

# Barry Atkinson

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3041037/publications.pdf>

Version: 2024-02-01

37  
papers

2,350  
citations

430874

18  
h-index

377865

34  
g-index

42  
all docs

42  
docs citations

42  
times ranked

4706  
citing authors

#	ARTICLE	IF	CITATIONS
1	Investigation of a SARS-CoV-2 Outbreak at an Automotive Manufacturing Site in England. <i>International Journal of Environmental Research and Public Health</i> , 2022, 19, 6400.	2.6	10
2	Infection of incompetent monkeypox virus contamination identified in domestic settings following an imported case of monkeypox into the <sc>UK</sc>. <i>Environmental Microbiology</i> , 2022, 24, 4561-4569.	3.8	45
3	Longitudinal antibody and T cell responses in Ebola virus disease survivors and contacts: an observational cohort study. <i>Lancet Infectious Diseases</i> , The, 2021, 21, 507-516.	9.1	26
4	Oropouche virus cases identified in Ecuador using an optimised qRT-PCR informed by metagenomic sequencing. <i>PLoS Neglected Tropical Diseases</i> , 2020, 14, e0007897.	3.0	10
5	SARS-CoV-2 shedding and infectivity. <i>Lancet</i> , The, 2020, 395, 1339-1340.	13.7	125
6	Title is missing!. , 2020, 14, e0007897.		0
7	Title is missing!. , 2020, 14, e0007897.		0
8	Title is missing!. , 2020, 14, e0007897.		0
9	Detection of Crimean-Congo Haemorrhagic Fever cases in a severe undifferentiated febrile illness outbreak in the Federal Republic of Sudan: A retrospective epidemiological and diagnostic cohort study. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007571.	3.0	10
10	Clinical and molecular epidemiology of Crimean-Congo hemorrhagic fever in Oman. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007100.	3.0	25
11	Emerging arboviruses of clinical importance in Central Asia. <i>Journal of General Virology</i> , 2018, 99, 1172-1184.	2.9	15
12	Lineage-dependent differences in the disease progression of Zika virus infection in type-I interferon receptor knockout (A129) mice. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0005704.	3.0	56
13	Presence and Persistence of Zika Virus RNA in Semen, United Kingdom, 2016. <i>Emerging Infectious Diseases</i> , 2017, 23, 611-615.	4.3	95
14	A Susceptible Mouse Model for Zika Virus Infection. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004658.	3.0	262
15	Serological and Virological Evidence of Crimean-Congo Haemorrhagic Fever Virus Circulation in the Human Population of Borno State, Northeastern Nigeria. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0005126.	3.0	28
16	Detection of Zika Virus in Semen. <i>Emerging Infectious Diseases</i> , 2016, 22, 940-940.	4.3	286
17	Complete Genome Sequence of Seoul Virus Strain Tchoupitoulas. <i>Genome Announcements</i> , 2016, 4, .	0.8	3
18	<i>Hyalomma</i> ticks on northward migrating birds in southern Spain: Implications for the risk of entry of Crimean-Congo haemorrhagic fever virus to Great Britain. <i>Journal of Vector Ecology</i> , 2016, 41, 128-134.	1.0	25

#	ARTICLE	IF	CITATIONS
19	Analysis of Diagnostic Findings From the European Mobile Laboratory in GuÃ©ckÃ©dou, Guinea, March 2014 Through March 2015. <i>Journal of Infectious Diseases</i> , 2016, 214, S250-S257.	4.0	32
20	Specificity, cross-reactivity, and function of antibodies elicited by Zika virus infection. <i>Science</i> , 2016, 353, 823-826.	12.6	675
21	Complete Genome Sequence of Zika Virus Isolated from Semen. <i>Genome Announcements</i> , 2016, 4, .	0.8	16
22	Importation of a <i>Hyalomma lusitanicum</i> tick into the UK on a dog. <i>Veterinary Record</i> , 2016, 179, 415-415.	0.3	14
23	Complete Genomic Sequence of Issyk-Kul Virus. <i>Genome Announcements</i> , 2015, 3, .	0.8	13
24	Crimean-Congo haemorrhagic fever virus in Kazakhstan (1948-2013). <i>International Journal of Infectious Diseases</i> , 2015, 38, 19-23.	3.3	30
25	A non-fatal case of hantavirus cardiopulmonary syndrome imported into the UK (ex Panama), July 2014. <i>Journal of Clinical Virology</i> , 2015, 67, 52-55.	3.1	5
26	Temporal and spatial analysis of the 2014â€“2015 Ebola virus outbreak in West Africa. <i>Nature</i> , 2015, 524, 97-101.	27.8	272
27	Whole-genome sequencing investigation of animal-skin-drum-associated UK anthrax cases reveals evidence of mixed populations and relatedness to a US case. <i>Microbial Genomics</i> , 2015, 1, e000039.	2.0	9
28	Diagnostic Testing for Hemorrhagic Fevers in Pakistan: 2007â€“2013. <i>American Journal of Tropical Medicine and Hygiene</i> , 2014, 91, 1243-1246.	1.4	13
29	Rapid molecular detection of Lujo virus RNA. <i>Journal of Virological Methods</i> , 2014, 195, 170-173.	2.1	5
30	Crimeanâ€“Congo hemorrhagic fever nosocomial infection in a immunosuppressed patient, Pakistan: Case report and virological investigation. <i>Journal of Medical Virology</i> , 2013, 85, 501-504.	5.0	17
31	Identification and analysis of Crimean-Congo hemorrhagic fever virus from human sera in Tajikistan. <i>International Journal of Infectious Diseases</i> , 2013, 17, e1031-e1037.	3.3	16
32	Genome Sequence of Ex-Afghanistan Crimean-Congo Hemorrhagic Fever Virus SCT Strain, from an Imported United Kingdom Case in October 2012. <i>Genome Announcements</i> , 2013, 1, .	0.8	17
33	Development of a Real-Time RT-PCR Assay for the Detection of Crimean-Congo Hemorrhagic Fever Virus. <i>Vector-Borne and Zoonotic Diseases</i> , 2012, 12, 786-793.	1.5	96
34	Ticks on northward migrating birds in southern Spain during Spring, 2011. <i>Journal of Vector Ecology</i> , 2012, 37, 478-480.	1.0	2
35	Crimean-Congo Hemorrhagic Fever in Tajikistan. <i>Vector-Borne and Zoonotic Diseases</i> , 2012, 12, 722-726.	1.5	29
36	Tick-Borne Encephalitis Virus, Kyrgyzstan. <i>Emerging Infectious Diseases</i> , 2011, 17, 876-879.	4.3	25

#	ARTICLE	IF	CITATIONS
37	The COVID-OUT study protocol: COVID-19 outbreak investigation to understand workplace SARS-CoV-2 transmission in the United Kingdom. Wellcome Open Research, 0, 6, 201.	1.8	8