

# FabrÃ-cio Souza Campos

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5242823/publications.pdf>

Version: 2024-02-01

73  
papers

794  
citations

516561

16  
h-index

610775

24  
g-index

82  
all docs

82  
docs citations

82  
times ranked

1105  
citing authors

#	ARTICLE	IF	CITATIONS
1	Faecal Virome Analysis of Wild Animals from Brazil. <i>Viruses</i> , 2019, 11, 803.	1.5	51
2	High prevalence of co-infections with bovine herpesvirus 1 and 5 found in cattle in southern Brazil. <i>Veterinary Microbiology</i> , 2009, 139, 67-73.	0.8	47
3	Complete Genome Sequence of Porcine Parvovirus 2 Recovered from Swine Sera. <i>Genome Announcements</i> , 2016, 4, .	0.8	40
4	Metagenomic Survey of Viral Diversity Obtained from Feces of Subantarctic and South American Fur Seals. <i>PLoS ONE</i> , 2016, 11, e0151921.	1.1	39
5	Adjuvant effect of green propolis on humoral immune response of bovines immunized with bovine herpesvirus type 5. <i>Veterinary Immunology and Immunopathology</i> , 2007, 116, 79-84.	0.5	36
6	A new marseillevirus isolated in Southern Brazil from <i>Limnoperna fortunei</i> . <i>Scientific Reports</i> , 2016, 6, 35237.	1.6	34
7	Neutralizing antibodies to bovine herpesvirus types 1 (BoHV-1) and 5 (BoHV-5) and its subtypes. <i>Veterinary Microbiology</i> , 2010, 142, 254-260.	0.8	33
8	Clinical, pathological, immunohistochemical and molecular characterization of feline chronic gingivostomatitis. <i>Journal of Feline Medicine and Surgery</i> , 2017, 19, 403-409.	0.6	32
9	Detection of bovine herpesvirus 1 and 5 in semen from Brazilian bulls. <i>Theriogenology</i> , 2011, 75, 1139-1145.	0.9	29
10	Detection of Alphacoronavirus in velvety free-tailed bats ( <i>Molossus molossus</i> ) and Brazilian free-tailed bats ( <i>Tadarida brasiliensis</i> ) from urban area of Southern Brazil. <i>Virus Genes</i> , 2013, 47, 164-167.	0.7	28
11	Real-Time Genomic Surveillance during the 2021 Re-Emergence of the Yellow Fever Virus in Rio Grande do Sul State, Brazil. <i>Viruses</i> , 2021, 13, 1976.	1.5	23
12	Comparison of the nasopharynx microbiome between influenza and non-influenza cases of severe acute respiratory infections: A pilot study. <i>Health Science Reports</i> , 2018, 1, e47.	0.6	22
13	First detection of porcine circovirus type 3 in Colombia and the complete genome sequence demonstrates the circulation of PCV 3a1 and PCV 3a2. <i>Veterinary Medicine and Science</i> , 2019, 5, 182-188.	0.6	22
14	Antibacterial activity of <i>Siparuna guianensis</i> essential oil mediated by impairment of membrane permeability and replication of pathogenic bacteria. <i>Industrial Crops and Products</i> , 2020, 146, 112142.	2.5	21
15	Prevalence of enterotoxin-encoding genes and antimicrobial resistance in coagulase-negative and coagulase-positive <i>Staphylococcus</i> isolates from black pudding. <i>Revista Da Sociedade Brasileira De Medicina Tropical</i> , 2012, 45, 579-585.	0.4	18
16	Detection of bovine herpesvirus 2 and bovine herpesvirus 4 DNA in trigeminal ganglia of naturally infected cattle by polymerase chain reaction. <i>Veterinary Microbiology</i> , 2014, 171, 182-188.	0.8	18
17	Detection of vanC 1 gene transcription in vancomycin-susceptible <i>Enterococcus faecalis</i> . <i>Memorias Do Instituto Oswaldo Cruz</i> , 2013, 108, 453-456.	0.8	17
18	Molecular detection and characterization of BK and JC polyomaviruses in urine samples of renal transplant patients in Southern Brazil. <i>Journal of Medical Virology</i> , 2015, 87, 522-528.	2.5	17

