

Juliana Domett Siqueira

List of Publications by Year in descending order

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Version: 2024-02-01

16
papers

368
citations

687363

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888059

17
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19
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19
docs citations

19
times ranked

922
citing authors

#	ARTICLE	IF	CITATIONS
1	Complex virome in feces from Amerindian children in isolated Amazonian villages. <i>Nature Communications</i> , 2018, 9, 4270.	12.8	51
2	Analysis of the cervical microbiome and potential biomarkers from postpartum HIV-positive women displaying cervical intraepithelial lesions. <i>Scientific Reports</i> , 2017, 7, 17364.	3.3	35
3	Composite Analysis of the Virome and Bacteriome of HIV/HPV Co-Infected Women Reveals Proxies for Immunodeficiency. <i>Viruses</i> , 2019, 11, 422.	3.3	30
4	SARS-CoV-2 genomic analyses in cancer patients reveal elevated intrahost genetic diversity. <i>Virus Evolution</i> , 2021, 7, veab013.	4.9	26
5	ENDEMIC INFECTION OF STRANDED SOUTHERN SEA OTTERS (<i>ENHYDRA LUTRIS NEREIS</i>) WITH NOVEL PARVOVIRUS, POLYOMAVIRUS, AND ADENOVIRUS. <i>Journal of Wildlife Diseases</i> , 2017, 53, 532-542.	0.8	25
6	HIV/HPV co-infection during pregnancy in southeastern Brazil: Prevalence, HPV types, cytological abnormalities and risk factors. <i>Gynecologic Oncology</i> , 2013, 128, 107-112.	1.4	23
7	Estimating HIV-1 Genetic Diversity in Brazil Through Next-Generation Sequencing. <i>Frontiers in Microbiology</i> , 2019, 10, 749.	3.5	23
8	New infections by SARS-CoV-2 variants of concern after natural infections and post-vaccination in Rio de Janeiro, Brazil. <i>Infection, Genetics and Evolution</i> , 2021, 94, 104998.	2.3	22
9	Identification of novel human papillomavirus lineages and sublineages in HIV/HPV-coinfected pregnant women by next-generation sequencing. <i>Virology</i> , 2016, 493, 202-208.	2.4	21
10	Assessment of the gorilla gut virome in association with natural simian immunodeficiency virus infection. <i>Retrovirology</i> , 2018, 15, 19.	2.0	21
11	Molecular diversity and polymerase gene genotypes of HIV-1 among treatment-naïve Cameroonian subjects with advanced disease. <i>Journal of Clinical Virology</i> , 2010, 48, 173-179.	3.1	17
12	Characterization of HIV-1 Near Full-Length Proviral Genome Quasispecies from Patients with Undetectable Viral Load Undergoing First-Line HAART Therapy. <i>Viruses</i> , 2017, 9, 392.	3.3	17
13	The effect of human leukocyte antigen G alleles on human papillomavirus infection and persistence in a cohort of HIV-positive pregnant women from Brazil. <i>Infection, Genetics and Evolution</i> , 2015, 34, 339-343.	2.3	16
14	Characterization and comparative analysis of a simian foamy virus complete genome isolated from Brazilian capuchin monkeys. <i>Virus Research</i> , 2015, 208, 1-6.	2.2	8
15	Bonafide, type-specific human papillomavirus persistence among HIV-positive pregnant women: predictive value for cytological abnormalities, a longitudinal cohort study. <i>Memorias Do Instituto Oswaldo Cruz</i> , 2016, 111, 120-127.	1.6	8
16	Characterisation of complete high- and low-risk human papillomavirus genomes isolated from cervical specimens in southern Brazil. <i>Memorias Do Instituto Oswaldo Cruz</i> , 2017, 112, 728-731.	1.6	7