

Shoya Iwanami

List of Publications by Year in Descending Order

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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.

18

papers

174

citations

8

h-index

13

g-index

27

ext. papers

385

ext. citations

5.4

avg, IF

2.69

L-index

#	Paper	IF	Citations
18	Antithetic effect of interferon- β n cell-free and cell-to-cell HIV-1 infection.. <i>PLoS Computational Biology</i> , 2022 , 18, e1010053	5	0
17	Detection of significant antiviral drug effects on COVID-19 using viral load and PCR-positive rate in randomized controlled trials. <i>Translational and Regulatory Sciences</i> , 2021 , 3, 85-88	0.3	
16	A quantitative model used to compare within-host SARS-CoV-2, MERS-CoV, and SARS-CoV dynamics provides insights into the pathogenesis and treatment of SARS-CoV-2. <i>PLoS Biology</i> , 2021 , 19, e3001128	9.7	25
15	Potential anti-COVID-19 agents, cepharanthine and nelfinavir, and their usage for combination treatment. <i>IScience</i> , 2021 , 24, 102367	6.1	34
14	Mefloquine, a Potent Anti-severe Acute Respiratory Syndrome-Related Coronavirus 2 (SARS-CoV-2) Drug as an Entry Inhibitor. <i>Frontiers in Microbiology</i> , 2021 , 12, 651403	5.7	6
13	Time variation in the probability of failing to detect a case of polymerase chain reaction testing for SARS-CoV-2 as estimated from a viral dynamics model. <i>Journal of the Royal Society Interface</i> , 2021 , 18, 20200947	4.1	1
12	Estimation of the incubation period of COVID-19 using viral load data. <i>Epidemics</i> , 2021 , 35, 100454	5.1	12
11	Detection of significant antiviral drug effects on COVID-19 with reasonable sample sizes in randomized controlled trials: A modeling study. <i>PLoS Medicine</i> , 2021 , 18, e1003660	11.6	8
10	Revisiting the guidelines for ending isolation for COVID-19 patients. <i>ELife</i> , 2021 , 10,	8.9	4
9	Incomplete antiviral treatment may induce longer durations of viral shedding during SARS-CoV-2 infection. <i>Life Science Alliance</i> , 2021 , 4,	5.8	2
8	Should a viral genome stay in the host cell or leave? A quantitative dynamics study of how hepatitis C virus deals with this dilemma. <i>PLoS Biology</i> , 2020 , 18, e3000562	9.7	2
7	Quantitative Immunology by Data Analysis Using Mathematical Models 2019 , 984-992		1
6	Revealing uninfected and infected target cell dynamics from peripheral blood data in highly and less pathogenic simian/human immunodeficiency virus infected Rhesus macaque. <i>Journal of Theoretical Biology</i> , 2019 , 479, 29-36	2.3	1
5	A highly pathogenic simian/human immunodeficiency virus effectively produces infectious virions compared with a less pathogenic virus in cell culture. <i>Theoretical Biology and Medical Modelling</i> , 2017 , 14, 9	2.3	11
4	Modelling SARS-CoV-2 Dynamics: Implications for Therapy		15
3	Inferring Timing of Infection Using Within-host SARS-CoV-2 Infection Dynamics Model: Are Imported Cases Truly Imported?		8
2	Multidrug treatment with nelfinavir and cepharanthine against COVID-19		35

1 Rethinking antiviral effects for COVID-19 in clinical studies: early initiation is key to successful treatment 5