

Jeffrey R Kugelman

List of Publications by Year in descending order

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Version: 2024-02-01

16
papers

2,519
citations

687363

13
h-index

940533

16
g-index

17
all docs

17
docs citations

17
times ranked

5335
citing authors

#	ARTICLE	IF	CITATIONS
1	Therapeutic efficacy of the small molecule GS-5734 against Ebola virus in rhesus monkeys. <i>Nature</i> , 2016, 531, 381-385.	27.8	1,245
2	Virus genomes reveal factors that spread and sustained the Ebola epidemic. <i>Nature</i> , 2017, 544, 309-315.	27.8	346
3	Ebola Virus Epidemiology, Transmission, and Evolution during Seven Months in Sierra Leone. <i>Cell</i> , 2015, 161, 1516-1526.	28.9	275
4	Standards for Sequencing Viral Genomes in the Era of High-Throughput Sequencing. <i>MBio</i> , 2014, 5, e01360-14.	4.1	89
5	Evolution and Spread of Ebola Virus in Liberia, 2014–2015. <i>Cell Host and Microbe</i> , 2015, 18, 659-669.	11.0	87
6	Nomenclature- and Database-Compatible Names for the Two Ebola Virus Variants that Emerged in Guinea and the Democratic Republic of the Congo in 2014. <i>Viruses</i> , 2014, 6, 4760-4799.	3.3	83
7	Monitoring of Ebola Virus Makona Evolution through Establishment of Advanced Genomic Capability in Liberia. <i>Emerging Infectious Diseases</i> , 2015, 21, 1135-1143.	4.3	79
8	Evaluation of the Potential Impact of Ebola Virus Genomic Drift on the Efficacy of Sequence-Based Candidate Therapeutics. <i>MBio</i> , 2015, 6, .	4.1	62
9	Reduced evolutionary rate in reemerged Ebola virus transmission chains. <i>Science Advances</i> , 2016, 2, e1600378.	10.3	62
10	Development of a coronavirus disease 2019 nonhuman primate model using airborne exposure. <i>PLoS ONE</i> , 2021, 16, e0246366.	2.5	52
11	Recent successes in therapeutics for Ebola virus disease: no time for complacency. <i>Lancet Infectious Diseases</i> , The, 2020, 20, e231-e237.	9.1	42
12	Error baseline rates of five sample preparation methods used to characterize RNA virus populations. <i>PLoS ONE</i> , 2017, 12, e0171333.	2.5	21
13	Genomic features of humoral immunity support tolerance model in Egyptian rousette bats. <i>Cell Reports</i> , 2021, 35, 109140.	6.4	19
14	Complete Coding Sequences of Eastern Equine Encephalitis Virus and Venezuelan Equine Encephalitis Virus Strains Isolated from Human Cases. <i>Genome Announcements</i> , 2015, 3, .	0.8	15
15	Molecular analysis of the 2012 Bundibugyo virus disease outbreak. <i>Cell Reports Medicine</i> , 2021, 2, 100351.	6.5	4
16	A Department of Defense Laboratory Consortium Approach to Next Generation Sequencing and Bioinformatics Training for Infectious Disease Surveillance in Kenya. <i>Frontiers in Genetics</i> , 2020, 11, 577563.	2.3	1