Cynthia Cardoso

List of Publications by Year
in descending order

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31 papers	783 citations	566801 15 h-index	525886 27 g-index
33 all docs	33 docs citations	33 times ranked	1238 citing authors

#	Article	IF	Citations
1	Pharmacogenetics of HIV therapy: challenges for tailoring treatment in genetically complex populations. Pharmacogenomics, 2022, 23, 157-159.	0.6	0
2	Genetic susceptibility to congenital Zika syndrome: Current research and future perspectives. , 2021, , 235-244.		0
3	Exome-Wide Search for Genes Associated With Central Nervous System Inflammatory Demyelinating Diseases Following CHIKV Infection: The Tip of the Iceberg. Frontiers in Genetics, 2021, 12, 639364.	1.1	8
4	Association between ACE2 and TMPRSS2 nasopharyngeal expression and COVID-19 respiratory distress. Scientific Reports, 2021, 11, 9658.	1.6	30
5	Whole-exome sequencing reveals insights into genetic susceptibility to Congenital Zika Syndrome. PLoS Neglected Tropical Diseases, 2021, 15, e0009507.	1.3	5
6	CYP2D6 Allele Frequency in Five Malaria Vivax Endemic Areas From Brazilian Amazon Region. Frontiers in Pharmacology, 2021, 12, 542342.	1.6	3
7	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
8	Polymorphisms at CYP enzymes, NR1I2 and NR1I3 in association with virologic response to antiretroviral therapy in Brazilian HIV-positive individuals. Pharmacogenomics Journal, 2021, , .	0.9	1
9	Differential Expression of Human MicroRNAs During Dengue Virus Infection in THP-1 Monocytes. Frontiers in Cellular and Infection Microbiology, 2021, 11, 714088.	1.8	2
10	Association between Maternal Non-Coding Interferon-Î» Polymorphisms and Congenital Zika Syndrome in a Cohort from Brazilian Northeast. Viruses, 2021, 13, 2253.	1.5	1
11	Congenital Zika Syndrome Is Associated With Interferon Alfa Receptor 1. Frontiers in Immunology, 2021, 12, 764746.	2.2	9
12	Association between MBL2 haplotypes and dengue severity in children from Rio de Janeiro, Brazil. Memorias Do Instituto Oswaldo Cruz, 2019, 114, e190004.	0.8	11
13	Variations in maternal adenylate cyclase genes are associated with congenital Zika syndrome in a cohort from Northeast, Brazil. Journal of Internal Medicine, 2019, 285, 215-222.	2.7	18
14	Polymorphism of IL10, IL4, CTLA4, and DAO Genes in Crossâ€Reactive Nonsteroidal Antiâ€inflammatory Drug Hypersensitivity. Journal of Clinical Pharmacology, 2018, 58, 107-113.	1.0	12
15	Hepatitis C Virus-Infected Responders and Relapsers to Treatment Show Similar Genetic Profiles ofIL28BandIL10Single Nucleotide Polymorphisms. BioMed Research International, 2018, 2018, 1-7.	0.9	4
16	Drug metabolism and transport gene polymorphisms and efavirenz adverse effects in Brazilian HIV-positive individuals. Journal of Antimicrobial Chemotherapy, 2018, 73, 2460-2467.	1.3	17
17	Host genetics and dengue fever. Infection, Genetics and Evolution, 2017, 56, 99-110.	1.0	41
18	Genetic polymorphisms of the IL6 and NOD2 genes are risk factors for inflammatory reactions in leprosy. PLoS Neglected Tropical Diseases, 2017, 11, e0005754.	1.3	42

#	Article	IF	Citations
19	Single Nucleotide Polymorphisms in Cellular Drug Transporters Are Associated with Intolerance to Antiretroviral Therapy in Brazilian HIV-1 Positive Individuals. PLoS ONE, 2016, 11, e0163170.	1.1	8
20	Association of Single Nucleotide Polymorphisms in the Lens Epithelium-Derived Growth Factor (LEDGF/p75) with HIV-1 Infection Outcomes in Brazilian HIV-1+ Individuals. PLoS ONE, 2014, 9, e101780.	1.1	1
21	NOD2 and CCDC122-LACC1 genes are associated with leprosy susceptibility in Brazilians. Human Genetics, 2014, 133, 1525-1532.	1.8	48
22	Candidate gene case-control and functional study shows macrophage inhibitory factor (MIF) polymorphism is associated with cutaneous leishmaniasis. Cytokine, 2013, 61, 168-172.	1.4	22
23	Tumor necrosis factor (TNF) and lymphotoxin-alpha (LTA) single nucleotide polymorphisms: Importance in ARDS in septic pediatric critically ill patients. Human Immunology, 2012, 73, 661-667.	1.2	28
24	Leprosy susceptibility: genetic variations regulate innate and adaptive immunity, and disease outcome. Future Microbiology, 2011, 6, 533-549.	1.0	93
25	Genetic and functional analysis of common MRC1 exon 7 polymorphisms in leprosy susceptibility. Human Genetics, 2010, 127, 337-348.	1.8	69
26	IFNG +874 T>A single nucleotide polymorphism is associated with leprosy among Brazilians. Human Genetics, 2010, 128, 481-490.	1.8	63
27	Genetic, epidemiological and biological analysis of interleukin-10 promoter single-nucleotide polymorphisms suggests a definitive role for â˜819C/T in leprosy susceptibility. Genes and Immunity, 2009, 10, 174-180.	2.2	58
28	IFNG +874T/A, IL10 -1082G/A and TNF -308G/A polymorphisms in association with tuberculosis susceptibility: a meta-analysis study. Human Genetics, 2008, 123, 477-484.	1.8	115
29	Suppression of T and B Cell Responses by Pterodon pubescens Seeds Ethanolic Extract. Pakistan Journal of Biological Sciences, 2008, 11, 2308-2313.	0.2	12
30	Immunosuppressive effects of Echinodorus macrophyllus aqueous extract. Journal of Ethnopharmacology, 2007, 111, 435-439.	2.0	25
31	Ninjurin 1 asp110ala single nucleotide polymorphism is associated with protection in leprosy nerve damage. Journal of Neuroimmunology, 2007, 190, 131-138.	1.1	28