

Rodrigo Jardim

List of Publications by Year in descending order

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Version: 2024-02-01

17
papers

300
citations

1306789

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940134

16
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20
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20
docs citations

20
times ranked

754
citing authors

#	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.	0.9	184
2	The Comparative Genomics and Phylogenomics of <i>Leishmania Amazonensis</i> Parasite. Evolutionary Bioinformatics, 2014, 10, EBO.S13759.	0.6	23
3	Multilocus Analysis Resolves the European Finch Epidemic Strain of <i>Trichomonas gallinae</i> and Suggests Introgression from Divergent Trichomonads. Genome Biology and Evolution, 2019, 11, 2391-2402.	1.1	17
4	Systematic Identification and Classification of β -Lactamases Based on Sequence Similarity Criteria: β -Lactamase Annotation. Evolutionary Bioinformatics, 2018, 14, 117693431879735.	0.6	15
5	Core of the saliva microbiome: an analysis of the MG-RAST data. BMC Oral Health, 2021, 21, 351.	0.8	11
6	Água de reuso: uma alternativa sustentável para o Brasil. Engenharia Sanitaria E Ambiental, 2020, 25, 791-808.	0.1	9
7	An Orthology-Based Analysis of Pathogenic Protozoa Impacting Global Health: An Improved Comparative Genomics Approach with Prokaryotes and Model Eukaryote Orthologs. OMICS A Journal of Integrative Biology, 2014, 18, 524-538.	1.0	8
8	Aporã virus, a novel mammarenavirus (Bunyavirales: Arenaviridae) related to highly pathogenic virus from South America. Memórias Do Instituto Oswaldo Cruz, 2019, 114, e180586.	0.8	7
9	STINGRAY: system for integrated genomic resources and analysis. BMC Research Notes, 2014, 7, 132.	0.6	5
10	Events linked to Geosmin and 2-methylisoborneol (2-MIB) in a Water Supply in the State of Rio de Janeiro, Brazil: a case study. International Journal of Hydrology, 2021, 5, 214-220.	0.2	4
11	Mining of potential drug targets through the identification of essential and analogous enzymes in the genomes of pathogens of <i>Glycine max</i> , <i>Zea mays</i> and <i>Solanum lycopersicum</i> . PLoS ONE, 2018, 13, e0197511.	1.1	3
12	Improved orthologous databases to ease protozoan targets inference. Parasites and Vectors, 2015, 8, 494.	1.0	2
13	Managing workflows on top of a cloud computing orchestrator for using heterogeneous environments on e-Science. International Journal of Web and Grid Services, 2017, 13, 375.	0.4	2
14	Evaluating FAIRness of Genomic Databases. Lecture Notes in Computer Science, 2020, , 128-137.	1.0	2
15	ProtozoaDB 2.0: A <i>Trypanosoma Brucei</i> Case Study. Pathogens, 2017, 6, 32.	1.2	1
16	Homology Inference Based on a Reconciliation Approach for the Comparative Genomics of Protozoa. Evolutionary Bioinformatics, 2018, 14, 117693431878513.	0.6	1
17	GAP: Enhancing Semantic Interoperability of Genomic Datasets and Provenance Through Nanopublications. Communications in Computer and Information Science, 2022, , 336-348.	0.4	0