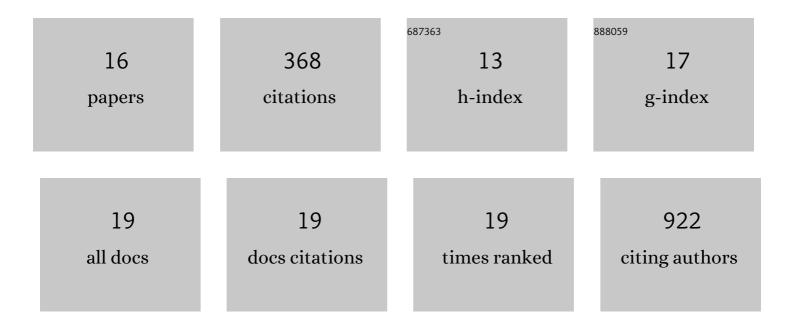
Juliana Domett Siqueira

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Complex virome in fecesÂfrom Amerindian childrenÂin isolated Amazonian villages. Nature Communications, 2018, 9, 4270.	12.8	51
2	Analysis of the cervical microbiome and potential biomarkers from postpartum HIV-positive women displaying cervical intraepithelial lesions. Scientific Reports, 2017, 7, 17364.	3.3	35
3	Composite Analysis of the Virome and Bacteriome of HIV/HPV Co-Infected Women Reveals Proxies for Immunodeficiency. Viruses, 2019, 11, 422.	3.3	30
4	SARS-CoV-2 genomic analyses in cancer patients reveal elevated intrahost genetic diversity. Virus Evolution, 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. Journal of Wildlife Diseases, 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. Gynecologic Oncology, 2013, 128, 107-112.	1.4	23
7	Estimating HIV-1 Genetic Diversity in Brazil Through Next-Generation Sequencing. Frontiers in Microbiology, 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. Infection, Genetics and Evolution, 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. Virology, 2016, 493, 202-208.	2.4	21
10	Assessment of the gorilla gut virome in association with natural simian immunodeficiency virus infection. Retrovirology, 2018, 15, 19.	2.0	21
11	Molecular diversity and polymerase gene genotypes of HIV-1 among treatment-naÃ ⁻ ve Cameroonian subjects with advanced disease. Journal of Clinical Virology, 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. Viruses, 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. Infection, Genetics and Evolution, 2015, 34, 339-343.	2.3	16
14	Characterization and comparative analysis of a simian foamy virus complete genome isolated from Brazilian capuchin monkeys. Virus Research, 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. Memorias Do Instituto Oswaldo Cruz, 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. Memorias Do Instituto Oswaldo Cruz, 2017, 112, 728-731.	1.6	7