#	ARTICLE	IF	CITATIONS
19	Analysis of single-nucleotide polymorphisms in the APOBEC3H gene of domestic cats ( <i>Felis catus</i> ) and their association with the susceptibility to feline immunodeficiency virus and feline leukemia virus infections. <i>Infection, Genetics and Evolution</i> , 2014, 27, 389-394.	1.0	16
20	Genome Characterization of the First Mimiviruses of Lineage C Isolated in Brazil. <i>Frontiers in Microbiology</i> , 2017, 8, 2562.	1.5	16
21	Soroprevalência de herpesvírus bovinos tipos 1 e/ou 5 no Estado do Rio Grande do Sul. <i>Pesquisa Veterinária Brasileira</i> , 2009, 29, 767-773.	0.5	16
22	Diverse gammacoronaviruses detected in wild birds from Madagascar. <i>European Journal of Wildlife Research</i> , 2015, 61, 635-639.	0.7	15
23	Influence of a subinhibitory concentration of vancomycin on the in vitro expression of virulence-related genes in the vancomycin-resistant <i>Enterococcus faecalis</i> . <i>Revista Da Sociedade Brasileira De Medicina Tropical</i> , 2015, 48, 617-621.	0.4	12
24	Preliminary characterization of some <i>Streptomyces</i> species isolated from a composting process and their antimicrobial potential. <i>World Journal of Microbiology and Biotechnology</i> , 2010, 26, 1847-1856.	1.7	11
25	Efficacy of an inactivated, recombinant bovine herpesvirus type 5 (BoHV-5) vaccine. <i>Veterinary Microbiology</i> , 2011, 148, 18-26.	0.8	9
26	Multiplex PCR followed by restriction length polymorphism analysis for the subtyping of bovine herpesvirus 5 isolates. <i>BMC Veterinary Research</i> , 2013, 9, 111.	0.7	9
27	Development of an Indirect ELISA for Serological Diagnosis of Bovine herpesvirus 5. <i>PLoS ONE</i> , 2016, 11, e0149134.	1.1	9
28	High Rate of Mutational Events in SARS-CoV-2 Genomes across Brazilian Geographical Regions, February 2020 to June 2021. <i>Viruses</i> , 2021, 13, 1806.	1.5	9
29	Genome sequence of bubaline alphaherpesvirus 1 (BuHV1) isolated in Australia in 1972. <i>Archives of Virology</i> , 2017, 162, 1169-1176.	0.9	8
30	Biodiesel and Bioplastic Production from Waste-Cooking-Oil Transesterification: An Environmentally Friendly Approach. <i>Energies</i> , 2022, 15, 1073.	1.6	8
31	Genomic Epidemiology of SARS-CoV-2 in Tocantins State and the Diffusion of P.1.7 and AY.99.2 Lineages in Brazil. <i>Viruses</i> , 2022, 14, 659.	1.5	8
32	Viral diversity in oral cavity from <i>Sapajus nigritus</i> by metagenomic analyses. <i>Brazilian Journal of Microbiology</i> , 2020, 51, 1941-1951.	0.8	7
33	Field Evaluation of Commercial Vaccines against Infectious Bovine Rhinotracheitis (Ibr) Virus Using Different Immunization Protocols. <i>Vaccines</i> , 2021, 9, 408.	2.1	7
34	Comparative genomic analysis and mosquito larvicidal activity of four <i>Bacillus thuringiensis</i> serovar israelensis strains. <i>Scientific Reports</i> , 2020, 10, 5518.	1.6	6
35	Fecal bacterial communities of wild black capuchin monkeys ( <i>Sapajus nigritus</i> ) from the Atlantic Forest biome in Southern Brazil are divergent from those of other non-human primates. <i>Current Research in Microbial Sciences</i> , 2021, 2, 100048.	1.4	6
36	Identification of potential new mosquito-associated viruses of adult <i>Aedes aegypti</i> mosquitoes from Tocantins state, Brazil. <i>Brazilian Journal of Microbiology</i> , 2022, 53, 51-62.	0.8	6

#	ARTICLE	IF	CITATIONS
37	Comparison between DNA Detection in Trigeminal Nerve Ganglia and Serology to Detect Cattle Infected with Bovine Herpesviruses Types 1 and 5. PLoS ONE, 2016, 11, e0155941.	1.1	5
38	Fungistatic activity of essential oils for the control of bipolaris leaf spot in maize. Journal of Medicinal Plants Research, 2019, 13, 280-287.	0.2	5
39	Control of papaya fruits anthracnose by essential oils of medicinal plants associated to different coatings. Journal of Medicinal Plants Research, 2020, 14, 239-246.	0.2	5
40	First Report of <i>Colletotrichum chrysophilum</i> Causing Cassava Anthracnose in Brazil. Plant Disease, 2021, 105, 1196-1196.	0.7	5
41	Serum neutralization with different types and subtypes of bovine herpesvirus 1 and 5. Pesquisa Veterinaria Brasileira, 2010, 30, 515-522.	0.5	4
42	Genome Sequence of KP-Rio/2015, a Novel Klebsiella pneumoniae ( Podoviridae ) Phage. Genome Announcements, 2016, 4, .	0.8	4
43	Secretory expression of bovine herpesvirus type 1/5 glycoprotein E in Pichia pastoris for the differential diagnosis of vaccinated or infected cattle. Protein Expression and Purification, 2017, 130, 21-27.	0.6	4
44	Multiplex RT-PCR identification of five viruses associated with the watermelon crops in the Brazilian Cerrado. African Journal of Microbiology Research, 2019, 13, 60-69.	0.4	4
45	Whole-genome analysis of natural interspecific recombinant between bovine alphaherpesviruses 1 and 5. Virus Research, 2022, 309, 198656.	1.1	4
46	A Brief History of Giant Viruses™ Studies in Brazilian Biomes. Viruses, 2022, 14, 191.	1.5	4
47	Special Issue “Viral Infections in Developing Countries”. Viruses, 2022, 14, 405.	1.5	4
48	Distribution and genetic diversity of the human polyomaviruses JC and BK in surface water and sewage treatment plant during 2009 in Porto Alegre, Southern Brazil. Brazilian Journal of Biology, 2017, 77, 459-468.	0.4	3
49	Prediction of fermentative parameters from mathematical modeling using thermotolerant probiotic yeast. Bioresource Technology Reports, 2020, 11, 100522.	1.5	3
50	Validation of conventional PCR-like alternative to SARS-CoV-2 detection with target nucleocapsid protein gene in naso-oropharyngeal samples. PLoS ONE, 2021, 16, e0257350.	1.1	3
51	No Evidence of SARS-CoV-2 Infection in Neotropical Primates Sampled During COVID-19 Pandemic in Minas Gerais and Rio Grande do Sul, Brazil. EcoHealth, 2021, 18, 414-420.	0.9	3
52	Efficacy of a gE-deleted, bovine herpesvirus 1 (BoHV-1) inactivated vaccine. Pesquisa Veterinaria Brasileira, 2009, 29, 545-551.	0.5	2
53	Complete Genome Sequences of Two Listeria monocytogenes Serovars, 1/2a and 4b, Isolated from Dairy Products in Brazil. Genome Announcements, 2015, 3, .	0.8	2
54	Modeling transmission dynamics of severe acute respiratory syndrome coronavirus 2 in São Paulo, Brazil. Revista Da Sociedade Brasileira De Medicina Tropical, 2021, 54, e05532020.	0.4	2

