FabrÃ-cio Souza Campos

List of Publications by Year in descending order

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516561 610775 73 794 16 24 g-index citations h-index papers 82 82 82 1105 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Identification of potential new mosquito-associated viruses of adult Aedes aegypti mosquitoes from Tocantins state, Brazil. Brazilian Journal of Microbiology, 2022, 53, 51-62.	0.8	6
2	Whole-genome analysis of natural interspecific recombinant between bovine alphaherpesviruses 1 and 5. Virus Research, 2022, 309, 198656.	1.1	4
3	Root biomass under different soil uses and native Cerrado in Tocantins, Brazil. Research, Society and Development, 2022, 11, e53811125333.	0.0	O
4	Biodiesel and Bioplastic Production from Waste-Cooking-Oil Transesterification: An Environmentally Friendly Approach. Energies, 2022, 15, 1073.	1.6	8
5	A Brief History of Giant Viruses' Studies in Brazilian Biomes. Viruses, 2022, 14, 191.	1.5	4
6	Complete Genome Sequences of Two Bovine Alphaherpesvirus 5 Subtype C Strains from Southeast Brazil. Microbiology Resource Announcements, 2022, , e0122821.	0.3	0
7	Special Issue "Viral Infections in Developing Countries― Viruses, 2022, 14, 405.	1.5	4
8	Genomic Epidemiology of SARS-CoV-2 in Tocantins State and the Diffusion of P.1.7 and AY.99.2 Lineages in Brazil. Viruses, 2022, 14, 659.	1.5	8
9	Occurrence of <i>Colletotrichum musicola</i> causing anthracnose on <i>Manihot esculenta</i> Plant Disease, 2022, , .	0.7	1
10	Special Issue "Emerging Viruses 2021: Surveillance, Prevention, Evolution and Control― Viruses, 2022, 14, 815.	1.5	0
11	First Report of <i>Colletotrichum chrysophillum</i> Causing Cassava Anthracnose in Brazil. Plant Disease, 2021, 105, 1196-1196.	0.7	5
12	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
13	Special Issue "Emerging Viruses 2020: Surveillance, Prevention, Evolution and Control― Viruses, 2021, 13, 251.	1.5	2
14	Field Evaluation of Commercial Vaccines against Infectious Bovine Rhinotracheitis (Ibr) Virus Using Different Immunization Protocols. Vaccines, 2021, 9, 408.	2.1	7
15	The complete chloroplast genome of Artocarpus altilis (Moraceae) and phylogenetic relationships. Mitochondrial DNA Part B: Resources, 2021, 6, 2291-2293.	0.2	1
16	High Rate of Mutational Events in SARS-CoV-2 Genomes across Brazilian Geographical Regions, February 2020 to June 2021. Viruses, 2021, 13, 1806.	1.5	9
17	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
18	Fecal bacterial communities of wild black capuchin monkeys (Sapajus nigritus) from the Atlantic Forest biome in Southern Brazil are divergent from those of other non-human primates. Current Research in Microbial Sciences, 2021, 2, 100048.	1.4	6

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19	Real-Time Genomic Surveillance during the 2021 Re-Emergence of the Yellow Fever Virus in Rio Grande do Sul State, Brazil. Viruses, 2021, 13, 1976.	1.5	23
20	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
21	Soil microbiological activity under different vegetation coverages in the Cerrado biome of Tocantins state. Ciencia Florestal, 2021, 31, 1494-1511.	0.1	O
22	Evaluation of theÂanti-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
23	Prediction of fermentative parameters from mathematical modeling using thermotolerant probiotic yeast. Bioresource Technology Reports, 2020, 11, 100522.	1.5	3
24	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
25	Viral diversity in oral cavity from Sapajus nigritus by metagenomic analyses. Brazilian Journal of Microbiology, 2020, 51, 1941-1951.	0.8	7
26	Antibacterial activity of Siparuna guianensis essential oil mediated by impairment of membrane permeability and replication of pathogenic bacteria. Industrial Crops and Products, 2020, 146, 112142.	2.5	21
27	Comparative genomic analysis and mosquito larvicidal activity of four Bacillus thuringiensis serovar israelensis strains. Scientific Reports, 2020, 10, 5518.	1.6	6
28	Faecal Virome Analysis of Wild Animals from Brazil. Viruses, 2019, 11, 803.	1.5	51
29	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
30	Complete Sequences of Two Plasmids Found in a Brazilian Bacillus thuringiensis Serovar israelensis Strain. Microbiology Resource Announcements, 2019, 8, .	0.3	0
31	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
32	First detection of porcine circovirus type 3 in Colombia and the complete genome sequence demonstrates the circulation of PCV 3a1 and PCV 3a2. Veterinary Medicine and Science, 2019, 5, 182-188.	0.6	22
33	Infecção latente pelo herpesvÃŧus 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
34	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
35	Comparison of the nasopharynx microbiome between influenza and nonâ€influenza cases of severe acute respiratory infections: A pilot study. Health Science Reports, 2018, 1, e47.	0.6	22
36	Genome Sequence of a New Siphoviridae Phage Found in a Brazilian Bacillus thuringiensis Serovar israelensis Strain. Genome Announcements, 2018, 6, .	0.8	0

