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	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
2	Leprosy susceptibility: genetic variations regulate innate and adaptive immunity, and disease outcome. Future Microbiology, 2011, 6, 533-549.	1.0	93
3	Genetic and functional analysis of common MRC1 exon 7 polymorphisms in leprosy susceptibility. Human Genetics, 2010, 127, 337-348.	1.8	69
4	IFNG +874 T>A single nucleotide polymorphism is associated with leprosy among Brazilians. Human Genetics, 2010, 128, 481-490.	1.8	63
5	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
6	NOD2 and CCDC122-LACC1 genes are associated with leprosy susceptibility in Brazilians. Human Genetics, 2014, 133, 1525-1532.	1.8	48
7	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
8	Host genetics and dengue fever. Infection, Genetics and Evolution, 2017, 56, 99-110.	1.0	41
9	Association between ACE2 and TMPRSS2 nasopharyngeal expression and COVID-19 respiratory distress. Scientific Reports, 2021, 11, 9658.	1.6	30
10	Ninjurin 1 asp110ala single nucleotide polymorphism is associated with protection in leprosy nerve damage. Journal of Neuroimmunology, 2007, 190, 131-138.	1.1	28
11	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
12	Immunosuppressive effects of Echinodorus macrophyllus aqueous extract. Journal of Ethnopharmacology, 2007, 111, 435-439.	2.0	25
13	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
14	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
15	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
16	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
17	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
18	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

#	Article	IF	Citations
19	Congenital Zika Syndrome Is Associated With Interferon Alfa Receptor 1. Frontiers in Immunology, 2021, 12, 764746.	2.2	9
20	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
21	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
22	Whole-exome sequencing reveals insights into genetic susceptibility to Congenital Zika Syndrome. PLoS Neglected Tropical Diseases, 2021, 15, e0009507.	1.3	5
23	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
24	CYP2D6 Allele Frequency in Five Malaria Vivax Endemic Areas From Brazilian Amazon Region. Frontiers in Pharmacology, 2021, 12, 542342.	1.6	3
25	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
26	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
27	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
28	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
29	Genetic susceptibility to congenital Zika syndrome: Current research and future perspectives. , 2021, , 235-244.		0
30	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
31	Pharmacogenetics of HIV therapy: challenges for tailoring treatment in genetically complex populations. Pharmacogenomics, 2022, 23, 157-159.	0.6	0