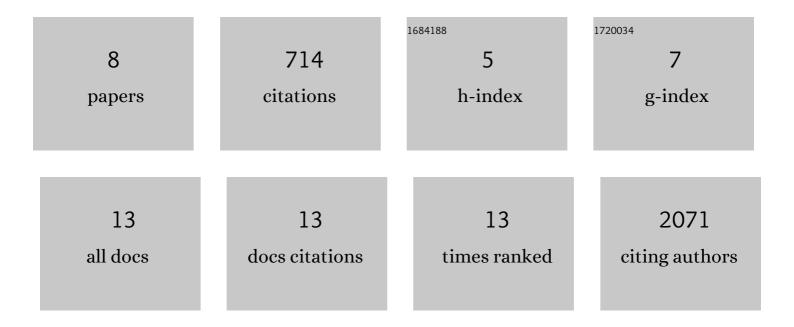
## **Clinton Paden**

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1773829/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Genomic surveillance reveals multiple introductions of SARS-CoV-2 into Northern California. Science, 2020, 369, 582-587.	12.6	253
2	Rapid, Sensitive, Full-Genome Sequencing of Severe Acute Respiratory Syndrome Coronavirus 2. Emerging Infectious Diseases, 2020, 26, 2401-2405.	4.3	126
3	Comprehensive viral enrichment enables sensitive respiratory virus genomic identification and analysis by next generation sequencing. Genome Research, 2018, 28, 869-877.	5.5	74
4	Zoonotic origin and transmission of Middle East respiratory syndrome coronavirus in the <scp>UAE</scp> . Zoonoses and Public Health, 2018, 65, 322-333.	2.2	56
5	Diversity of Middle East respiratory syndrome coronaviruses in 109 dromedary camels based on full-genome sequencing, Abu Dhabi, United Arab Emirates. Emerging Microbes and Infections, 2017, 6, 1-10.	6.5	34
6	Isolation and growth characterization of novel full length and deletion mutant human MERS-CoV strains from clinical specimens collected during 2015. Journal of General Virology, 2019, 100, 1523-1529.	2.9	5
7	Introduction, Transmission Dynamics, and Fate of Early Severe Acute Respiratory Syndrome Coronavirus 2 Lineages in Santa Clara County, California. Journal of Infectious Diseases, 2021, 224, 207-217.	4.0	2
8	Unrecognized introductions of SARS-CoV-2 into the US state of Georgia shaped the early epidemic. Virus Evolution, 2022, 8, veac011.	4.9	2