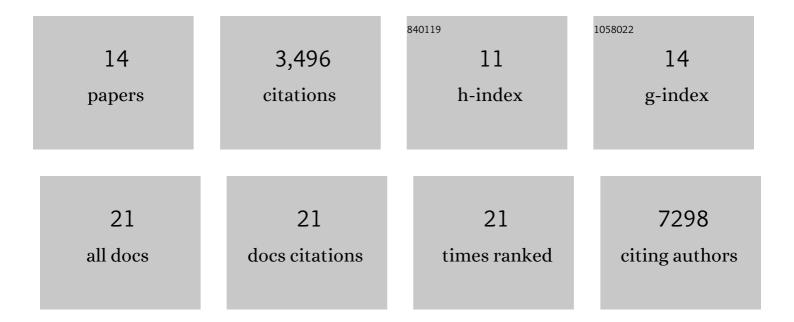
Alicia Sotomayor-GonzÃ;lez

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6668999/publications.pdf

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#	Article	IF	CITATIONS
1	Predominance of antibody-resistant SARS-CoV-2 variants in vaccine breakthrough cases from the San Francisco Bay Area, California. Nature Microbiology, 2022, 7, 277-288.	5.9	37
2	The B.1.427/1.429 (epsilon) SARS-CoV-2 variants are more virulent than ancestral B.1 (614G) in Syrian hamsters. PLoS Pathogens, 2022, 18, e1009914.	2.1	26
3	Neutralizing immunity in vaccine breakthrough infections from the SARS-CoV-2 Omicron and Delta variants. Cell, 2022, 185, 1539-1548.e5.	13.5	126
4	Contact-Tracing Outcomes Among Household Contacts of Fully Vaccinated Coronavirus Disease 2019 (COVID-19) Patients: San Francisco, California, 29 January–2 July 2021. Clinical Infectious Diseases, 2022, 75, e267-e275.	2.9	12
5	Limited cross-variant immunity from SARS-CoV-2 Omicron without vaccination. Nature, 2022, 607, 351-355.	13.7	143
6	COVID-19 Variant Detection with a High-Fidelity CRISPR-Cas12 Enzyme. Journal of Clinical Microbiology, 2022, 60, .	1.8	18
7	A diagnostic host response biosignature for COVID-19 from RNA profiling of nasal swabs and blood. Science Advances, 2021, 7, .	4.7	79
8	Transmission, infectivity, and neutralization of a spike L452R SARS-CoV-2 variant. Cell, 2021, 184, 3426-3437.e8.	13.5	424
9	SARS-CoV-2 seroprevalence and neutralizing activity in donor and patient blood. Nature Communications, 2020, 11, 4698.	5.8	124
10	Associations of Early COVID-19 Cases in San Francisco With Domestic and International Travel. Clinical Infectious Diseases, 2020, 71, 2976-2980.	2.9	8
11	Genomic surveillance reveals multiple introductions of SARS-CoV-2 into Northern California. Science, 2020, 369, 582-587.	6.0	253
12	CRISPR–Cas12-based detection of SARS-CoV-2. Nature Biotechnology, 2020, 38, 870-874.	9.4	1,884
13	Phylogenetic Analysis and Characterization of the Complete Hepatitis E Virus Genome (Zoonotic) Tj ETQq1 1 0.7	84314 rgE 1.5	ST 4Overlock
14	Isolation and characterization of porcine epidemic diarrhea virus associated with the 2014 disease outbreak in Mexico: case report. BMC Veterinary Research, 2016, 12, 132.	0.7	25