## Barry Atkinson

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3041037/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Specificity, cross-reactivity, and function of antibodies elicited by Zika virus infection. Science, 2016, 353, 823-826.	12.6	675
2	Detection of Zika Virus in Semen. Emerging Infectious Diseases, 2016, 22, 940-940.	4.3	286
3	Temporal and spatial analysis of the 2014–2015 Ebola virus outbreak in West Africa. Nature, 2015, 524, 97-101.	27.8	272
4	A Susceptible Mouse Model for Zika Virus Infection. PLoS Neglected Tropical Diseases, 2016, 10, e0004658.	3.0	262
5	SARS-CoV-2 shedding and infectivity. Lancet, The, 2020, 395, 1339-1340.	13.7	125
6	Development of a Real-Time RT-PCR Assay for the Detection of Crimean-Congo Hemorrhagic Fever Virus. Vector-Borne and Zoonotic Diseases, 2012, 12, 786-793.	1.5	96
7	Presence and Persistence of Zika Virus RNA in Semen, United Kingdom, 2016. Emerging Infectious Diseases, 2017, 23, 611-615.	4.3	95
8	Lineage-dependent differences in the disease progression of Zika virus infection in type-I interferon receptor knockout (A129) mice. PLoS Neglected Tropical Diseases, 2017, 11, e0005704.	3.0	56
9	Infectionâ€competent monkeypox virus contamination identified in domestic settings following an imported case of monkeypox into the <scp>UK</scp> . Environmental Microbiology, 2022, 24, 4561-4569.	3.8	45
10	Analysis of Diagnostic Findings From the European Mobile Laboratory in Guéckédou, Guinea, March 2014 Through March 2015. Journal of Infectious Diseases, 2016, 214, S250-S257.	4.0	32
11	Crimean-Congo haemorrhagic fever virus in Kazakhstan (1948-2013). International Journal of Infectious Diseases, 2015, 38, 19-23.	3.3	30
12	Crimean-Congo Hemorrhagic Fever in Tajikistan. Vector-Borne and Zoonotic Diseases, 2012, 12, 722-726.	1.5	29
13	Serological and Virological Evidence of Crimean-Congo Haemorrhagic Fever Virus Circulation in the Human Population of Borno State, Northeastern Nigeria. PLoS Neglected Tropical Diseases, 2016, 10, e0005126.	3.0	28
14	Longitudinal antibody and T cell responses in Ebola virus disease survivors and contacts: an observational cohort study. Lancet Infectious Diseases, The, 2021, 21, 507-516.	9.1	26
15	Tick-Borne Encephalitis Virus, Kyrgyzstan. Emerging Infectious Diseases, 2011, 17, 876-879.	4.3	25
16	<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. Journal of Vector Ecology, 2016, 41, 128-134.	1.0	25
17	Clinical and molecular epidemiology of Crimean-Congo hemorrhagic fever in Oman. PLoS Neglected Tropical Diseases, 2019, 13, e0007100.	3.0	25
18	Crimean–Congo hemorrhagic fever nosocomial infection in a immunosuppressed patient, Pakistan: Case report and virological investigation. Journal of Medical Virology, 2013, 85, 501-504.	5.0	17

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19	Genome Sequence of Ex-Afghanistan Crimean-Congo Hemorrhagic Fever Virus SCT Strain, from an Imported United Kingdom Case in October 2012. Genome Announcements, 2013, 1, .	0.8	17
20	Identification and analysis of Crimean-Congo hemorrhagic fever virus from human sera in Tajikistan. International Journal of Infectious Diseases, 2013, 17, e1031-e1037.	3.3	16
21	Complete Genome Sequence of Zika Virus Isolated from Semen. Genome Announcements, 2016, 4, .	0.8	16
22	Emerging arboviruses of clinical importance in Central Asia. Journal of General Virology, 2018, 99, 1172-1184.	2.9	15
23	Importation of a <i>Hyalomma lusitanicum</i> tick into the UK on a dog. Veterinary Record, 2016, 179, 415-415.	0.3	14
24	Diagnostic Testing for Hemorrhagic Fevers in Pakistan: 2007–2013. American Journal of Tropical Medicine and Hygiene, 2014, 91, 1243-1246.	1.4	13
25	Complete Genomic Sequence of Issyk-Kul Virus. Genome Announcements, 2015, 3, .	0.8	13
26	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. PLoS Neglected Tropical Diseases, 2019, 13, e0007571.	3.0	10
27	Oropouche virus cases identified in Ecuador using an optimised qRT-PCR informed by metagenomic sequencing. PLoS Neglected Tropical Diseases, 2020, 14, e0007897.	3.0	10
28	Investigation of a SARS-CoV-2 Outbreak at an Automotive Manufacturing Site in England. International Journal of Environmental Research and Public Health, 2022, 19, 6400.	2.6	10
29	Whole-genome sequencing investigation of animal-skin-drum-associated UK anthrax cases reveals evidence of mixed populations and relatedness to a US case. Microbial Genomics, 2015, 1, e000039.	2.0	9
30	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
31	Rapid molecular detection of Lujo virus RNA. Journal of Virological Methods, 2014, 195, 170-173.	2.1	5
32	A non-fatal case of hantavirus cardiopulmonary syndrome imported into the UK (ex Panama), July 2014. Journal of Clinical Virology, 2015, 67, 52-55.	3.1	5
33	Complete Genome Sequence of Seoul Virus Strain Tchoupitoulas. Genome Announcements, 2016, 4, .	0.8	3
34	Ticks on northward migrating birds in southern Spain during Spring, 2011. Journal of Vector Ecology, 2012, 37, 478-480.	1.0	2
35	Title is missing!. , 2020, 14, e0007897.		0

#	Article	IF	CITATIONS
37	Title is missing!. , 2020, 14, e0007897.		0