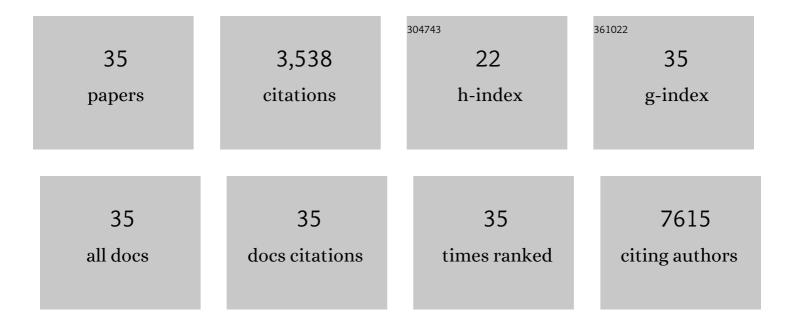
Isabel Garcia-Dorival

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Acute kidney injury in patients hospitalized with COVID-19 from the ISARIC WHO CCP-UK Study: a prospective, multicentre cohort study. Nephrology Dialysis Transplantation, 2022, 37, 271-284.	0.7	48
2	New insights into the role of endosomal proteins for African swine fever virus infection. PLoS Pathogens, 2022, 18, e1009784.	4.7	19
3	Implementation of corticosteroids in treatment of COVID-19 in the ISARIC WHO Clinical Characterisation Protocol UK: prospective, cohort study. The Lancet Digital Health, 2022, 4, e220-e234.	12.3	20
4	Procalcitonin Is Not a Reliable Biomarker of Bacterial Coinfection in People With Coronavirus Disease 2019 Undergoing Microbiological Investigation at the Time of Hospital Admission. Open Forum Infectious Diseases, 2022, 9, ofac179.	0.9	10
5	Analysis of an Ebola virus disease survivor whose host and viral markers were predictive of death indicates the effectiveness of medical countermeasures and supportive care. Genome Medicine, 2021, 13, 5.	8.2	9
6	Comparison of rhesus and cynomolgus macaques as an infection model for COVID-19. Nature Communications, 2021, 12, 1260.	12.8	115
7	Identification of potential inhibitors of protein-protein interaction useful to fight against Ebola and other highly pathogenic viruses. Antiviral Research, 2021, 186, 105011.	4.1	15
8	Inflammatory profiles across the spectrum of disease reveal a distinct role for GM-CSF in severe COVID-19. Science Immunology, 2021, 6, .	11.9	161
9	Risk of adverse outcomes in patients with underlying respiratory conditions admitted to hospital with COVID-19: a national, multicentre prospective cohort study using the ISARIC WHO Clinical Characterisation Protocol UK. Lancet Respiratory Medicine,the, 2021, 9, 699-711.	10.7	122
10	Development and validation of the ISARIC 4C Deterioration model for adults hospitalised with COVID-19: a prospective cohort study. Lancet Respiratory Medicine,the, 2021, 9, 349-359.	10.7	161
11	Importance of patient bed pathways and length of stay differences in predicting COVID-19 hospital bed occupancy in England. BMC Health Services Research, 2021, 21, 566.	2.2	22
12	African Swine Fever Virus Ubiquitin-Conjugating Enzyme Is an Immunomodulator Targeting NF-κB Activation. Viruses, 2021, 13, 1160.	3.3	25
13	Changes in in-hospital mortality in the first wave of COVID-19: a multicentre prospective observational cohort study using the WHO Clinical Characterisation Protocol UK. Lancet Respiratory Medicine,the, 2021, 9, 773-785.	10.7	78
14	Characterisation of in-hospital complications associated with COVID-19 using the ISARIC WHO Clinical Characterisation Protocol UK: a prospective, multicentre cohort study. Lancet, The, 2021, 398, 223-237.	13.7	110
15	Non-steroidal anti-inflammatory drug use and outcomes of COVID-19 in the ISARIC Clinical Characterisation Protocol UK cohort: a matched, prospective cohort study. Lancet Rheumatology, The, 2021, 3, e498-e506.	3.9	58
16	Expression of the Ebola Virus VP24 Protein Compromises the Integrity of the Nuclear Envelope and Induces a Laminopathy-Like Cellular Phenotype. MBio, 2021, 12, e0097221.	4.1	6
