LHVGGil

List of Publications by Year in descending order

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47 papers

1,429 citations

331642 21 h-index 330122 37 g-index

48 all docs

48 docs citations

48 times ranked

2397 citing authors

#	Article	IF	Citations
1	Serologic evidence of West Nile virus and Saint Louis encephalitis virus in horses from Southern Brazil. Brazilian Journal of Microbiology, 2021, 52, 1021-1027.	2.0	4
2	Absence of norovirus contamination in shellfish harvested and commercialized in the Northeast coast of Brazil. Brazilian Journal of Medical and Biological Research, 2020, 53, e9529.	1.5	4
3	Seroprevalence of selected flaviviruses in free-living and captive capuchin monkeys in the state of Pernambuco, Brazil. Transboundary and Emerging Diseases, 2018, 65, 1094-1097.	3.0	25
4	Current status, challenges and perspectives in the development of vaccines against yellow fever, dengue, Zika and chikungunya viruses. Acta Tropica, 2018, 182, 257-263.	2.0	52
5	Development and characterization of a packaging cell line for pseudo-infectious yellow fever virus particle generation. Revista Da Sociedade Brasileira De Medicina Tropical, 2018, 51, 66-70.	0.9	2
6	A scoping review of Chikungunya virus infection: epidemiology, clinical characteristics, viral co-circulation complications, and control. Acta Tropica, 2018, 188, 213-224.	2.0	91
7	Viral immunogenicity determines epidemiological fitness in a cohort of DENV-1 infection in Brazil. PLoS Neglected Tropical Diseases, 2018, 12, e0006525.	3.0	17
8	Serum cytokine/chemokine profiles in patients with dengue fever (DF) and dengue hemorrhagic fever (FHD) by using protein array. Journal of Clinical Virology, 2017, 89, 39-45.	3.1	19
9	Risk Analysis and Occurrence of Hepatitis E Virus (HEV) in Domestic Swine in Northeast Brazil. Food and Environmental Virology, 2017, 9, 256-259.	3.4	9
10	Systems Biology Reveals NS4B-Cyclophilin A Interaction: A New Target to Inhibit YFV Replication. Journal of Proteome Research, 2017, 16, 1542-1555.	3.7	17
11	Construction and characterization of a recombinant yellow fever virus stably expressing Gaussia luciferase. Anais Da Academia Brasileira De Ciencias, 2017, 89, 2119-2130.	0.8	7
12	Hydrocephalus and arthrogryposis in an immunocompetent mouse model of ZIKA teratogeny: A developmental study. PLoS Neglected Tropical Diseases, 2017, 11, e0005363.	3.0	43
13	Perspectives on the Zika outbreak: herd immunity, antibody-dependent enhancement and vaccine. Revista Do Instituto De Medicina Tropical De Sao Paulo, 2017, 59, e21.	1.1	8
14	Primary dengue haemorrhagic fever in patients from northeast of Brazil is associated with high levels of interferon- \hat{l}^2 during acute phase. Memorias Do Instituto Oswaldo Cruz, 2016, 111, 378-384.	1.6	20
15	Use of homologous recombination in yeast to create chimeric bovine viral diarrhea virus cDNA clones. Brazilian Journal of Microbiology, 2016, 47, 993-999.	2.0	4
16	In vivo near-infrared fluorescence imaging of Leishmania amazonensis expressing infrared fluorescence protein (iRFP) for real-time monitoring of cutaneous leishmaniasis in mice. Journal of Microbiological Methods, 2016, 130, 189-195.	1.6	14
17	Dengue in the State of Rio Grande do Norte, Brazil, 2010–2012. Tropical Medicine and International Health, 2015, 20, 1707-1710.	2.3	5
18	Full-length infectious clone of a low passage dengue virus serotype 2 from Brazil. Memorias Do Instituto Oswaldo Cruz, 2015, 110, 677-683.	1.6	4

