

Fengyu Hu

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

49
papers

3,469
citations

12
h-index

54
g-index

54
ext. papers

4,586
ext. citations

9.7
avg. IF

5.38
L-index

#	Paper	IF	Citations
49	Temporal dynamics in viral shedding and transmissibility of COVID-19. <i>Nature Medicine</i> , 2020 , 26, 672-675	50.5	2459
48	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	18.9	272
47	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	31.7	180
46	Human infections with recently-emerging highly pathogenic H7N9 avian influenza virus in China. <i>Journal of Infection</i> , 2017 , 75, 71-75	18.9	115
45	Transmission, viral kinetics and clinical characteristics of the emergent SARS-CoV-2 Delta VOC in Guangzhou, China. <i>EClinicalMedicine</i> , 2021 , 40, 101129	11.3	62
44	Single-cell analysis reveals bronchoalveolar epithelial dysfunction in COVID-19 patients. <i>Protein and Cell</i> , 2020 , 11, 680-687	7.2	41
43	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	4.9	30
42	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, e24	18.9	27
41	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	19.7	23
40	Protective humoral and cellular immune responses to SARS-CoV-2 persist up to 1 year after recovery. <i>Nature Communications</i> , 2021 , 12, 4984	17.4	20
39	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	19.7	12
38	Cell Culture Models and Animal Models for HBV Study. <i>Advances in Experimental Medicine and Biology</i> , 2020 , 1179, 109-135	3.6	11
37	Molecular and serological characterization of SARS-CoV-2 infection among COVID-19 patients. <i>Virology</i> , 2020 , 551, 26-35	3.6	11
36	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	18.9	9
35	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	19.7	7
34	Longitudinal Peripheral Blood Transcriptional Analysis Reveals Molecular Signatures of Disease Progression in COVID-19 Patients. <i>Journal of Immunology</i> , 2021 , 206, 2146-2159	5.3	7
33	Gemykibivirus Genome in Lower Respiratory Tract of Elderly Woman With Unexplained Acute Respiratory Distress Syndrome. <i>Clinical Infectious Diseases</i> , 2019 , 69, 861-864	11.6	6

32	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	21	6
31	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	4.9	5
30	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	18.9	5
29	Factors associated with clinical outcomes in patients with Coronavirus Disease 2019 in Guangzhou, China. <i>Journal of Clinical Virology</i> , 2020 , 133, 104661	14.5	5
28	Characterization of small genomic regions of the hepatitis B virus should be performed with more caution. <i>Virology Journal</i> , 2018 , 15, 188	6.1	5
27	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	2.6	3
26	Evolutionary pathways to NS5A inhibitor resistance in genotype 1 hepatitis C virus. <i>Antiviral Research</i> , 2018 , 158, 45-51	10.8	3
25	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	2.6	3
24	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	4	3
23	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	3.7	3
22	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, e120	18.9	3
21	Complicated genotypes circulating among treatment naïve HIV-1 patients in Guangzhou, China. <i>Infection, Genetics and Evolution</i> , 2021 , 87, 104673	4.5	3
20	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	1.6	2
19	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 , 13, 4389-4394	4.2	2
18	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	15.4	2
17	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	5.1	2
16	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	1.3	2
15	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	6.1	2

14	Analysis of B cell receptor repertoires reveals key signatures of systemic B cell response after SARS-CoV-2 infection. <i>Journal of Virology</i> , 2021 , JVI0160021	6.6	2
13	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	14.5	1
12	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	5.9	1
11	Genetic Characteristics of HIV-1 CRF12_BF First Identified in Guangdong Province, China. <i>AIDS Research and Human Retroviruses</i> , 2021 , 37, 157-161	1.6	1
10	Difference of Intra-host 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	5.9	1
9	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 , 14, 2977-2989	4.2	1
8	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	5.7	1
7	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	1.6	0
6	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	4.5	0
5	Persistent chronic immune activation in HIV/HBV-coinfected patients after antiretroviral therapy. <i>Journal of Viral Hepatitis</i> , 2021 , 28, 1355-1361	3.4	0
4	Comparison and Correlation of Genetic Variability of the HBV Pre-S Region in HIV/HBV Co-Infected Patients: Quasispecies Perspective. <i>Infection and Drug Resistance</i> , 2020 , 13, 4327-4334	4.2	
3	Genetic characteristics of HIV-1 CRF06_cpx and CRF56_cpx strains isolated in Guangzhou, China. <i>Archives of Virology</i> , 2021 , 166, 2853-2857	2.6	
2	Kinetics of Severity Biomarkers and Immunological Features of Methylprednisolone Therapy for Severe COVID-19 Patients. <i>Frontiers in Immunology</i> , 2022 , 13, 758946	8.4	
1	Distribution of nontuberculous mycobacteria in patients with and without HIV/AIDS in Chongqing. <i>HIV Medicine</i> , 2022 , 23 Suppl 1, 54-63	2.7	