open Forum Infectious Diseases, 2018, 5, ofy262.	0.9	5
812	Control of Carbapenemase-producing Enterobacteriaceae: Beyond the Hospital. EClinicalMedicine, 2018, 6, 3-4.	7.1	5
813	Improving the specific diagnosis of trematode, cestode and nematode infections by a multiplex single-tube real-time PCR assay. Journal of Clinical Pathology, 2019, 72, 487-492.	2.0	5
814	Early triple antiviral therapy for COVID-19 â€" Authors' reply. Lancet, The, 2020, 396, 1488.	13.7	5
815	SPINK6 inhibits human airway serine proteases and restricts influenza virus activation. EMBO Molecular Medicine, 2022, 14, e14485.	6.9	5
816	Multiplex metal-detection based assay (MMDA) for COVID-19 diagnosis and identification of disease severity biomarkers. Chemical Science, 2022, 13, 3216-3226.	7.4	5
817	Subinhibitory Concentrations of Antibiotics Exacerbate Staphylococcal Infection by Inducing Bacterial Virulence. Microbiology Spectrum, 2022, 10, .	3.0	5
818	Air dispersal of respiratory viruses other than severe acute respiratory coronavirus virus 2 (SARS-CoV-2) and the implication on hospital infection control. Infection Control and Hospital Epidemiology, 2023, 44, 768-773.	1.8	5
819	Characterization of a novel insertion sequence, IS Bp1 , in Burkholderia pseudomallei. Archives of Microbiology, 2002, 177, 267-273.	2.2	4
820	Clinical Deterioration in Community Acquired Infections Associated with Lymphocyte Upsurge in Immunocompetent Hosts. Scandinavian Journal of Infectious Diseases, 2004, 36, 743-751.	1.5	4
821	Rapid diagnosis of multidrug-resistant smear-positive pulmonary tuberculosis. International Journal of Antimicrobial Agents, 2010, 35, 202-203.	2.5	4
822	Equine rhinitis B viruses in horse fecal samples from the Middle East. Virology Journal, 2016, 13, 94.	3.4	4
823	Gastrointestinal colonization with multiple New Delhi metallo- \hat{l}^2 -lactamase-producing Enterobacteriaceae isolates in the same patient: a potential challenge in outbreak investigation. Journal of Hospital Infection, 2016, 92, 108-109.	2.9	4
824	Fatal pancytopenia due to albendazole treatment for strongyloidiasis. IDCases, 2018, 12, 112-116.	0.9	4
825	Evaluation of RealStar® Alpha Herpesvirus PCR Kit for Detection of HSV-1, HSV-2, and VZV in Clinical Specimens. BioMed Research International, 2019, 2019, 1-6.	1.9	4
826	Comparative performance of two commercial sample-to-result systems for hepatitis C virus quantitation and genotyping. Expert Review of Molecular Diagnostics, 2020, 20, 1253-1258.	3.1	4
827	The Severe Acute Respiratory Syndrome. , 2007, , 163-193.		4
828	MBEToolbox 2.0: an enhanced version of a MATLAB toolbox for molecular biology and evolution. Evolutionary Bioinformatics, 2007, 2, 179-82.	1.2	4

#	Article	IF	Citations
829	Pyoderma gangrenosum with pulmonary involvement: a pulmonary special report and literature review. Expert Review of Respiratory Medicine, 2022, 16, 149-159.	2.5	4
830	MBEToolbox 2.0: An enhanced version of a MATLAB toolbox for Molecular Biology and Evolution. Evolutionary Bioinformatics, 2006, 2, 117693430600200.	1.2	3
831	A Real-Time Reverse Transcriptase-Polymer Chain Reaction To Evaluate Natural History of Viral Shedding in Outpatient Children and Adolescents With Pandemic 2009 Influenza A(H1N1). Chest, 2010, 138, 457-458.	0.8	3
832	Complication of Corticosteroid Treatment by Acute <i>Plasmodium malariae</i> Infection Confirmed by Small-Subunit rRNA Sequencing. Journal of Clinical Microbiology, 2010, 48, 4313-4316.	3.9	3
833	Brittle tail syndrome is an emerging infection in horses caused by a keratinolytic fungus Equicapillimyces hongkongensis gen. nov., sp. nov. Veterinary Microbiology, 2012, 155, 399-408.	1.9	3
834	Solving the mystery of H7N9 by crystal balls. Cell Research, 2013, 23, 1335-1336.	12.0	3
835	Circulating cytomegalic cells in a patient with advanced HIV presenting with cytomegalovirus rhinosinusitis. Journal of Clinical Virology, 2015, 65, 87-89.	3.1	3
836	Anaconda paramyxovirus infection in an adult green anaconda after prolonged incubation: Pathological characterization and whole genome sequence analysis. Infection, Genetics and Evolution, 2017, 51, 239-244.	2.3	3
837	Variants of SARS Coronavirus-2 and Their Potential Impact on the Future of the COVID-19 Pandemic. Zoonoses, 2021, 1 , .	1.1	3
838	Severe fever with thrombocytopenia syndrome virus (SFTSV)-host interactome screen identifies viral nucleoprotein-associated host factors as potential antiviral targets. Computational and Structural Biotechnology Journal, 2021, 19, 5568-5577.	4.1	3
839	In-House Immunofluorescence Assay for Detection of SARS-CoV-2 Antigens in Cells from Nasopharyngeal Swabs as a Diagnostic Method for COVID-19. Diagnostics, 2021, 11, 2346.	2.6	3
840	Response to Evidence in favor of the essentiality of human cell membrane-bound ACE2 and against soluble ACE2 for SARS-CoV-2 infectivity. Cell, 2022, 185, 1840-1841.	28.9	3
841	Viral Infections, an Overview with a Focus on Prevention of Transmission., 2017,, 368-377.		2
842	Nosocomial transmission of hepatitis C virus in a liver transplant center in Hong Kong: implication of reusable blood collection tube holder as the vehicle for transmission. Infection Control and Hospital Epidemiology, 2018, 39, 1170-1177.	1.8	2
843	Risk of Hepatitis E among Persons Who Inject Drugs in Hong Kong: A Qualitative and Quantitative Serological Analysis. Microorganisms, 2020, 8, 675.	3.6	2
844	Logistical feasibility and potential benefits of a population-wide passive immunotherapy program during an influenza pandemic. Influenza and Other Respiratory Viruses, 2011, 5, 226-9.	3 . 4	2
845	Evaluation of an Antigen Detection Rapid Diagnostic Test for Detection of SARS-CoV-2 in Clinical Samples. Covid, 2021, 1, 775-783.	1.5	2
846	Broad-spectrum Respiratory Virus Entry Inhibitors. Advances in Experimental Medicine and Biology, 2022, 1366, 137-153.	1.6	2

