

Fengyu Hu

List of Publications by Year in descending order

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Version: 2024-02-01

51
papers

5,595
citations

471371

17
h-index

214721

47
g-index

54
all docs

54
docs citations

54
times ranked

12914
citing authors

#	ARTICLE	IF	CITATIONS
1	Temporal dynamics in viral shedding and transmissibility of COVID-19. <i>Nature Medicine</i> , 2020, 26, 672-675.	15.2	3,838
2	Detectable 2019-nCoV viral RNA in blood is a strong indicator for the further clinical severity. <i>Emerging Microbes and Infections</i> , 2020, 9, 469-473.	3.0	339
3	Efficacy and Safety of Lopinavir/Ritonavir or Arbidol in Adult Patients with Mild/Moderate COVID-19: An Exploratory Randomized Controlled Trial. <i>Med</i> , 2020, 1, 105-113.e4.	2.2	256
4	Transmission, viral kinetics and clinical characteristics of the emergent SARS-CoV-2 Delta VOC in Guangzhou, China. <i>EClinicalMedicine</i> , 2021, 40, 101129.	3.2	176
5	Human infections with recently-emerging highly pathogenic H7N9 avian influenza virus in China. <i>Journal of Infection</i> , 2017, 75, 71-75.	1.7	143
6	Protective humoral and cellular immune responses to SARS-CoV-2 persist up to 1 year after recovery. <i>Nature Communications</i> , 2021, 12, 4984.	5.8	100
7	Single-cell analysis reveals bronchoalveolar epithelial dysfunction in COVID-19 patients. <i>Protein and Cell</i> , 2020, 11, 680-687.	4.8	75
8	Prognostic Factors for COVID-19 Pneumonia Progression to Severe Symptoms Based on Earlier Clinical Features: A Retrospective Analysis. <i>Frontiers in Medicine</i> , 2020, 7, 557453.	1.2	60
9	COVID-19 induces new-onset insulin resistance and lipid metabolic dysregulation via regulation of secreted metabolic factors. <i>Signal Transduction and Targeted Therapy</i> , 2021, 6, 427.	7.1	55
10	Serum levels of soluble vascular cell adhesion molecules may correlate with the severity of dengue virus-1 infection in adults. <i>Emerging Microbes and Infections</i> , 2015, 4, 1-7.	3.0	37
11	Association between detectable SARS-CoV-2 RNA in anal swabs and disease severity in patients with coronavirus disease 2019. <i>Journal of Medical Virology</i> , 2021, 93, 794-802.	2.5	34
12	Prognostic roles of KL-6 in disease severity and lung injury in COVID-19 patients: A longitudinal retrospective analysis. <i>Journal of Medical Virology</i> , 2021, 93, 2505-2512.	2.5	28
13	Cell Culture Models and Animal Models for HBV Study. <i>Advances in Experimental Medicine and Biology</i> , 2020, 1179, 109-135.	0.8	28
14	Longitudinal Peripheral Blood Transcriptional Analysis Reveals Molecular Signatures of Disease Progression in COVID-19 Patients. <i>Journal of Immunology</i> , 2021, 206, 2146-2159.	0.4	25
15	Analysis of B Cell Receptor Repertoires Reveals Key Signatures of the Systemic B Cell Response after SARS-CoV-2 Infection. <i>Journal of Virology</i> , 2022, 96, JVI0160021.	1.5	24
16	Molecular and serological characterization of SARS-CoV-2 infection among COVID-19 patients. <i>Virology</i> , 2020, 551, 26-35.	1.1	21
17	Transmitted drug resistance and transmission clusters among HIV-1 treatment-naïve patients in Guangdong, China: a cross-sectional study. <i>Virology Journal</i> , 2021, 18, 181.	1.4	16
18	Gemykibivirus Genome in Lower Respiratory Tract of Elderly Woman With Unexplained Acute Respiratory Distress Syndrome. <i>Clinical Infectious Diseases</i> , 2019, 69, 861-864.	2.9	14