#	ARTICLE	IF	CITATIONS
55	Special Issue "Emerging Viruses 2020: Surveillance, Prevention, Evolution and Control" Viruses, 2021, 13, 251.	1.5	2
56	Detection of bovine herpesvirus 1 and 5 in trigeminal ganglia of beef cattle in Uruguay. Archivos De Medicina Veterinaria, 2014, 46, 451-455.	0.2	2
57	Complete Genome Sequence of Enterococcus faecalis Strain P8-1 Isolated from Wild Magellanic Penguin ( Spheniscus magellanicus ) Feces on the South Coast of Brazil. Genome Announcements, 2016, 4, .	0.8	1
58	A Spontaneous Deletion of the US1.67/US2 Genomic Region on the Bovine Herpesvirus 1 Strain Cooper. Genome Announcements, 2016, 4, .	0.8	1
59	Osteochondroma in a young cat infected by feline leukemia virus. Ciencia Rural, 2017, 47, .	0.3	1
60	Evaluation of the Anti-apoptotic activity of bovine alphaherpesvirus type 5 US3 protein kinase in insect cells using a recombinant baculovirus. Brazilian Journal of Microbiology, 2020, 51, 827-835.	0.8	1
61	The complete chloroplast genome of Artocarpus altilis (Moraceae) and phylogenetic relationships. Mitochondrial DNA Part B: Resources, 2021, 6, 2291-2293.	0.2	1
62	Infecção latente pelo herpesvírus bovino tipo 1 em búfalos (Bubalus bubalis) no Rio Grande do Sul. Arquivo Brasileiro De Medicina Veterinaria E Zootecnia, 2019, 71, 1236-1242.	0.1	1
63	Occurrence of Colletotrichum musicola causing anthracnose on Manihot esculenta. Plant Disease, 2022, , .	0.7	1
64	Metagenomic Analysis of RNA Viruses in Watermelon. Journal of Experimental Agriculture International, 0, , 26-36.	0.3	1
65	Culture optimization of Escherichia coli for expression of gE protein from bovine herpesvirus 1 and 5. BMC Proceedings, 2014, 8, .	1.8	0
66	Genome Sequence of a New Siphoviridae Phage Found in a Brazilian Bacillus thuringiensis Serovar israelensis Strain. Genome Announcements, 2018, 6, .	0.8	0
67	Complete Sequences of Two Plasmids Found in a Brazilian Bacillus thuringiensis Serovar israelensis Strain. Microbiology Resource Announcements, 2019, 8, .	0.3	0
68	The constitutive expression of the V gene of Parainfluenza virus 5 affects the growth properties of bovine herpesvirus 5. Brazilian Archives of Biology and Technology, 2014, 57, 45-47.	0.5	0
69	Screening of beneficial bacteria associated with Calophyllum brasiliense Cambess so as to develop microbial inoculants for agriculture. Plant OMICS, 2019, , 93-101.	0.4	0
70	Soil microbiological activity under different vegetation coverages in the Cerrado biome of Tocantins state. Ciencia Florestal, 2021, 31, 1494-1511.	0.1	0
71	Root biomass under different soil uses and native Cerrado in Tocantins, Brazil. Research, Society and Development, 2022, 11, e53811125333.	0.0	0
72	Complete Genome Sequences of Two Bovine Alphaherpesvirus 5 Subtype C Strains from Southeast Brazil. Microbiology Resource Announcements, 2022, , e0122821.	0.3	0

#	ARTICLE	IF	CITATIONS
73	Special Issue "Emerging Viruses 2021: Surveillance, Prevention, Evolution and Control" Viruses, 2022, 14, 815.	1.5	0