

Isabelle Casademont

List of Publications by Year in descending order

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Version: 2024-02-01

17
papers

994
citations

623734

14
h-index

888059

17
g-index

17
all docs

17
docs citations

17
times ranked

1849
citing authors

#	ARTICLE	IF	CITATIONS
1	A variant in the CD209 promoter is associated with severity of dengue disease. <i>Nature Genetics</i> , 2005, 37, 507-513.	21.4	267
2	Positively Selected <i>G6PD</i> -Mahidol Mutation Reduces <i>Plasmodium vivax</i> Density in Southeast Asians. <i>Science</i> , 2009, 326, 1546-1549.	12.6	150
3	Autochthonous Japanese Encephalitis with Yellow Fever Coinfection in Africa. <i>New England Journal of Medicine</i> , 2017, 376, 1483-1485.	27.0	99
4	The large form of human 2,5-Oligoadenylate Synthetase (OAS3) exerts antiviral effect against Chikungunya virus. <i>Virology</i> , 2009, 384, 216-222.	2.4	90
5	Increased adaptive immune responses and proper feedback regulation protect against clinical dengue. <i>Science Translational Medicine</i> , 2017, 9, .	12.4	68
6	A Modified mRNA Vaccine Targeting Immunodominant NS Epitopes Protects Against Dengue Virus Infection in HLA Class I Transgenic Mice. <i>Frontiers in Immunology</i> , 2019, 10, 1424.	4.8	59
7	Genetic Determination and Linkage Mapping of <i>Plasmodium falciparum</i> Malaria Related Traits in Senegal. <i>PLoS ONE</i> , 2008, 3, e2000.	2.5	49
8	Heritability of the Human Infectious Reservoir of Malaria Parasites. <i>PLoS ONE</i> , 2010, 5, e11358.	2.5	39
9	High Anti-Dengue Virus Activity of the <i>OAS</i> Gene Family Is Associated With Increased Severity of Dengue. <i>Journal of Infectious Diseases</i> , 2015, 212, 2011-2020.	4.0	37
10	A Blood RNA Signature Detecting Severe Disease in Young Dengue Patients at Hospital Arrival. <i>Journal of Infectious Diseases</i> , 2018, 217, 1690-1698.	4.0	27
11	Impact of Mosquito Bites on Asexual Parasite Density and Gametocyte Prevalence in Asymptomatic Chronic <i>Plasmodium falciparum</i> Infections and Correlation with IgE and IgG Titers. <i>Infection and Immunity</i> , 2012, 80, 2240-2246.	2.2	25
12	Host Genetic Variation Does Not Determine Spatio-Temporal Patterns of European Bat 1 Lyssavirus. <i>Genome Biology and Evolution</i> , 2017, 9, 3202-3213.	2.5	19
13	Joint ancestry and association test indicate two distinct pathogenic pathways involved in classical dengue fever and dengue shock syndrome. <i>PLoS Neglected Tropical Diseases</i> , 2018, 12, e0006202.	3.0	17
14	Population genetics-informed meta-analysis in seven genes associated with risk to dengue fever disease. <i>Infection, Genetics and Evolution</i> , 2018, 62, 60-72.	2.3	16
15	Interaction of RNA viruses of the natural virome with the African malaria vector, <i>Anopheles coluzzii</i> . <i>Scientific Reports</i> , 2019, 9, 6319.	3.3	15
16	Heritability of <i>P. falciparum</i> and <i>P. vivax</i> Malaria in a Karen Population in Thailand. <i>PLoS ONE</i> , 2008, 3, e3887.	2.5	13
17	Detecting multi-way epistasis in family-based association studies. <i>Briefings in Bioinformatics</i> , 2016, 18, bbw039.	6.5	4