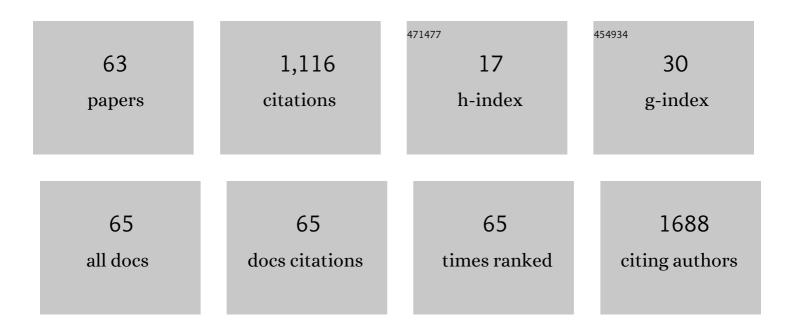
Alexandro Guterres

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	2020 taxonomic update for phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. Archives of Virology, 2020, 165, 3023-3072.	2.1	184
2	Hantavirus Reservoirs: Current Status with an Emphasis on Data from Brazil. Viruses, 2014, 6, 1929-1973.	3.3	76
3	Cat-scratch disease: ocular manifestations and visual outcome. International Ophthalmology, 2010, 30, 553-558.	1.4	55
4	What is the potential function of microRNAs as biomarkers and therapeutic targets in COVID-19?. Infection, Genetics and Evolution, 2020, 85, 104417.	2.3	55
5	Zoonotic pathogens in Atlantic Forest wild rodents in Brazil: Bartonella and Coxiella infections. Acta Tropica, 2017, 168, 64-73.	2.0	51
6	Coxiella and Bartonella spp. in bats (Chiroptera) captured in the Brazilian Atlantic Forest biome. BMC Veterinary Research, 2018, 14, 279.	1.9	41
7	Microorganisms in ticks (Acari: Ixodidae) collected on marsupials and rodents from Santa Catarina, ParanÃ _i and Mato Grosso do Sul states, Brazil. Ticks and Tick-borne Diseases, 2017, 8, 90-98.	2.7	39
8	Bartonella spp. infection in HIV positive individuals, their pets and ectoparasites in Rio de Janeiro, Brazil: Serological and molecular study. Acta Tropica, 2010, 115, 137-141.	2.0	35
9	Molecular identification of the agent of Q fever – Coxiella burnetii – in domestic animals in State of Rio de Janeiro, Brazil. Revista Da Sociedade Brasileira De Medicina Tropical, 2014, 47, 231-234.	0.9	31
10	Hantaviruses and a neglected environmental determinant. One Health, 2018, 5, 27-33.	3.4	30
11	Molecular Identification of Q Fever in Patients with a Suspected Diagnosis of Dengue in Brazil in 2013–2014. American Journal of Tropical Medicine and Hygiene, 2016, 94, 1090-1094.	1.4	26
12	Ecological study of hantavirus infection in wild rodents in an endemic area in Brazil. Acta Tropica, 2014, 131, 1-10.	2.0	22
13	Misinterpretation of viral load in COVID-19 clinical outcomes. Virus Research, 2021, 296, 198340.	2.2	21
14	Population Ecology of Hantavirus Rodent Hosts in Southern Brazil. American Journal of Tropical Medicine and Hygiene, 2014, 91, 249-257.	1.4	20
15	Rickettsia bellii, Rickettsia amblyommii, and Laguna Negra hantavirus in an Indian reserve in the Brazilian Amazon. Parasites and Vectors, 2014, 7, 191.	2.5	19
16	Detection of different South American hantaviruses. Virus Research, 2015, 210, 106-113.	2.2	19
17	First molecular detection of Coxiella burnetii in Brazilian artisanal cheese: a neglected food safety hazard in ready-to-eat raw-milk product. Brazilian Journal of Infectious Diseases, 2020, 24, 208-212.	0.6	18
18	Phylogenetic analysis of the S segment from Juquitiba hantavirus: Identification of two distinct lineages in Oligoryzomys nigripes. Infection, Genetics and Evolution, 2013, 18, 262-268.	2.3	17

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19	Hemoparasites in ticks of wild birds of Serra dos Órgãos National Park, state of Rio de Janeiro, Brazil. Brazilian Journal of Veterinary Parasitology, 2019, 28, 238-244.	0.7	17
20	Investigation of Bartonella spp. in brazilian mammals with emphasis on rodents and bats from the Atlantic Forest. International Journal for Parasitology: Parasites and Wildlife, 2020, 13, 80-89.	1.5	17