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37	Clinical, pathological, immunohistochemical and molecular characterization of feline chronic gingivostomatitis. Journal of Feline Medicine and Surgery, 2017, 19, 403-409.	0.6	32
38	Genome sequence of bubaline alphaherpesvirus 1 (BuHV1) isolated in Australia in 1972. Archives of Virology, 2017, 162, 1169-1176.	0.9	8
39	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
40	Genome Characterization of the First Mimiviruses of Lineage C Isolated in Brazil. Frontiers in Microbiology, 2017, 8, 2562.	1.5	16
41	Osteochondroma in a young cat infected by feline leukemia virus. Ciencia Rural, 2017, 47, .	0.3	1
42	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
43	Development of an Indirect ELISA for Serological Diagnosis of Bovine herpesvirus 5. PLoS ONE, 2016, 11, e0149134.	1.1	9
44	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
45	A new marseillevirus isolated in Southern Brazil from Limnoperna fortunei. Scientific Reports, 2016, 6, 35237.	1.6	34
46	Genome Sequence of KP-Rio/2015, a Novel Klebsiella pneumoniae (Podoviridae) Phage. Genome Announcements, 2016, 4, .	0.8	4
47	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
48	Complete Genome Sequence of Porcine Parvovirus 2 Recovered from Swine Sera. Genome Announcements, 2016, 4, .	0.8	40
49	A Spontaneous Deletion of the US1.67/US2 Genomic Region on the Bovine Herpesvirus 1 Strain Cooper. Genome Announcements, 2016, 4, .	0.8	1
50	Metagenomic Survey of Viral Diversity Obtained from Feces of Subantarctic and South American Fur Seals. PLoS ONE, 2016, 11, e0151921.	1.1	39
51	Molecular detection and characterization of BK and JC polyomaviruses in urine samples of renal transplant patients in Southern Brazil. Journal of Medical Virology, 2015, 87, 522-528.	2.5	17
52	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
53	Influence of a subinhibitory concentration of vancomycin on the in vitro expression of virulence-related genes in the vancomycin-resistant Enterococcus faecalis. Revista Da Sociedade Brasileira De Medicina Tropical, 2015, 48, 617-621.	0.4	12
54	Diverse gammacoronaviruses detected in wild birds from Madagascar. European Journal of Wildlife Research, 2015, 61, 635-639.	0.7	15

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55	Culture optimization of Escherichia coli for expression of gE proteinfrombovine herpesvirus 1 and 5 . BMC Proceedings, $2014,8,.$	1.8	O
56	Analysis of single-nucleotide polymorphisms in the APOBEC3H gene of domestic cats (Felis catus) and their association with the susceptibility to feline immunodeficiency virus and feline leukemia virus infections. Infection, Genetics and Evolution, 2014, 27, 389-394.	1.0	16
57	Detection of bovine herpesvirus 2 and bovine herpesvirus 4 DNA in trigeminal ganglia of naturally infected cattle by polymerase chain reaction. Veterinary Microbiology, 2014, 171, 182-188.	0.8	18
58	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
59	The constitutive expression of the ν 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	O
60	Multiplex PCR followed by restriction length polymorphism analysis for the subtyping of bovine herpesvirus 5 isolates. BMC Veterinary Research, 2013, 9, 111.	0.7	9
61	Detection of Alphacoronavirus in velvety free-tailed bats (Molossus molossus) and Brazilian free-tailed bats (Tadarida brasiliensis) from urban area of Southern Brazil. Virus Genes, 2013, 47, 164-167.	0.7	28
62	Detection of vanC 1 gene transcription in vancomycin-susceptible Enterococcus faecalis. Memorias Do Instituto Oswaldo Cruz, 2013, 108, 453-456.	0.8	17
63	Prevalence of enterotoxin-encoding genes and antimicrobial resistance in coagulase-negative and coagulase-positive Staphylococcus isolates from black pudding. Revista Da Sociedade Brasileira De Medicina Tropical, 2012, 45, 579-585.	0.4	18
64	Detection of bovine herpesvirus 1 and 5 in semen from Brazilian bulls. Theriogenology, 2011, 75, 1139-1145.	0.9	29
65	Efficacy of an inactivated, recombinant bovine herpesvirus type 5 (BoHV-5) vaccine. Veterinary Microbiology, 2011, 148, 18-26.	0.8	9
66	Preliminary characterization of some Streptomyces species isolated from a composting process and their antimicrobial potential. World Journal of Microbiology and Biotechnology, 2010, 26, 1847-1856.	1.7	11
67	Neutralizing antibodies to bovine herpesvirus types 1 (BoHV-1) and 5 (BoHV-5) and its subtypes. Veterinary Microbiology, 2010, 142, 254-260.	0.8	33
68	Serum neutralization with different types and subtypes of bovine herpesvirus 1 and 5. Pesquisa Veterinaria Brasileira, 2010, 30, 515-522.	0.5	4
69	Efficacy of a gE-deleted, bovine herpesvirus 1 (BoHV-1) inactivated vaccine. Pesquisa Veterinaria Brasileira, 2009, 29, 545-551.	0.5	2
70	High prevalence of co-infections with bovine herpesvirus 1 and 5 found in cattle in southern Brazil. Veterinary Microbiology, 2009, 139, 67-73.	0.8	47
71	Soroprevalência de herpesvÃfus bovinos tipos 1 e/ou 5 no Estado do Rio Grande do Sul. Pesquisa Veterinaria Brasileira, 2009, 29, 767-773.	0.5	16
72	Adjuvant effect of green propolis on humoral immune response of bovines immunized with bovine herpesvirus type 5. Veterinary Immunology and Immunopathology, 2007, 116, 79-84.	0.5	36

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73	Metagenomic Analysis of RNA Viruses in Watermelon. Journal of Experimental Agriculture International, 0, , 26-36.	0.3	1