

Candelaria Vergara

List of Publications by Year in descending order

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Version: 2024-02-01

20
papers

1,037
citations

759233

12
h-index

677142

22
g-index

24
all docs

24
docs citations

24
times ranked

2598
citing authors

#	ARTICLE	IF	CITATIONS
1	Post-sequelae symptoms and comorbidities after COVID-19. <i>Journal of Medical Virology</i> , 2022, 94, 2060-2066.	5.0	14
2	Trans-ancestral fine-mapping of MHC reveals key amino acids associated with spontaneous clearance of hepatitis C in HLA-DQ1. <i>American Journal of Human Genetics</i> , 2022, 109, 299-310.	6.2	6
3	Benchmarking statistical methods for analyzing parent-child dyads in genetic association studies. <i>Genetic Epidemiology</i> , 2022, 46, 266-284.	1.3	2
4	A Multi-ancestry Sex-Stratified Genome-Wide Association Study of Spontaneous Clearance of Hepatitis C Virus. <i>Journal of Infectious Diseases</i> , 2021, 223, 2090-2098.	4.0	5
5	Whole genome sequencing identifies novel genetic mutations in patients with eczema herpeticum. <i>Allergy: European Journal of Allergy and Clinical Immunology</i> , 2021, 76, 2510-2523.	5.7	20
6	Decreased Activated CD4 ⁺ T Cell Repertoire Diversity After Antiretroviral Therapy in HIV-1/HCV Coinfection Correlates with CD4 ⁺ T Cell Recovery. <i>Viral Immunology</i> , 2021, 34, 622-631.	1.3	2
7	Multi-ancestry fine mapping of interferon lambda and the outcome of acute hepatitis C virus infection. <i>Genes and Immunity</i> , 2020, 21, 348-359.	4.1	5
8	Association of HLA-DRB1*09:01 with tIgE levels among African-ancestry individuals with asthma. <i>Journal of Allergy and Clinical Immunology</i> , 2020, 146, 147-155.	2.9	14
9	Multi-Ancestry Genome-Wide Association Study of Spontaneous Clearance of Hepatitis C Virus. <i>Gastroenterology</i> , 2019, 156, 1496-1507.e7.	1.3	32
10	Association study in African-admixed populations across the Americas recapitulates asthma risk loci in non-African populations. <i>Nature Communications</i> , 2019, 10, 880.	12.8	71
11	Assembly of a pan-genome from deep sequencing of 910 humans of African descent. <i>Nature Genetics</i> , 2019, 51, 30-35.	21.4	276
12	Genotype imputation performance of three reference panels using African ancestry individuals. <i>Human Genetics</i> , 2018, 137, 281-292.	3.8	38
13	Association analysis of exome variants and refraction, axial length, and corneal curvature in a European-American population. <i>Human Mutation</i> , 2018, 39, 1973-1979.	2.5	3
14	CMPK2 and BCL-G are associated with type 1 interferon-induced HIV restriction in humans. <i>Science Advances</i> , 2018, 4, eaat0843.	10.3	64
15	A continuum of admixture in the Western Hemisphere revealed by the African Diaspora genome. <i>Nature Communications</i> , 2016, 7, 12522.	12.8	136
16	Polymorphisms in melanoma differentiation-associated gene 5 are not associated with clearance of hepatitis C virus in a European American population. <i>Hepatology</i> , 2016, 63, 1061-1062.	7.3	3
17	Genome-Wide Association Study Identification of Novel Loci Associated with Airway Responsiveness in Chronic Obstructive Pulmonary Disease. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2015, 53, 226-234.	2.9	27
18	African Ancestry is a Risk Factor for Asthma and High Total IgE Levels in African Admixed Populations. <i>Genetic Epidemiology</i> , 2013, 37, 393-401.	1.3	46

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19	A genome-wide association study on African-ancestry populations for asthma. Journal of Allergy and Clinical Immunology, 2010, 125, 336-346.e4.	2.9	213
20	Gene Encoding Duffy Antigen/Receptor for Chemokines Is Associated with Asthma and IgE in Three Populations. American Journal of Respiratory and Critical Care Medicine, 2008, 178, 1017-1022.	5.6	51