LÃ©a Cristina Castellucci

List of Publications by Year
in descending order

20 papers	522 citations	933447 10 h-index	752698 20 g-index
20 all docs	20 docs citations	20 times ranked	685 citing authors

#	Article	IF	Citations
1	Influence of Obesity on Clinical Manifestations and Response to Therapy in Cutaneous Leishmaniasis Caused by Leishmania braziliensis. Clinical Infectious Diseases, 2021, 73, 1020-1026.	5.8	3
2	FLI1 gene influences lesion size and skin test may predict therapeutic response in cutaneous leishmaniasis. Memorias Do Instituto Oswaldo Cruz, 2020, 115, e190361.	1.6	2
3	Serum immune markers as triggers of reactional episodes in multibacillary patients with leprosy. Leprosy Review, 2020, 91, 393-402.	0.3	1
4	Polymorphism in the interleukin-10 gene is associated with overactive bladder phenotype associated with HTLV-1 infection. Revista Da Sociedade Brasileira De Medicina Tropical, 2019, 52, e20180481.	0.9	6
5	The miRNA 361-3p, a Regulator of GZMB and TNF Is Associated With Therapeutic Failure and Longer Time Healing of Cutaneous Leishmaniasis Caused by L. (viannia) braziliensis. Frontiers in Immunology, 2018, 9, 2621.	4.8	25
6	Whole blood profiling of leprosy type 1(reversal) reactions highlights prominence of innate immune response genes. BMC Infectious Diseases, 2018, 18, 422.	2.9	6
7	Polymorphisms in genes TLR1, 2 and 4 are associated with differential cytokine and chemokine serum production in patients with leprosy. Memorias Do Instituto Oswaldo Cruz, 2017, 112, 260-268.	1.6	10
8	The â^' 308 bp TNF gene polymorphism influences tumor necrosis factor expression in leprosy patients in Bahia State, Brazil. Infection, Genetics and Evolution, 2016, 39, 147-154.	2.3	8
9	Wound healing genes and susceptibility to cutaneous leishmaniasis in Brazil: Role of COL1A1. Infection, Genetics and Evolution, 2015, 30, 225-229.	2.3	13
10	The role of ERBB2 gene polymorphisms in leprosy susceptibility. Brazilian Journal of Infectious Diseases, 2015, 19, 206-208.	0.6	5
11	Host genetic factors in American cutaneous leishmaniasis: a critical appraisal of studies conducted in an endemic area of Brazil. Memorias Do Instituto Oswaldo Cruz, 2014, 109, 279-288.	1.6	37
12	Wound healing genes and susceptibility to cutaneous leishmaniasis in Brazil. Infection, Genetics and Evolution, 2012, 12, 1102-1110.	2.3	31
13	FLI1 polymorphism affects susceptibility to cutaneous leishmaniasis in Brazil. Genes and Immunity, 2011, 12, 589-594.	4.1	27
14	The â^'2518bp promoter polymorphism at CCL2/MCP1 influences susceptibility to mucosal but not localized cutaneous leishmaniasis in Brazil. Infection, Genetics and Evolution, 2010, 10, 607-613.	2.3	34
15	CXCR1 and SLC11A1polymorphisms affect susceptibility to cutaneous leishmaniasis in Brazil: a case-control and family-based study. BMC Medical Genetics, 2010, 11, 10.	2.1	48
16	Evidence for associations between the purinergic receptor P2X7 (P2RX7) and toxoplasmosis. Genes and Immunity, 2010, 11, 374-383.	4.1	95
17	Candidate gene analysis of ocular toxoplasmosis in Brazil: evidence for a role for toll-like receptor 9 (TLR9). Memorias Do Instituto Oswaldo Cruz, 2009, 104, 1187-1190.	1.6	45
18	IL6â^'174 G/C Promoter Polymorphism Influences Susceptibility to Mucosal but Not Localized Cutaneous Leishmaniasis in Brazil. Journal of Infectious Diseases, 2006, 194, 519-527.	4.0	87

#	Article	IF	Citations
19	FAMILIAL AGGREGATION OF MUCOSAL LEISHMANIASIS IN NORTHEAST BRAZIL. American Journal of Tropical Medicine and Hygiene, 2005, 73, 69-73.	1.4	28
20	Familial aggregation of mucosal leishmaniasis in northeast Brazil. American Journal of Tropical Medicine and Hygiene, 2005, 73, 69-73.	1.4	11