## Pablo MartÃ-nez de Salazar

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Near real-time surveillance of the SARS-CoV-2 epidemic with incomplete data. PLoS Computational Biology, 2022, 18, e1009964.	3.2	8
2	Estimating internationally imported cases during the early COVID-19 pandemic. Nature Communications, 2021, 12, 311.	12.8	35
3	Estimating lengths-of-stay of hospitalised COVID-19 patients using a non-parametric model: a case study in Galicia (Spain). Epidemiology and Infection, 2021, 149, e102.	2.1	9
4	Lockdowns result in changes in human mobility which may impact the epidemiologic dynamics of SARS-CoV-2. Scientific Reports, 2021, 11, 6995.	3.3	42
5	High coverage COVID-19 mRNA vaccination rapidly controls SARS-CoV-2 transmission in long-term care facilities. Communications Medicine, 2021, 1, .	4.2	16
6	The association between gold mining and malaria in Guyana: a statistical inference and time-series analysis. Lancet Planetary Health, The, 2021, 5, e731-e738.	11.4	7
7	A nowcasting framework for correcting for reporting delays in malaria surveillance. PLoS Computational Biology, 2021, 17, e1009570.	3.2	4
8	Lockdown measures and relative changes in the age-specific incidence of SARS-CoV-2 in Spain. Epidemiology and Infection, 2020, 148, e268.	2.1	5
9	Estimating clinical severity of COVID-19 from the transmission dynamics in Wuhan, China. Nature Medicine, 2020, 26, 506-510.	30.7	1,067
10	Identifying Locations with Possible Undetected Imported Severe Acute Respiratory Syndrome Coronavirus 2 Cases by Using Importation Predictions. Emerging Infectious Diseases, 2020, 26, 1465-1469.	4.3	32
11	Using observational data to quantify bias of traveller-derived COVID-19 prevalence estimates in Wuhan, China. Lancet Infectious Diseases, The, 2020, 20, 803-808.	9.1	58
12	Practical considerations for measuring the effective reproductive number, Rt. PLoS Computational Biology, 2020, 16, e1008409.	3.2	343