17	Co-infections, secondary infections, and antimicrobial use in patients hospitalised with COVID-19 during the first pandemic wave from the ISARIC WHO CCP-UK study: a multicentre, prospective cohort study. Lancet Microbe, The, 2021, 2, e354-e365.	7.3	216
18	Identification of Niemann-Pick C1 protein as a potential novel SARS-CoV-2 intracellular target. Antiviral Research, 2021, 194, 105167.	4.1	19

#	Article	IF	CITATIONS
19	Outcome of Hospitalization for COVID-19 in Patients with Interstitial Lung Disease. An International Multicenter Study. American Journal of Respiratory and Critical Care Medicine, 2020, 202, 1656-1665.	5.6	171
20	Amplicon-Based Detection and Sequencing of SARS-CoV-2 in Nasopharyngeal Swabs from Patients With COVID-19 and Identification of Deletions in the Viral Genome That Encode Proteins Involved in Interferon Antagonism. Viruses, 2020, 12, 1164.	3.3	51
21	An Investigation of the Effect of Transfected Defective, Ebola Virus Genomes on Ebola Replication. Frontiers in Cellular and Infection Microbiology, 2020, 10, 159.	3.9	4
22	Quantification of Ebola virus replication kinetics in vitro. PLoS Computational Biology, 2020, 16, e1008375.	3.2	10
23	Laboratory Findings, Compassionate Use of Favipiravir, and Outcome in Patients With Ebola Virus Disease, Guinea, 2015—A Retrospective Observational Study. Journal of Infectious Diseases, 2019, 220, 195-202.	4.0	38
24	Identification of a small molecule inhibitor of Ebola virus genome replication and transcription using in silico screening. Antiviral Research, 2018, 156, 46-54.	4.1	14
25	Transcriptomic signatures differentiate survival from fatal outcomes in humans infected with Ebola virus. Genome Biology, 2017, 18, 4.	8.8	115
26	A comparison of host gene expression signatures associated with infection in vitro by the Makona and Ecran (Mayinga) variants of Ebola virus. Scientific Reports, 2017, 7, 43144.	3.3	21
27	Deep Sequencing of RNA from Blood and Oral Swab Samples Reveals the Presence of Nucleic Acid from a Number of Pathogens in Patients with Acute Ebola Virus Disease and Is Consistent with Bacterial Translocation across the Gut. MSphere, 2017, 2, .	2.9	30
28	Different features of Vδ2 T and NK cells in fatal and non-fatal human Ebola infections. PLoS Neglected Tropical Diseases, 2017, 11, e0005645.	3.0	46
29	Antiviral Screening of Multiple Compounds against Ebola Virus. Viruses, 2016, 8, 277.	3.3	37
30	Unique human immune signature of Ebola virus disease in Guinea. Nature, 2016, 533, 100-104.	27.8	170
31	Elucidation of the Cellular Interactome of Ebola Virus Nucleoprotein and Identification of Therapeutic Targets. Journal of Proteome Research, 2016, 15, 4290-4303.	3.7	43
32	Real-time, portable genome sequencing for Ebola surveillance. Nature, 2016, 530, 228-232.	27.8	1,179
33	Temporal and spatial analysis of the 2014–2015 Ebola virus outbreak in West Africa. Nature, 2015, 524, 97-101.	27.8	272
34	Elucidating variations in the nucleotide sequence of Ebola virus associated with increasing pathogenicity. Genome Biology, 2014, 15, 540.	8.8	44
35	Elucidation of the Ebola Virus VP24 Cellular Interactome and Disruption of Virus Biology through Targeted Inhibition of Host-Cell Protein Function. Journal of Proteome Research, 2014, 13, 5120-5135.	3.7	79