## Tara Alpert

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8895756/publications.pdf

Version: 2024-02-01

840776 888059 1,969 19 11 17 h-index citations g-index papers 26 26 26 5049 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Longitudinal Immune Profiling of a Severe Acute Respiratory Syndrome Coronavirus 2 Reinfection in a Solid Organ Transplant Recipient. Journal of Infectious Diseases, 2022, 225, 374-384.	4.0	7
2	A stem-loop RNA RIG-I agonist protects against acute and chronic SARS-CoV-2 infection in mice. Journal of Experimental Medicine, 2022, 219, .	8.5	46
3	Sequencing SARS-CoV-2 genomes from saliva. Virus Evolution, 2022, 8, veab098.	4.9	4
4	An outbreak of SARSâ€CoVâ€2 on a transplant unit in the early vaccination era. Transplant Infectious Disease, 2022, 24, .	1.7	5
5	Comparative transmissibility of SARS-CoV-2 variants Delta and Alpha in New England, USA. Cell Reports Medicine, 2022, 3, 100583.	6.5	101
6	Combining genomic and epidemiological data to compare the transmissibility of SARS-CoV-2 variants Alpha and lota. Communications Biology, 2022, 5, 439.	4.4	9
7	Abstract S03-03: Cancer patients display diminished viral RNA clearance and altered T cell responses during SARS-CoV-2 infection. , 2021, , .		0
8	Early introductions and transmission of SARS-CoV-2 variant B.1.1.7 in the United States. Cell, 2021, 184, 2595-2604.e13.	28.9	113
9	Multiplex qPCR discriminates variants of concern to enhance global surveillance of SARS-CoV-2. PLoS Biology, 2021, 19, e3001236.	5.6	200
10	Impact of circulating SARS-CoV-2 variants on mRNA vaccine-induced immunity. Nature, 2021, 600, 523-529.	27.8	194
11	COVID-19 Outcomes and Genomic Characterization of SARS-CoV-2 Isolated From Veterans in New England States: Retrospective Analysis. Jmirx Med, 2021, 2, e31503.	0.4	5
12	Authors' Response to Peer Reviews of "COVID-19 Outcomes and Genomic Characterization of SARS-CoV-2 Isolated From Veterans in New England States: Retrospective Analysisâ€, Jmirx Med, 2021, 2, e35515.	0.4	0
13	Widespread Transcriptional Readthrough Caused by Nab2 Depletion Leads to Chimeric Transcripts with Retained Introns. Cell Reports, 2020, 33, 108324.	6.4	26
14	Real-time public health communication of local SARS-CoV-2 genomic epidemiology. PLoS Biology, 2020, 18, e3000869.	5.6	15
15			
10	Coast-to-Coast Spread of SARS-CoV-2 during the Early Epidemic in the United States. Cell, 2020, 181, 990-996.e5.	28.9	321
16		28.9	108
	990-996.e5.  Acute encephalopathy with elevated CSF inflammatory markers as the initial presentation of COVID-19.		

#	Article	IF	CITATIONS
19	Perfect timing: splicing and transcription rates in living cells. Wiley Interdisciplinary Reviews RNA, 2017, 8, e1401.	6.4	70