

Catherine B Pratt

List of Publications by Year in Descending Order

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

19 papers	548 citations	12 h-index	21 g-index
21 ext. papers	799 ext. citations	24.2 avg, IF	2.49 L-index

#	Paper	IF	Citations
19	A year of genomic surveillance reveals how the SARS-CoV-2 pandemic unfolded in Africa. <i>Science</i> , 2021 , 374, 423-431	33.3	35
18	Genome Analysis for Sequence Variants in SARS-CoV-2 Among Asymptomatic Individuals in a Long-term Care Facility. <i>JAMA Network Open</i> , 2021 , 4, e217939	10.4	2
17	Ebola Virus Transmission Initiated by Relapse of Systemic Ebola Virus Disease. <i>New England Journal of Medicine</i> , 2021 , 384, 1240-1247	59.2	16
16	Integration of genomic sequencing into the response to the Ebola virus outbreak in Nord Kivu, Democratic Republic of the Congo. <i>Nature Medicine</i> , 2021 , 27, 710-716	50.5	3
15	Assessment of a Program for SARS-CoV-2 Screening and Environmental Monitoring in an Urban Public School District. <i>JAMA Network Open</i> , 2021 , 4, e2126447	10.4	5
14	"Super-Spreaders" and Person-to-Person Transmission of Andes Virus in Argentina. <i>New England Journal of Medicine</i> , 2020 , 383, 2230-2241	59.2	17
13	Prospective Cohort Study of Next-Generation Sequencing as a Diagnostic Modality for Unexplained Encephalitis in Children. <i>Journal of the Pediatric Infectious Diseases Society</i> , 2020 , 9, 326-333	4.8	17
12	Medical countermeasures during the 2018 Ebola virus disease outbreak in the North Kivu and Ituri Provinces of the Democratic Republic of the Congo: a rapid genomic assessment. <i>Lancet Infectious Diseases, The</i> , 2019 , 19, 648-657	25.5	36
11	2018 Ebola virus disease outbreak in Équateur Province, Democratic Republic of the Congo: a retrospective genomic characterisation. <i>Lancet Infectious Diseases, The</i> , 2019 , 19, 641-647	25.5	17
10	Lassa virus circulating in Liberia: a retrospective genomic characterisation. <i>Lancet Infectious Diseases, The</i> , 2019 , 19, 1371-1378	25.5	21
9	Genomic characterisation of human monkeypox virus in Nigeria. <i>Lancet Infectious Diseases, The</i> , 2018 , 18, 246	25.5	17
8	Whole-Genome Sequences of Variants of Sterne and Their Toxin Gene Deletion Mutants. <i>Genome Announcements</i> , 2017 , 5,		3
7	Temporal and spatial analysis of the 2014-2015 Ebola virus outbreak in West Africa. <i>Nature</i> , 2015 , 524, 97-101	50.4	229
6	Proteomics-identified Bvg-activated autotransporters protect against bordetella pertussis in a mouse model. <i>PLoS ONE</i> , 2014 , 9, e105011	3.7	32
5	Plasticity of fimbrial genotype and serotype within populations of Bordetella pertussis: analysis by paired flow cytometry and genome sequencing. <i>Microbiology (United Kingdom)</i> , 2014 , 160, 2030-2044	2.9	5
4	Antibody responses to Bordetella pertussis Fim2 or Fim3 following immunization with a whole-cell, two-component, or five-component acellular pertussis vaccine and following pertussis disease in children in Sweden in 1997 and 2007. <i>Vaccine Journal</i> , 2014 , 21, 165-73		12
3	Nasopharyngeal colonization by Neisseria lactamica and induction of protective immunity against Neisseria meningitidis. <i>Clinical Infectious Diseases</i> , 2011 , 52, 70-7	11.6	57

2	Characterization of the key antigenic components and pre-clinical immune responses to a meningococcal disease vaccine based on Neisseria lactamica outer membrane vesicles. <i>Hum Vaccin</i> , 2008 , 4, 23-30	22
1	Operationalizing genomic epidemiology during the Nord-Kivu Ebola outbreak, Democratic Republic of the Congo	2