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19	Molecular phylogeny, diversity, symbiosis and discover of bioactive compounds of endophytic fungi associated with the medicinal Amazonian plant Carapa guianensis Aublet (Meliaceae). Biochemical Systematics and Ecology, 2015, 59, 36-44.	1.3	49
20	Efficient assembly of full-length infectious clone of Brazilian IBDV isolate by homologous recombination in yeast. Brazilian Journal of Microbiology, 2014, 45, 1555-1563.	2.0	4
21	Diversity Patterns, Ecology and Biological Activities of Fungal Communities Associated with the Endemic Macroalgae Across the Antarctic Peninsula. Microbial Ecology, 2014, 67, 775-787.	2.8	106
22	Insertion and stable expression of Gaussia luciferase gene by the genome of bovine viral diarrhea virus. Research in Veterinary Science, 2014, 97, 439-448.	1.9	5
23	A two-plasmid strategy for engineering a dengue virus type 3 infectious clone from primary Brazilian isolate. Anais Da Academia Brasileira De Ciencias, 2014, 86, 1749-1759.	0.8	10
24	Complement factor H gene (CFH) polymorphisms C-257T, G257A and haplotypes are associated with protection against severe dengue phenotype, possible related with high CFH expression. Human Immunology, 2013, 74, 1225-1230.	2.4	21
25	HLA-B*44 Is Associated with Dengue Severity Caused by DENV-3 in a Brazilian Population. Journal of Tropical Medicine, 2013, 2013, 1-11.	1.7	20
26	Identification of Conserved and HLA Promiscuous DENV3 T-Cell Epitopes. PLoS Neglected Tropical Diseases, 2013, 7, e2497.	3.0	39
27	Construction and characterisation of a complete reverse genetics system of dengue virus type 3. Memorias Do Instituto Oswaldo Cruz, 2013, 108, 983-991.	1.6	12
28	Construction of yellow fever virus subgenomic replicons by yeast-based homologous recombination cloning technique. Anais Da Academia Brasileira De Ciencias, 2013, 85, 159-168.	0.8	7
29	Potential biomarkers for the clinical prognosis of severe dengue. Memorias Do Instituto Oswaldo Cruz, 2013, 108, 755-762.	1.6	14
30	Immune transcript variations among Aedes aegypti populations with distinct susceptibility to dengue virus serotype 2. Acta Tropica, 2012, 124, 113-119.	2.0	24
31	Description of a Prospective 17DD Yellow Fever Vaccine Cohort in Recife, Brazil. American Journal of Tropical Medicine and Hygiene, 2011, 85, 739-747.	1.4	39
32	Classification of Dengue Fever Patients Based on Gene Expression Data Using Support Vector Machines. PLoS ONE, 2010, 5, e11267.	2.5	36
33	Early molecular markers predictive of dengue hemorrhagic fever. Anais Da Academia Brasileira De Ciencias, 2009, 81, 671-677.	0.8	7
34	Identification of Continuous Human B-Cell Epitopes in the Envelope Glycoprotein of Dengue Virus Type 3 (DENV-3). PLoS ONE, 2009, 4, e7425.	2.5	23
35	Effect of the viral protein Nproon virulence of bovine viral diarrhea virus and induction of interferon type I in calves. American Journal of Veterinary Research, 2009, 70, 1117-1123.	0.6	7
36	Alternative Complement Pathway Deregulation Is Correlated with Dengue Severity. PLoS ONE, 2009, 4, e6782.	2.5	95

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37	Gene Expression Profiling during Early Acute Febrile Stage of Dengue Infection Can Predict the Disease Outcome. PLoS ONE, 2009, 4, e7892.	2.5	77
38	Modulation of PKR activity in cells infected by bovine viral diarrhea virus. Virus Research, 2006, 116, 69-77.	2.2	25
39	A Novel, Highly Selective Inhibitor of Pestivirus Replication That Targets the Viral RNA-Dependent RNA Polymerase. Journal of Virology, 2006, 80, 149-160.	3.4	78
40	The Amino-Terminal Domain of Bovine Viral Diarrhea Virus N pro Protein Is Necessary for Alpha/Beta Interferon Antagonism. Journal of Virology, 2006, 80, 900-911.	3 . 4	82
41	Involvement of a Bovine Viral Diarrhea Virus NS5B Locus in Virion Assembly. Journal of Virology, 2004, 78, 9612-9623.	3.4	21
42	The envelope glycoprotein E2 is a determinant of cell culture tropism in ruminant pestiviruses. Journal of General Virology, 2003, 84, 1269-1274.	2.9	69
43	Role of bovine viral diarrhea virus biotype in the establishment of fetal infections. American Journal of Veterinary Research, 2002, 63, 1455-1463.	0.6	33
44	Phylogenetic analysis of Brazilian bovine viral diarrhea virus type 2 (BVDV-2) isolates: evidence for a subgenotype within BVDV-2. Virus Research, 2002, 87, 51-60.	2.2	126
45	Clinical, pathological and antigenic aspects of bovine viral diarrhea virus (BVDV) type 2 isolates identified in Brazil. Veterinary Microbiology, 2000, 77, 175-183.	1.9	44
46	Identificação do vÃŧus da Diarréia Viral Bovina tipo 2 (BVDV-2) no sul do Brasil. Pesquisa Veterinaria Brasileira, 2000, 20, 85-89.	0.5	1
47	Caracterização preliminar de amostras do vÃrus da Diarréia Viral Bovina (BVDV) isoladas no Brasil. Pesquisa Veterinaria Brasileira, 1998, 18, 84-92.	0.5	10