

# Jedson Ferreira Cardoso

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4158138/publications.pdf>

Version: 2024-02-01

14  
papers

1,228  
citations

840776

11  
h-index

1058476

14  
g-index

14  
all docs

14  
docs citations

14  
times ranked

3242  
citing authors

#	ARTICLE	IF	CITATIONS
1	Differential expression analysis and profiling of hepatic miRNA and isomiRNA in dengue hemorrhagic fever. <i>Scientific Reports</i> , 2021, 11, 5554.	3.3	12
2	High Genomic Variability in Equine Infectious Anemia Virus Obtained from Naturally Infected Horses in Pantanal, Brazil: An Endemic Region Case. <i>Viruses</i> , 2020, 12, 207.	3.3	7
3	Comparative analysis of the transcriptome of the Amazonian fish species <i>Colossoma macropomum</i> (tambaqui) and hybrid tambacu by next generation sequencing. <i>PLoS ONE</i> , 2019, 14, e0212755.	2.5	12
4	Clinical and molecular diagnosis of <i>Chlamydia</i> in captive parrots in Pará State, Brazil. <i>Seminário de Ciências Agrárias</i> , 2019, 40, 2603.	0.3	2
5	Oropouche orthobunyavirus: Genetic characterization of full-length genomes and development of molecular methods to discriminate natural reassortments. <i>Infection, Genetics and Evolution</i> , 2019, 68, 16-22.	2.3	16
6	Description and phylogeny of the mitochondrial genome of <i>Sabethes chloropterus</i> , <i>Sabethes glaucodaemon</i> and <i>Sabethes belisarioi</i> (Diptera: Culicidae). <i>Genomics</i> , 2019, 111, 607-611.	2.9	24
7	De novo transcriptome based on next-generation sequencing reveals candidate genes with sex-specific expression in <i>Arapaima gigas</i> (Schinz, 1822), an ancient Amazonian freshwater fish. <i>PLoS ONE</i> , 2018, 13, e0206379.	2.5	13
8	Oropouche Virus: Clinical, Epidemiological, and Molecular Aspects of a Neglected Orthobunyavirus. <i>American Journal of Tropical Medicine and Hygiene</i> , 2017, 96, 16-0672.	1.4	81
9	Identification and characterization of the expression profile of the microRNAs in the Amazon species <i>Colossoma macropomum</i> by next generation sequencing. <i>Genomics</i> , 2017, 109, 67-74.	2.9	14
10	Characterization of the Bujaru, frijoles and Tapara antigenic complexes into the sandfly fever group and two unclassified phleboviruses from Brazil. <i>Journal of General Virology</i> , 2017, 98, 585-594.	2.9	15
11	Zika virus epidemic in Brazil. I. Fatal disease in adults: Clinical and laboratorial aspects. <i>Journal of Clinical Virology</i> , 2016, 85, 56-64.	3.1	74
12	Zika virus in the Americas: Early epidemiological and genetic findings. <i>Science</i> , 2016, 352, 345-349.	12.6	877
13	Genome-Wide Study of the Defective Sucrose Fermenter Strain of <i>Vibrio cholerae</i> from the Latin American Cholera Epidemic. <i>PLoS ONE</i> , 2012, 7, e37283.	2.5	13
14	Phylogeography of Dengue Virus Serotype 4, Brazil, 2010–2011. <i>Emerging Infectious Diseases</i> , 2012, 18, 1858-1864.	4.3	68