

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