21	Characterization of rickettsia rickettsii in a case of Fatal Brazilian spotted fever in the city of Rio de Janeiro, Brazil. Brazilian Journal of Infectious Diseases, 2008, 12, 149-51.	0.6	16
22	Rio Mamore Virus and Hantavirus Pulmonary Syndrome, Brazil. Emerging Infectious Diseases, 2014, 20, 1568-1570.	4.3	16
23	Serologic evidence of the exposure of small mammals to spotted-fever Rickettsia and Rickettsia bellii in Minas Gerais, Brazil. Journal of Infection in Developing Countries, 2016, 10, 275-282.	1.2	16
24	Characterization of Juquitiba Virus in Oligoryzomys fornesi from Brazilian Cerrado. Viruses, 2014, 6, 1473-1482.	3.3	15
25	Xapuri virus, a novel mammarenavirus: natural reassortment and increased diversity between New World viruses. Emerging Microbes and Infections, 2018, 7, 1-10.	6.5	15
26	Effect of Convalescent Plasma in Critically Ill Patients With COVID-19: An Observational Study. Frontiers in Medicine, 2021, 8, 630982.	2.6	15
27	Fatal spotted fever group rickettsiosis due to Rickettsia conorii conorii mimicking a hemorrhagic viral fever in a South African traveler in Brazil. Ticks and Tick-borne Diseases, 2010, 1, 149-150.	2.7	14
28	Detection of the first incidence of Akodon paranaensis naturally infected with the Jabora virus strain (Hantavirus) in Brazil. Memorias Do Instituto Oswaldo Cruz, 2012, 107, 424-428.	1.6	14
29	Hantavirus pulmonary syndrome in a highly endemic area of Brazil. Epidemiology and Infection, 2016, 144, 1096-1106.	2.1	14
30	Hantavirus pulmonary syndrome and rodent reservoirs in the savanna-like biome of Brazil's southeastern region. Epidemiology and Infection, 2016, 144, 1107-1116.	2.1	14
31	New bunya-like viruses: Highlighting their relations. Infection, Genetics and Evolution, 2017, 49, 164-173.	2.3	13
32	A Fatal Hantavirus Pulmonary Syndrome Misdiagnosed as Dengue: An Investigation into the First Reported Case in Rio de Janeiro State, Brazil. American Journal of Tropical Medicine and Hygiene, 2017, 97, 125-129.	1.4	12
33	Detection of Latino virus (Arenaviridae: Mammarenavirus) naturally infecting Calomys callidus. Acta Tropica, 2018, 179, 17-24.	2.0	12
34	Morphological, molecular and phylogenetic characterization of Borrelia theileri in Rhipicephalus microplus. Brazilian Journal of Veterinary Parasitology, 2018, 27, 555-561.	0.7	12
35	Malaria and Hantavirus Pulmonary Syndrome in Gold Mining in the Amazon Region, Brazil. International Journal of Environmental Research and Public Health, 2019, 16, 1852.	2.6	12
36	Clinical and epidemiological use of nested PCR targeting the repetitive element IS 1111 associated with the transposase gene from Coxiella burnetii. Brazilian Journal of Microbiology, 2018, 49, 138-143.	2.0	11

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37	Co-circulation of Clade C New World Arenaviruses: New geographic distribution and host species. Infection, Genetics and Evolution, 2015, 33, 242-245.	2.3	10
38	Evaluation of HBV-Like Circulation in Wild and Farm Animals from Brazil and Uruguay. International Journal of Environmental Research and Public Health, 2019, 16, 2679.	2.6	8
39	Cyclin A in nonfunctioning pituitary adenomas. Endocrine, 2020, 70, 380-387.	2.3	8
40	Co-circulation of Araraquara and Juquitiba Hantavirus in Brazilian Cerrado. Microbial Ecology, 2018, 75, 783-789.	2.8	8
