

Igor Siveroni

List of Publications by Year in descending order

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Version: 2024-02-01

12
papers

6,676
citations

759055

12
h-index

1199470

12
g-index

17
all docs

17
docs citations

17
times ranked

13297
citing authors

#	ARTICLE	IF	CITATIONS
1	Additive Uncorrelated Relaxed Clock Models for the Dating of Genomic Epidemiology Phylogenies. <i>Molecular Biology and Evolution</i> , 2021, 38, 307-317.	3.5	28
2	Evaluating the Effects of SARS-CoV-2 Spike Mutation D614G on Transmissibility and Pathogenicity. <i>Cell</i> , 2021, 184, 64-75.e11.	13.5	843
3	Genomic epidemiology of a densely sampled COVID-19 outbreak in China. <i>Virus Evolution</i> , 2021, 7, veaa102.	2.2	18
4	Genetic evidence for the association between COVID-19 epidemic severity and timing of non-pharmaceutical interventions. <i>Nature Communications</i> , 2021, 12, 2188.	5.8	23
5	Changes in symptomatology, reinfection, and transmissibility associated with the SARS-CoV-2 variant B.1.1.7: an ecological study. <i>Lancet Public Health</i> , The, 2021, 6, e335-e345.	4.7	269
6	State-level tracking of COVID-19 in the United States. <i>Nature Communications</i> , 2020, 11, 6189.	5.8	104
7	Estimating the effects of non-pharmaceutical interventions on COVID-19 in Europe. <i>Nature</i> , 2020, 584, 257-261.	13.7	2,558
8	Evidence of initial success for China exiting COVID-19 social distancing policy after achieving containment. <i>Wellcome Open Research</i> , 2020, 5, 81.	0.9	62
9	Evidence of initial success for China exiting COVID-19 social distancing policy after achieving containment. <i>Wellcome Open Research</i> , 2020, 5, 81.	0.9	81
10	BEAST 2.5: An advanced software platform for Bayesian evolutionary analysis. <i>PLoS Computational Biology</i> , 2019, 15, e1006650.	1.5	2,484
11	Bayesian phylodynamic inference with complex models. <i>PLoS Computational Biology</i> , 2018, 14, e1006546.	1.5	64
12	A UML-based static verification framework for security. <i>Requirements Engineering</i> , 2010, 15, 95-118.	2.1	21