

# Clifton D Mckee

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5338531/publications.pdf>

Version: 2024-02-01

24  
papers

630  
citations

687220

13  
h-index

642610

23  
g-index

29  
all docs

29  
docs citations

29  
times ranked

667  
citing authors

#	ARTICLE	IF	CITATIONS
1	Ecology, evolution and spillover of coronaviruses from bats. Nature Reviews Microbiology, 2022, 20, 299-314.	13.6	108
2	Curating the Evidence About COVID-19 for Frontline Public Health and Clinical Care: The Novel Coronavirus Research Compendium. Public Health Reports, 2022, 137, 197-202.	1.3	2
3	Why do some coronaviruses become pandemic threats when others do not?. PLoS Biology, 2022, 20, e3001652.	2.6	3
4	The Ecology of Nipah Virus in Bangladesh: A Nexus of Land-Use Change and Opportunistic Feeding Behavior in Bats. Viruses, 2021, 13, 169.	1.5	41
5	Bats are key hosts in the radiation of mammal-associated Bartonella bacteria. Infection, Genetics and Evolution, 2021, 89, 104719.	1.0	23
6	Comparison of sand fly trapping approaches for vector surveillance of Leishmania and Bartonella species in ecologically distinct, endemic regions of Peru. PLoS Neglected Tropical Diseases, 2021, 15, e0009517.	1.3	3
7	Seasonality of Date Palm Sap Feeding Behavior by Bats in Bangladesh. EcoHealth, 2021, 18, 359-371.	0.9	2
8	Bartonella species in medically important mosquitoes, Central Europe. Parasitology Research, 2020, 119, 2713-2717.	0.6	4
9	Longitudinal Study of Bacterial Infectious Agents in a Community of Small Mammals in New Mexico. Vector-Borne and Zoonotic Diseases, 2020, 20, 496-508.	0.6	9
10	Trypanosoma (Herpetosoma) diversity in rodents and lagomorphs of New Mexico with a focus on epizootological aspects of infection in Southern Plains woodrats (Neotoma micropus). PLoS ONE, 2020, 15, e0244803.	1.1	2
11	Host Phylogeny, Geographic Overlap, and Roost Sharing Shape Parasite Communities in European Bats. Frontiers in Ecology and Evolution, 2019, 7, .	1.1	34
12	Model-guided suggestions for targeted surveillance based on cattle shipments in the U.S.. Preventive Veterinary Medicine, 2018, 150, 52-59.	0.7	11
13	Survey of Parasitic Bacteria in Bat Bugs, Colorado. Journal of Medical Entomology, 2018, 55, 237-241.	0.9	8
14	Genotyping of Bartonella bacteria and their animal hosts: current status and perspectives. Parasitology, 2018, 145, 543-562.	0.7	58
15	Human Exposure to Novel Bartonella Species from Contact with Fruit Bats. Emerging Infectious Diseases, 2018, 24, 2317-2323.	2.0	41
16	Transmission pathways and spillover of an erythrocytic bacterial pathogen from domestic cats to wild felids. Ecology and Evolution, 2018, 8, 9779-9792.	0.8	23
17	Acquisition of Bartonella elizabethae by Experimentally Exposed Oriental Rat Fleas (Xenopsylla Tj ETQq1 1 0.784314 rgBT /Overlock 10 Entomology, 2018, 55, 1292-1298.	0.9	5
18	Human Exposure to Novel Bartonella Species from Contact with Fruit Bats. Emerging Infectious Diseases, 2018, 24, 2317-2323.	2.0	3

#	ARTICLE	IF	CITATIONS
19	Diversity and phylogenetic relationships among Bartonella strains from Thai bats. PLoS ONE, 2017, 12, e0181696.	1.1	30
20	Prevalence, diversity, and host associations of Bartonella strains in bats from Georgia (Caucasus). PLoS Neglected Tropical Diseases, 2017, 11, e0005428.	1.3	52
21	Molecular Survey of Bacterial Zoonotic Agents in Bats from the Country of Georgia (Caucasus). PLoS ONE, 2017, 12, e0171175.	1.1	45
22	Phylogenetic and geographic patterns of bartonella host shifts among bat species. Infection, Genetics and Evolution, 2016, 44, 382-394.	1.0	44
23	Genes located in a chromosomal inversion are correlated with territorial song in white-throated sparrows. Genes, Brain and Behavior, 2015, 14, 641-654.	1.1	43
24	Classification of Bartonella Strains Associated with Straw-Colored Fruit Bats (Eidolon helvum) across Africa Using a Multi-locus Sequence Typing Platform. PLoS Neglected Tropical Diseases, 2015, 9, e0003478.	1.3	29