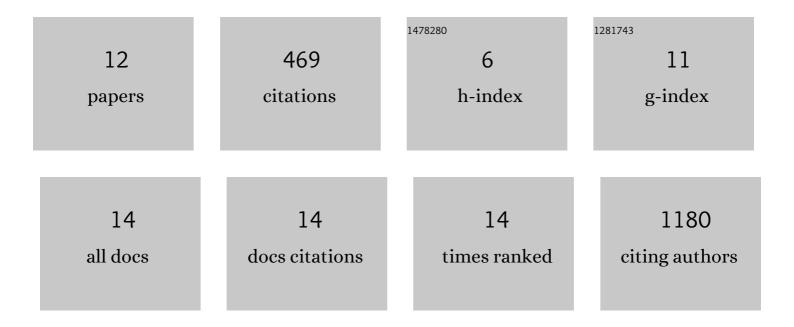
Mateus H Gouveia

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/818883/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Origin and dynamics of admixture in Brazilians and its effect on the pattern of deleterious mutations. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 8696-8701.	3.3	206
2	Genomic ancestry and ethnoracial self-classification based on 5,871 community-dwelling Brazilians (The Epigen Initiative). Scientific Reports, 2015, 5, 9812.	1.6	115
3	Origins, Admixture Dynamics, and Homogenization of the African Gene Pool in the Americas. Molecular Biology and Evolution, 2020, 37, 1647-1656.	3.5	43
4	Evaluating the Causal Link Between Malaria Infection and Endemic Burkitt Lymphoma in Northern Uganda: A Mendelian Randomization Study. EBioMedicine, 2017, 25, 58-65.	2.7	37
5	Genetic signatures of gene flow and malaria-driven natural selection in sub-Saharan populations of the "endemic Burkitt Lymphoma belt". PLoS Genetics, 2019, 15, e1008027.	1.5	23
6	Trans-ethnic meta-analysis identifies new loci associated with longitudinal blood pressure traits. Scientific Reports, 2021, 11, 4075.	1.6	13
7	Reconstructed lost Native American populations from Eastern Brazil are shaped by differential Jê/Tupi ancestry. Genome Biology and Evolution, 2019, 11, 2593-2604.	1.1	8
8	Genetics of cognitive trajectory in Brazilians: 15 years of follow-up from the BambuÃ-Epigen Cohort Study of Aging. Scientific Reports, 2019, 9, 18085.	1.6	6
9	Endemic Burkitt Lymphoma in second-degree relatives in Northern Uganda: in-depth genome-wide analysis suggests clues about genetic susceptibility. Leukemia, 2021, 35, 1209-1213.	3.3	5
10	Genomic African and Native American Ancestry and 15â€Year Cognitive Trajectory: Bambui Study, Brazil. Journal of the American Geriatrics Society, 2018, 66, 1956-1962.	1.3	2
11	The Genetics of Circulating Resistin Level, A Biomarker for Cardiovascular Diseases, Is Informed by Mendelian Randomization and the Unique Characteristics of African Genomes. Circulation Genomic and Precision Medicine, 2020, 13, 488-503.	1.6	1
12	Putative pathogen-selected polymorphisms in the PKLR gene are associated with mycobacterial susceptibility in Brazilian and African populations. PLoS Neglected Tropical Diseases, 2021, 15, e0009434.	1.3	0