41	Detection of Rickettsia spp. in ring-tailed coatis (Nasua nasua) and ticks of the Iguaçu National Park, Brazilian Atlantic Rainforest. Ticks and Tick-borne Diseases, 2022, 13, 101891.	2.7	8
42	Aporé virus, a novel mammarenavirus (Bunyavirales: Arenaviridae) related to highly pathogenic virus from South America. Memorias Do Instituto Oswaldo Cruz, 2019, 114, e180586.	1.6	7
43	The mystery of the phylogeographic structural pattern in rodent-borne hantaviruses. Molecular Phylogenetics and Evolution, 2019, 136, 35-43.	2.7	7
44	First serological evidence of hantavirus among febrile patients in Mozambique. International Journal of Infectious Diseases, 2017, 61, 51-55.	3.3	5
45	Is the evolution of Hantavirus driven by its host?. Infection, Genetics and Evolution, 2015, 35, 142-143.	2.3	3
46	Genetic diversity of Anaplasma marginale in calves with anaplasmosis on farms in Minas Gerais, Brazil. Ticks and Tick-borne Diseases, 2021, 12, 101552.	2.7	3
47	Hantavirus pulmonary syndrome in children: case report and case series from an endemic area of Brazil. Revista Do Instituto De Medicina Tropical De Sao Paulo, 2019, 61, e65.	1.1	3
48	Serological Survey of Rabies Virus Infection among Bats in Brazil. Virus Reviews & Research: Journal of the Brazilian Society for Virology, 0, 23, 1.	0.1	3
49	Letter. Revista Da Sociedade Brasileira De Medicina Tropical, 2018, 51, 881-882.	0.9	3
50	Seroprevalence of rodent-borne viruses in Afro-descendent communities in Brazil. Revista Do Instituto De Medicina Tropical De Sao Paulo, 2019, 61, e66.	1.1	3
51	Out of the shadows, into the spotlight: Invisible zoonotic diseases in Brazil. The Lancet Regional Health Americas, 2022, 8, 100202.	2.6	3
52	What is the minimum length of gltA gene required for phylogenetic analyzes in Bartonella?. Research in Microbiology, 2019, 170, 60-64.	2.1	2
53	MicroRNAs and Mammarenaviruses: Modulating Cellular Metabolism. Cells, 2020, 9, 2525.	4.1	2
54	Telomerase expression in clinically non-functioning pituitary adenomas. Endocrine, 2021, 72, 208-215.	2.3	2

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55	Bartonella spp infections diagnosed between 2005 and 2009 by the National Rickettsial Reference Laboratory in Rio de Janeiro, Brazil. International Journal of Infectious Diseases, 2010, 14, e373.	3.3	1
56	Novel variants of human herpesvirus 2 from Brazilian HIV-1 coinfected subjects. Memorias Do Instituto Oswaldo Cruz, 2018, 113, e180328.	1.6	1
57	USP8 Somatic Mutations in Cushing's Disease and Silent Corticotropinomas. Journal of the Endocrine Society, 2021, 5, A651-A651.	0.2	1
58	Viral Loads of SARS-CoV-2 in Young Children. JAMA Pediatrics, 2021, 175, 528.	6.2	1
59	Preliminary selection and evaluation of the binding of aptamers against a Hantavirus antigen using fluorescence spectroscopy and modeling. AIP Conference Proceedings, 2015, , .	0.4	0
60	Orthohantavirus Survey in Indigenous Lands in a Savannah-Like Biome, Brazil. Viruses, 2021, 13, 1122.	3.3	0
61	Barreiras de proteção essenciais para atividades no campo. , 0, , 145-152.		0
62	A Retrospective Survey of Rodent-borne Viruses in Rural Populations of Brazilian Amazon. Revista Da Sociedade Brasileira De Medicina Tropical, 2020, 53, e20190511.	0.9	0
63	The importance of determining the amount of â€~therapeutic units'Âbefore using convalescent plasma. Future Virology, 2021, 16, 791-794.	1.8	Ο