

# Joseph G Chappell

## List of Publications by Year in Descending Order

**Source:** <https://exaly.com/author-pdf/6322029/joseph-g-chappell-publications-by-year.pdf>

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This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

10  
papers

65  
citations

4  
h-index

7  
g-index

13  
ext. papers

115  
ext. citations

6.6  
avg, IF

1.81  
L-index

#	Paper	IF	Citations
10	Retrospective screening of routine respiratory samples revealed undetected community transmission and missed intervention opportunities for SARS-CoV-2 in the United Kingdom. <i>Journal of General Virology</i> , <b>2021</b> , 102,	4.9	6
9	Two doses of the SARS-CoV-2 BNT162b2 vaccine enhance antibody responses to variants in individuals with prior SARS-CoV-2 infection. <i>Science Translational Medicine</i> , <b>2021</b> , 13, eabj0847	17.5	16
8	Retrieval of the Complete Coding Sequence of the UK-Endemic Tatenale Orthohantavirus Reveals Extensive Strain Variation and Supports Its Classification as a Novel Species. <i>Viruses</i> , <b>2020</b> , 12,	6.2	3
7	Perceptions and Experiences of the University of Nottingham Pilot SARS-CoV-2 Asymptomatic Testing Service: A Mixed-Methods Study. <i>International Journal of Environmental Research and Public Health</i> , <b>2020</b> , 18,	4.6	15
6	Discovery and Prevalence of Divergent RNA Viruses in European Field Voles and Rabbits. <i>Viruses</i> , <b>2019</b> , 12,	6.2	2
5	Shared Common Ancestry of Rodent Alphacoronaviruses Sampled Globally. <i>Viruses</i> , <b>2019</b> , 11,	6.2	16
4	Identification of Infectious Agents in High-Throughput Sequencing Data Sets Is Easily Achievable Using Free, Cloud-Based Bioinformatics Platforms. <i>Journal of Clinical Microbiology</i> , <b>2019</b> , 57,	9.7	3
3	Retrospective screening of routine respiratory samples revealed undetected community transmission and missed intervention opportunities for SARS-CoV-2 in the United Kingdom		2
2	Potent anti-SARS-CoV-2 Antibody Responses are Associated with Better Prognosis in Hospital Inpatient COVID-19 Disease		1
1	Retrieval of the Complete Coding Sequence of the UK-Endemic Tatenale Orthohantavirus Reveals Extensive Strain Variation and Supports its Classification as a Novel Species		1