#	Article	IF	Citations
847	Penicillium cbrysogenum Infection in a Cotton Farmer with Acute Myeloid Leukemia. Infectious Diseases in Clinical Practice, 1997, 6, 482-483.	0.3	1
848	Corrigendum to "Genomic and experimental evidence for a potential sexual cycle in the pathogenic thermal dimorphic fungus Penicillium marneffei ―[FEBS Lett. 580 (2006) 3409-3416]. FEBS Letters, 2006, 580, 4976-4977.	2.8	1
849	Prognostication of Methicillin-resistant Staphylococcus Aureus (MRSA) patient survival., 2011,,.		1
850	Backbone and side-chain 1H, 13C and 15N assignments of the PPIase domain of macrophage infectivity potentiator (Mip) protein from Coxiella burnetii. Biomolecular NMR Assignments, 2014, 8, 173-176.	0.8	1
851	No evidence for a superior platform to develop therapeutic antibodies rapidly in response to MERS-CoV and other emerging viruses. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, E5115-E5115.	7.1	1
852	Epstein–Barr virusâ€positive Tâ€cellâ€associated colitis mimicking inflammatory bowel disease: clinicopathological study of two cases. Histopathology, 2016, 68, 465-468.	2.9	1
853	Paired Heavy and Light Chain Signatures Contribute to Potent SARS-CoV-2 Neutralization in Public Antibody Responses. SSRN Electronic Journal, 0, , .	0.4	1
854	A Unique Pattern of Staphylococcal Scalded Skin Syndrome-Like Erosions in Patients with Atopic Dermatitis: Dermatitis flammeus. Skinmed, 2018, 16, 309-313.	0.0	1
855	The impact of personal coaching on influenza vaccination among healthcare workers before and during COVID-19 pandemic. Vaccine, 2022, , .	3.8	1
856	Macrolides as Immunomodulatory Agents: Review and Future Directions. Medicinal Chemistry Reviews Online, 2004, 1, 151-161.	0.1	0
857	Corrigendum to "96 weeks combination of adefovir dipivoxil plus emtricitabine vs. adefovir dipivoxil monotherapy in the treatment of chronic hepatitis Bâ€₊ Journal of Hepatology, 2009, 50, 1283-1284.	3.7	0
858	Pathogenesis of pandemic H1N1 2009 influenza virus infection and the implication on management. Frontiers of Medicine in China, 2010, 4, 147-156.	0.1	0
859	Logistic regression analysis for Predicting Methicillin-resistant Staphylococcus Aureus (MRSA) in-hospital mortality. , 2011, , .		0
860	Disseminated cryptococcosis mimicking a lymphoma. European Journal of Haematology, 2012, 88, 275-276.	2.2	0
861	Reply to Leow. Journal of Infectious Diseases, 2013, 208, 1351-1352.	4.0	0
862	Response. Chest, 2014, 145, 435.	0.8	0
863	Response. Chest, 2018, 154, 999.	0.8	0
864	Another avian influenza A subtype jumping into human: this time is H7N4. Science Bulletin, 2018, 63, 1025-1026.	9.0	0

#	Article	IF	CITATIONS
865	Clinical practice guidelines for the provision of renal service in Hong Kong: Infection Control in Renal Service. Nephrology, 2019, 24, 98-129.	1.6	0
866	Reply. Hepatology, 2020, 72, 1155-1156.	7.3	0
867	In Silico Structure-Based Design of Antiviral Peptides Targeting the Severe Fever with Thrombocytopenia Syndrome Virus Glycoprotein Gn. Viruses, 2021, 13, 2047.	3.3	0
868	Viral Infections, an Overview with a Focus on Prevention of Transmission., 2016,,.		0
869	Title is missing!. , 2020, 16, e1008611.		O
870	Title is missing!. , 2020, 16, e1008611.		0
871	Title is missing!. , 2020, 16, e1008611.		0
872	Title is missing!. , 2020, 16, e1008611.		0