

# Aurora Fernández-García

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5986556/publications.pdf>

Version: 2024-02-01

18  
papers

1,727  
citations

1163065

8  
h-index

1281846

11  
g-index

19  
all docs

19  
docs citations

19  
times ranked

4988  
citing authors

#	ARTICLE	IF	CITATIONS
1	Prevalence of SARS-CoV-2 in Spain (ENE-COVID): a nationwide, population-based seroepidemiological study. <i>Lancet, The</i> , 2020, 396, 535-544.	13.7	1,465
2	Infection fatality risk for SARS-CoV-2 in community dwelling population of Spain: nationwide seroepidemiological study. <i>BMJ, The</i> , 2020, 371, m4509.	6.0	150
3	Shift within age-groups of mumps incidence, hospitalizations and severe complications in a highly vaccinated population. Spain, 1998–2014. <i>Vaccine</i> , 2017, 35, 4339-4345.	3.8	21
4	Measles virus genotype D4 strains with non-standard length M-F non-coding region circulated during the major outbreaks of 2011-2012 in Spain. <i>PLoS ONE</i> , 2018, 13, e0199975.	2.5	14
5	Measles in Vaccinated People: Epidemiology and Challenges in Surveillance and Diagnosis in the Post-Elimination Phase. Spain, 2014–2020. <i>Viruses</i> , 2021, 13, 1982.	3.3	14
6	Duration of immunity to measles, rubella and mumps during the first year of life. <i>Vaccine</i> , 2019, 37, 4164-4171.	3.8	13
7	ENE-COVID nationwide serosurvey served to characterize asymptomatic infections and to develop a symptom-based risk score to predict COVID-19. <i>Journal of Clinical Epidemiology</i> , 2021, 139, 240-254.	5.0	12
8	Transmission dynamics of HIV-1 subtype B in the Basque Country, Spain. <i>Infection, Genetics and Evolution</i> , 2016, 40, 91-97.	2.3	11
9	Evolution of antibodies against SARS-CoV-2 over seven months: Experience of the nationwide seroprevalence ENE-COVID study in Spain. <i>Journal of Clinical Virology</i> , 2022, 149, 105130.	3.1	9
10	Last cases of rubella and congenital rubella syndrome in Spain, 1997–2016: The success of a vaccination program. <i>Vaccine</i> , 2019, 37, 169-175.	3.8	7
11	Increase of Diversity of Mumps Virus Genotype G SH Variants Circulating Among a Highly Immunized Population: Spain, 2007–2019. <i>Journal of Infectious Diseases</i> , 2022, 227, 151-160.	4.0	4
12	Mind your Ps: A probabilistic model to aid the interpretation of molecular epidemiology data. <i>EBioMedicine</i> , 2022, 79, 103989.	6.1	1
13	Respuesta a «Implementación de técnicas moleculares para el diagnóstico de parotiditis epidémica». <i>Enfermedades Infecciosas Y Microbiología Clínica</i> , 2019, 37, 67-68.	0.5	0
14	Reply to «Implementation of molecular techniques for diagnosis of mumps». <i>Enfermedades Infecciosas Y Microbiología Clínica (English Ed)</i> , 2019, 37, 67-68.	0.3	0
15	Brote de parotiditis relacionado con el fumar en un narguile de uso público. <i>Enfermedades Infecciosas Y Microbiología Clínica</i> , 2020, 39, 503-503.	0.5	0
16	Mumps virus outbreak related to a water pipe (narghile) shared smoking. <i>Enfermedades Infecciosas Y Microbiología Clínica (English Ed)</i> , 2021, 39, 503-505.	0.3	0
17	Respuesta a «Brote de sarampión-modificado en personal sanitario tras exposición a un caso de sarampión clásico». <i>Enfermedades Infecciosas Y Microbiología Clínica</i> , 2021, 40, 344-344.	0.5	0
18	Reply to «Modified measles outbreak in vaccinated healthcare workers exposed to primary measles case». <i>Enfermedades Infecciosas Y Microbiología Clínica (English Ed)</i> , 2022, 40, 344-344.	0.3	0