

Scott Hennigan

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1868138/publications.pdf>

Version: 2024-02-01

12
papers

545
citations

1478505

6
h-index

1872680

6
g-index

12
all docs

12
docs citations

12
times ranked

1432
citing authors

#	ARTICLE	IF	CITATIONS
1	Zika virus evolution and spread in the Americas. <i>Nature</i> , 2017, 546, 411-415.	27.8	323
2	Capturing sequence diversity in metagenomes with comprehensive and scalable probe design. <i>Nature Biotechnology</i> , 2019, 37, 160-168.	17.5	96
3	Transmission from vaccinated individuals in a large SARS-CoV-2 Delta variant outbreak. <i>Cell</i> , 2022, 185, 485-492.e10.	28.9	38
4	Combining genomics and epidemiology to track mumps virus transmission in the United States. <i>PLoS Biology</i> , 2020, 18, e3000611.	5.6	37
5	Large-Scale Complete-Genome Sequencing and Phylodynamic Analysis of Eastern Equine Encephalitis Virus Reveals Source-Sink Transmission Dynamics in the United States. <i>Journal of Virology</i> , 2018, 92, .	3.4	31
6	Jamestown Canyon virus in Massachusetts: clinical case series and vector screening. <i>Emerging Microbes and Infections</i> , 2020, 9, 903-912.	6.5	20
7	Combining genomics and epidemiology to track mumps virus transmission in the United States. , 2020, 18, e3000611.		0
8	Combining genomics and epidemiology to track mumps virus transmission in the United States. , 2020, 18, e3000611.		0
9	Combining genomics and epidemiology to track mumps virus transmission in the United States. , 2020, 18, e3000611.		0
10	Combining genomics and epidemiology to track mumps virus transmission in the United States. , 2020, 18, e3000611.		0
11	Combining genomics and epidemiology to track mumps virus transmission in the United States. , 2020, 18, e3000611.		0
12	Combining genomics and epidemiology to track mumps virus transmission in the United States. , 2020, 18, e3000611.		0