

# Cara E Brook

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9228821/publications.pdf>

Version: 2024-02-01

25  
papers

1,350  
citations

623188

14  
h-index

580395

25  
g-index

35  
all docs

35  
docs citations

35  
times ranked

2479  
citing authors

#	ARTICLE	IF	CITATIONS
1	Bats as “special” reservoirs for emerging zoonotic pathogens. <i>Trends in Microbiology</i> , 2015, 23, 172-180.	3.5	358
2	A year of genomic surveillance reveals how the SARS-CoV-2 pandemic unfolded in Africa. <i>Science</i> , 2021, 374, 423-431.	6.0	144
3	Possibility for reverse zoonotic transmission of SARS-CoV-2 to free-ranging wildlife: A case study of bats. <i>PLoS Pathogens</i> , 2020, 16, e1008758.	2.1	127
4	Ecology, evolution and spillover of coronaviruses from bats. <i>Nature Reviews Microbiology</i> , 2022, 20, 299-314.	13.6	108
5	Accelerated viral dynamics in bat cell lines, with implications for zoonotic emergence. <i>ELife</i> , 2020, 9, .	2.8	91
6	<i>Bartonella</i> spp. in Fruit Bats and Blood-Feeding Ectoparasites in Madagascar. <i>PLoS Neglected Tropical Diseases</i> , 2015, 9, e0003532.	1.3	71
7	Host phylogenetic distance drives trends in virus virulence and transmissibility across the animal–human interface. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2019, 374, 20190296.	1.8	64
8	The science of the host–virus network. <i>Nature Microbiology</i> , 2021, 6, 1483-1492.	5.9	59
9	Blueprint for a pop-up SARS-CoV-2 testing lab. <i>Nature Biotechnology</i> , 2020, 38, 791-797.	9.4	50
10	Disentangling serology to elucidate henipavirus and filovirus transmission in Madagascar fruit bats. <i>Journal of Animal Ecology</i> , 2019, 88, 1001-1016.	1.3	36
11	Spatial heterogeneity in projected leprosy trends in India. <i>Parasites and Vectors</i> , 2015, 8, 542.	1.0	23
12	Population viability and harvest sustainability for Madagascar lemurs. <i>Conservation Biology</i> , 2019, 33, 99-111.	2.4	23
13	Bats host the most virulent “but not the most dangerous” zoonotic viruses. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, e2113628119.	3.3	22
14	Optimizing COVID-19 control with asymptomatic surveillance testing in a university environment. <i>Epidemics</i> , 2021, 37, 100527.	1.5	21
15	Elucidating transmission dynamics and host-parasite-vector relationships for rodent-borne <i>Bartonella</i> spp. in Madagascar. <i>Epidemics</i> , 2017, 20, 56-66.	1.5	19
16	Population trends for two Malagasy fruit bats. <i>Biological Conservation</i> , 2019, 234, 165-171.	1.9	15
17	Modeling the burden of poultry disease on the rural poor in Madagascar. <i>One Health</i> , 2015, 1, 60-65.	1.5	14
18	Introduction of rubella-containing-vaccine to Madagascar: implications for roll-out and local elimination. <i>Journal of the Royal Society Interface</i> , 2016, 13, 20151101.	1.5	14

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19	Babesial infection in the Madagascan flying fox, <i>Pteropus rufus</i> Geoffroy, 1803. <i>Parasites and Vectors</i> , 2019, 12, 51.	1.0	14
20	A review of mechanistic models of viral dynamics in bat reservoirs for zoonotic disease. <i>Pathogens and Global Health</i> , 2020, 114, 407-425.	1.0	13
21	Full Genome Nobecovirus Sequences From Malagasy Fruit Bats Define a Unique Evolutionary History for This Coronavirus Clade. <i>Frontiers in Public Health</i> , 2022, 10, 786060.	1.3	13
22	Do gastrointestinal microbiomes play a role in bats' unique viral hosting capacity?. <i>Trends in Microbiology</i> , 2022, 30, 632-642.	3.5	9
23	The zoonotic potential of bat-borne coronaviruses. <i>Emerging Topics in Life Sciences</i> , 2020, 4, 365-381.	1.1	8
24	Cross-sectional cycle threshold values reflect epidemic dynamics of COVID-19 in Madagascar. <i>Epidemics</i> , 2022, 38, 100533.	1.5	8
25	A batty concept goes viral. <i>Nature Ecology and Evolution</i> , 2019, 3, 1620-1621.	3.4	0