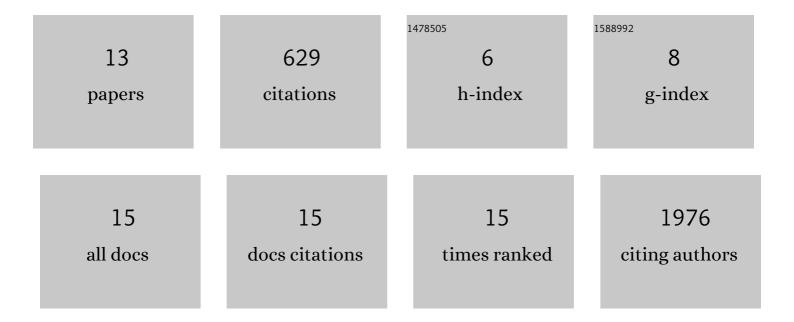
## Talia Kustin

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/324094/publications.pdf Version: 2024-02-01



TALIA KUSTIN

#	Article	IF	CITATIONS
1	Evidence for increased breakthrough rates of SARS-CoV-2 variants of concern in BNT162b2-mRNA-vaccinated individuals. Nature Medicine, 2021, 27, 1379-1384.	30.7	296
2	Full genome viral sequences inform patterns of SARS-CoV-2 spread into and within Israel. Nature Communications, 2020, 11, 5518.	12.8	115
3	Biased Mutation and Selection in RNA Viruses. Molecular Biology and Evolution, 2021, 38, 575-588.	8.9	43
4	A method to identify respiratory virus infections in clinical samples using next-generation sequencing. Scientific Reports, 2019, 9, 2606.	3.3	32
5	In vivo engineered B cells secrete high titers of broadly neutralizing anti-HIV antibodies in mice. Nature Biotechnology, 2022, 40, 1241-1249.	17.5	29
6	Drivers of within-host genetic diversity in acute infections of viruses. PLoS Pathogens, 2020, 16, e1009029.	4.7	25
7	Inferring Protein Function in an Emerging Virus: Detection of the Nucleoprotein in Tilapia Lake Virus. Journal of Virology, 2022, 96, JVI0175721.	3.4	11
8	The association between SARSâ€CoVâ€⊋ infection and late pregnancy loss. International Journal of Gynecology and Obstetrics, 2021, , .	2.3	4
9	Drivers of within-host genetic diversity in acute infections of viruses. , 2020, 16, e1009029.		0
10	Drivers of within-host genetic diversity in acute infections of viruses. , 2020, 16, e1009029.		0
11	Drivers of within-host genetic diversity in acute infections of viruses. , 2020, 16, e1009029.		0
12	Drivers of within-host genetic diversity in acute infections of viruses. , 2020, 16, e1009029.		0
13	Drivers of within-host genetic diversity in acute infections of viruses. , 2020, 16, e1009029.		0