

# Jantien A Backer

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9177625/publications.pdf>

Version: 2024-02-01

10  
papers

1,704  
citations

1163117

8  
h-index

1372567

10  
g-index

17  
all docs

17  
docs citations

17  
times ranked

3403  
citing authors

#	ARTICLE	IF	CITATIONS
1	Transmissibility of SARS-CoV-2 among fully vaccinated individuals. <i>Lancet Infectious Diseases</i> , The, 2022, 22, 16-17.	9.1	5
2	Shorter serial intervals in SARS-CoV-2 cases with Omicron BA.1 variant compared with Delta variant, the Netherlands, 13 to 26 December 2021. <i>Eurosurveillance</i> , 2022, 27, .	7.0	99
3	Impact of physical distancing measures against COVID-19 on contacts and mixing patterns: repeated cross-sectional surveys, the Netherlands, 2016–17, April 2020 and June 2020. <i>Eurosurveillance</i> , 2021, 26, .	7.0	59
4	Associations Between Measures of Social Distancing and Severe Acute Respiratory Syndrome Coronavirus 2 Seropositivity: A Nationwide Population-based Study in the Netherlands. <i>Clinical Infectious Diseases</i> , 2021, 73, 2318-2321.	5.8	40
5	SOCRATES-CoMix: a platform for timely and open-source contact mixing data during and in between COVID-19 surges and interventions in over 20 European countries. <i>BMC Medicine</i> , 2021, 19, 254.	5.5	45
6	Vaccinating children against influenza: overall cost-effective with potential for undesirable outcomes. <i>BMC Medicine</i> , 2020, 18, 11.	5.5	10
7	Incubation period of 2019 novel coronavirus (2019-nCoV) infections among travellers from Wuhan, China, 20–28 January 2020. <i>Eurosurveillance</i> , 2020, 25, .	7.0	1,234
8	Visual tools to assess the plausibility of algorithm-identified infectious disease clusters: an application to mumps data from the Netherlands dating from January 2009 to June 2016. <i>Eurosurveillance</i> , 2019, 24, .	7.0	2
9	Simultaneous inference of phylogenetic and transmission trees in infectious disease outbreaks. <i>PLoS Computational Biology</i> , 2017, 13, e1005495.	3.2	93
10	Spatiotemporal Analysis of the 2014 Ebola Epidemic in West Africa. <i>PLoS Computational Biology</i> , 2016, 12, e1005210.	3.2	36