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19	Significantly reduced abilities to cross-neutralize SARS-CoV-2 variants by sera from convalescent COVID-19 patients infected by Delta or early strains. <i>Cellular and Molecular Immunology</i> , 2021, 18, 2560-2562.	4.8	11
20	Metagenomic analysis identified co-infection with human rhinovirus C and bocavirus 1 in an adult suffering from severe pneumonia. <i>Journal of Infection</i> , 2018, 76, 311-313.	1.7	10
21	Complicated genotypes circulating among treatment naïve HIV-1 patients in Guangzhou, China. <i>Infection, Genetics and Evolution</i> , 2021, 87, 104673.	1.0	10
22	Factors associated with clinical outcomes in patients with Coronavirus Disease 2019 in Guangzhou, China. <i>Journal of Clinical Virology</i> , 2020, 133, 104661.	1.6	9
23	HIV-1 Drug Resistance and Genetic Transmission Networks Among MSM Failing Antiretroviral Therapy in South China 2014–2019. <i>Infection and Drug Resistance</i> , 2021, Volume 14, 2977-2989.	1.1	9
24	Changing Epidemiology of Hepatitis C Virus Genotype among Patients with Human Immunodeficiency Virus/Hepatitis C Virus Co-Infection in China. <i>PLoS ONE</i> , 2016, 11, e0161844.	1.1	9
25	Characterization of small genomic regions of the hepatitis B virus should be performed with more caution. <i>Virology Journal</i> , 2018, 15, 188.	1.4	8
26	Evidence that the second human pegivirus (HPgV-2) is primarily a lymphotropic virus and can replicate independent of HCV replication. <i>Emerging Microbes and Infections</i> , 2020, 9, 485-495.	3.0	8
27	Glycopeptide Antibiotic Teicoplanin Inhibits Cell Entry of SARS-CoV-2 by Suppressing the Proteolytic Activity of Cathepsin L. <i>Frontiers in Microbiology</i> , 2022, 13, 884034.	1.5	8
28	Naturally occurring antiviral drug resistance in HIV patients who are mono-infected or co-infected with HBV or HCV in China. <i>Journal of Medical Virology</i> , 2018, 90, 1246-1256.	2.5	7
29	Outcome of Sentinel Hospital-based and CDC-based ART Service Delivery: A Prospective Open Cohort of People Living with HIV in China. <i>Scientific Reports</i> , 2017, 7, 42637.	1.6	6
30	Absence of Integrase Inhibitor-Associated Resistance Among Antiretroviral Therapy-Naïve HIV-1-Infected Adults in Guangdong Province, China, in 2018. <i>Infection and Drug Resistance</i> , 2020, Volume 13, 4389-4394.	1.1	6
31	Characteristics of drug resistance in HIV-1 CRF55_01B from ART-experienced patients in Guangdong, China. <i>Journal of Antimicrobial Chemotherapy</i> , 2020, 75, 1925-1931.	1.3	6
32	The Association of TLR2, TLR3, and TLR9 Gene Polymorphisms With Susceptibility to Talaromycosis Among Han Chinese AIDS Patients in Guangdong. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021, 11, 625461.	1.8	6
33	Mucosal Priming with a Recombinant Influenza A Virus-Vectored Vaccine Elicits T-Cell and Antibody Responses to HIV-1 in Mice. <i>Journal of Virology</i> , 2021, 95, .	1.5	6
34	Expression and characterization of a <i>Talaromyces marneffei</i> active phospholipase B expressed in a <i>Pichia pastoris</i> expression system. <i>Emerging Microbes and Infections</i> , 2016, 5, 1-6.	3.0	5
35	Genotypic distribution and hepatic fibrosis among HIV/HCV co-infected individuals in Southern China: a retrospective cross-sectional study. <i>BMC Infectious Diseases</i> , 2015, 15, 401.	1.3	4
36	Dual Therapy with Lopinavir/Ritonavir plus Lamivudine could be a Viable Alternative for Antiretroviral-Therapy-Naïve Adults with HIV-1 Infection Regardless of HIV Viral Load or Subgenotype in Resource-Limited Settings: A Randomised, Open-Label and Non-Inferiority Study from China. <i>Indian Journal of Medical Microbiology</i> , 2018, 36, 513-516.	0.3	4

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37	Genetic characterization of a novel HIV-1 CRF01_AE/CRF07_BC recombinant form among men who have sex with men in Guangdong, China. <i>Archives of Virology</i> , 2018, 163, 3093-3097.	0.9	4
38	A novel one-step quantitative reverse transcription PCR assay for selective amplification of hepatitis B virus pregenomic RNA from a mixture of HBV DNA and RNA in serum. <i>Archives of Virology</i> , 2019, 164, 2683-2690.	0.9	4
39	Difference of Intrahost Dynamics of the Second Human Pegivirus and Hepatitis C Virus in HPgV-2/HCV-Coinfected Patients. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021, 11, 728415.	1.8	4
40	Evolutionary pathways to NS5A inhibitor resistance in genotype 1 hepatitis C virus. <i>Antiviral Research</i> , 2018, 158, 45-51.	1.9	3
41	Reduction of anti-HIV antibody responses in subjects receiving antiretroviral therapy during chronic HIV-1 infection. <i>Journal of Clinical Virology</i> , 2020, 128, 104414.	1.6	3
42	Genetic Characteristics of HIV-1 CRF12_BF First Identified in Guangdong Province, China. <i>AIDS Research and Human Retroviruses</i> , 2021, 37, 157-161.	0.5	3
43	Persistent chronic immune activation in HIV/HBV coinfected patients after antiretroviral therapy. <i>Journal of Viral Hepatitis</i> , 2021, 28, 1355-1361.	1.0	3
44	Characterization of Four Nearly Full-Length Genomic Sequences of HIV-1 Subtype G Identified in Guangdong Province, China. <i>AIDS Research and Human Retroviruses</i> , 2019, 35, 668-672.	0.5	2
45	Near Full-Length Genomic Characterization of a Novel HIV-1 B/C Recombinant Form Identified in Guangdong Province, China. <i>AIDS Research and Human Retroviruses</i> , 2021, 37, 566-571.	0.5	1
46	Impact of HBeAg status on immune recovery in HIV/HBV co-infected patients after highly active antiretroviral therapy. <i>Infection, Genetics and Evolution</i> , 2021, 91, 104781.	1.0	1
47	<p>Comparison and Correlation of Genetic Variability of the HBV Pre-S Region in HIV/HBV Co-Infected Patients: Quasispecies Perspective</p>. <i>Infection and Drug Resistance</i> , 2020, Volume 13, 4327-4334.	1.1	0
48	Genetic characteristics of HIV-1 CRF06_cpx and CRF56_cpx strains isolated in Guangzhou, China. <i>Archives of Virology</i> , 2021, 166, 2853-2857.	0.9	0
49	Kinetics of Severity Biomarkers and Immunological Features of Methylprednisolone Therapy for Severe COVID-19 Patients. <i>Frontiers in Immunology</i> , 2022, 13, 758946.	2.2	0
50	Distribution of nontuberculous mycobacteria in patients with and without HIV/AIDS in Chongqing. <i>HIV Medicine</i> , 2022, 23, 54-63.	1.0	0
51	OUP accepted manuscript. <i>Journal of Antimicrobial Chemotherapy</i> , 2022, , .	1.3	0