

Tara Alpert

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.

24
papers

1,060
citations

11
h-index

26
g-index

26
ext. papers

1,601
ext. citations

17.4
avg, IF

4.25
L-index

#	Paper	IF	Citations
24	Coast-to-Coast Spread of SARS-CoV-2 during the Early Epidemic in the United States. <i>Cell</i> , 2020 , 181, 990-996.e5	56.2	235
23	SARS-CoV-2 infection of the placenta. <i>Journal of Clinical Investigation</i> , 2020 , 130, 4947-4953	15.9	230
22	Splicing and transcription touch base: co-transcriptional spliceosome assembly and function. <i>Nature Reviews Molecular Cell Biology</i> , 2017 , 18, 637-650	48.7	173
21	Multiplex qPCR discriminates variants of concern to enhance global surveillance of SARS-CoV-2. <i>PLoS Biology</i> , 2021 , 19, e3001236	9.7	85
20	Acute encephalopathy with elevated CSF inflammatory markers as the initial presentation of COVID-19. <i>BMC Neurology</i> , 2020 , 20, 248	3.1	78
19	Impact of circulating SARS-CoV-2 variants on mRNA vaccine-induced immunity. <i>Nature</i> , 2021 ,	50.4	71
18	Early introductions and transmission of SARS-CoV-2 variant B.1.1.7 in the United States. <i>Cell</i> , 2021 , 184, 2595-2604.e13	56.2	55
17	Perfect timing: splicing and transcription rates in living cells. <i>Wiley Interdisciplinary Reviews RNA</i> , 2017 , 8, e1401	9.3	43
16	PCR assay to enhance global surveillance for SARS-CoV-2 variants of concern 2021 ,		15
15	Real-time public health communication of local SARS-CoV-2 genomic epidemiology. <i>PLoS Biology</i> , 2020 , 18, e3000869	9.7	12
14	A stem-loop RNA RIG-I agonist protects against acute and chronic SARS-CoV-2 infection in mice. <i>Journal of Experimental Medicine</i> , 2022 , 219,	16.6	11
13	Impact of circulating SARS-CoV-2 variants on mRNA vaccine-induced immunity in uninfected and previously infected individuals		11
12	Early introductions and community transmission of SARS-CoV-2 variant B.1.1.7 in the United States 2021 ,		9
11	Comparative transmissibility of SARS-CoV-2 variants Delta and Alpha in New England, USA.. <i>Cell Reports Medicine</i> , 2022 , 3, 100583	18	9
10	Widespread Transcriptional Readthrough Caused by Nab2 Depletion Leads to Chimeric Transcripts with Retained Introns. <i>Cell Reports</i> , 2020 , 33, 108324	10.6	6
9	Longitudinal immune profiling of a SARS-CoV-2 reinfection in a solid organ transplant recipient. <i>Journal of Infectious Diseases</i> , 2021 ,	7	4
8	Combining genomic and epidemiological data to compare the transmissibility of SARS-CoV-2 lineages 2021 ,		4

7	COVID-19 Outcomes and Genomic Characterization of SARS-CoV-2 Isolated From Veterans in New England States: Retrospective Analysis.. <i>Jmirx Med</i> , 2021 , 2, e31503	0.2	3
6	Variant abundance estimation for SARS-CoV-2 in wastewater using RNA-Seq quantification 2021 ,		3
5	An outbreak of SARS-CoV-2 on a transplant unit in the early vaccination era.. <i>Transplant Infectious Disease</i> , 2021 ,	2.7	1
4	Longitudinal immune profiling of a SARS-CoV-2 reinfection in a solid organ transplant recipient 2021 ,		1
3	Combining genomic and epidemiological data to compare the transmissibility of SARS-CoV-2 variants Alpha and Iota.. <i>Communications Biology</i> , 2022 , 5, 439	6.7	1
2	Sequencing SARS-CoV-2 genomes from saliva.. <i>Virus Evolution</i> , 2022 , 8, veab098	3.7	0
1	Authors' Response to Peer Reviews of "COVID-19 Outcomes and Genomic Characterization of SARS-CoV-2 Isolated From Veterans in New England States: Retrospective Analysis" <i>Jmirx Med</i> , 2021 , 2, e35515	0.2	