

# César G Albariño

## List of Publications by Year in descending order

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32  
papers

1,818  
citations

394421

19  
h-index

414414

32  
g-index

33  
all docs

33  
docs citations

33  
times ranked

2300  
citing authors

#	ARTICLE	IF	CITATIONS
1	Isolation of Genetically Diverse Marburg Viruses from Egyptian Fruit Bats. <i>PLoS Pathogens</i> , 2009, 5, e1000536.	4.7	549
2	Marburg Virus Infection Detected in a Common African Bat. <i>PLoS ONE</i> , 2007, 2, e764.	2.5	330
3	Recovery of Recombinant Crimean Congo Hemorrhagic Fever Virus Reveals a Function for Non-structural Glycoproteins Cleavage by Furin. <i>PLoS Pathogens</i> , 2015, 11, e1004879.	4.7	61
4	Lassa and Ebola virus inhibitors identified using minigenome and recombinant virus reporter systems. <i>Antiviral Research</i> , 2016, 136, 9-18.	4.1	61
5	High Diversity and Ancient Common Ancestry of Lymphocytic Choriomeningitis Virus. <i>Emerging Infectious Diseases</i> , 2010, 16, 1093-1100.	4.3	59
6	Inhibitors of cellular kinases with broad-spectrum antiviral activity for hemorrhagic fever viruses. <i>Antiviral Research</i> , 2015, 120, 40-47.	4.1	59
7	Statins Suppress Ebola Virus Infectivity by Interfering with Glycoprotein Processing. <i>MBio</i> , 2018, 9, .	4.1	58
8	25-Hydroxycholesterol Inhibition of Lassa Virus Infection through Aberrant GP1 Glycosylation. <i>MBio</i> , 2016, 7, .	4.1	55
9	Characterisation of infectious Ebola virus from the ongoing outbreak to guide response activities in the Democratic Republic of the Congo: a phylogenetic and in vitro analysis. <i>Lancet Infectious Diseases</i> , The, 2019, 19, 1023-1032.	9.1	48
10	Efficient Rescue of Recombinant Lassa Virus Reveals the Influence of S Segment Noncoding Regions on Virus Replication and Virulence. <i>Journal of Virology</i> , 2011, 85, 4020-4024.	3.4	46
11	Development of a reverse genetics system to generate a recombinant Ebola virus Makona expressing a green fluorescent protein. <i>Virology</i> , 2015, 484, 259-264.	2.4	45
12	The lipid moiety of brincidofovir is required for in vitro antiviral activity against Ebola virus. <i>Antiviral Research</i> , 2016, 125, 71-78.	4.1	44
13	Development of a reverse genetics system to generate recombinant Marburg virus derived from a bat isolate. <i>Virology</i> , 2013, 446, 230-237.	2.4	42
14	Identification of broadly neutralizing monoclonal antibodies against Crimean-Congo hemorrhagic fever virus. <i>Antiviral Research</i> , 2017, 146, 112-120.	4.1	40
15	Reverse Genetics Generation of Chimeric Infectious Junin/Lassa Virus Is Dependent on Interaction of Homologous Glycoprotein Stable Signal Peptide and G2 Cytoplasmic Domains. <i>Journal of Virology</i> , 2011, 85, 112-122.	3.4	38
16	Novel activities by ebolavirus and marburgvirus interferon antagonists revealed using a standardized in vitro reporter system. <i>Virology</i> , 2017, 501, 147-165.	2.4	38
17	Antibody-Mediated Virus Neutralization Is Not a Universal Mechanism of Marburg, Ebola, or Sosuga Virus Clearance in Egyptian Rousette Bats. <i>Journal of Infectious Diseases</i> , 2019, 219, 1716-1721.	4.0	28
18	Use of a Scalable Replicon-Particle Vaccine to Protect Against Lethal Lassa Virus Infection in the Guinea Pig Model. <i>Journal of Infectious Diseases</i> , 2018, 217, 1957-1966.	4.0	26

#	ARTICLE	IF	CITATIONS
19	Transcriptional analysis of viral mRNAs reveals common transcription patterns in cells infected by five different filoviruses. <i>PLoS ONE</i> , 2018, 13, e0201827.	2.5	22
20	Recombinant Marburg viruses containing mutations in the IID region of VP35 prevent inhibition of Host immune responses. <i>Virology</i> , 2015, 476, 85-91.	2.4	21
21	Rousette Bat Dendritic Cells Overcome Marburg Virus-Mediated Antiviral Responses by Upregulation of Interferon-Related Genes While Downregulating Proinflammatory Disease Mediators. <i>MSphere</i> , 2019, 4, .	2.9	20
22	Marburg Virus Persistence on Fruit as a Plausible Route of Bat to Primate Filovirus Transmission. <i>Viruses</i> , 2021, 13, 2394.	3.3	20
23	The S Genome Segment Is Sufficient to Maintain Pathogenicity in Intra-Clade Lassa Virus Reassortants in a Guinea Pig Model. <i>Frontiers in Cellular and Infection Microbiology</i> , 2018, 8, 240.	3.9	18
24	Insights into Reston virus spillovers and adaption from virus whole genome sequences. <i>PLoS ONE</i> , 2017, 12, e0178224.	2.5	14
25	Small Interfering RNA Inhibition of Andes Virus Replication. <i>PLoS ONE</i> , 2014, 9, e99764.	2.5	12
26	Preliminary Evaluation of the Effect of Investigational Ebola Virus Disease Treatments on Viral Genome Sequences. <i>Journal of Infectious Diseases</i> , 2016, 214, S333-S341.	4.0	11
27	Virus fitness differences observed between two naturally occurring isolates of Ebola virus Makona variant using a reverse genetics approach. <i>Virology</i> , 2016, 496, 237-243.	2.4	10
28	Rapid Determination of Ebolavirus Infectivity in Clinical Samples Using a Novel Reporter Cell Line. <i>Journal of Infectious Diseases</i> , 2017, 216, 1380-1385.	4.0	10
29	Development of a reverse genetics system for Sosuga virus allows rapid screening of antiviral compounds. <i>PLoS Neglected Tropical Diseases</i> , 2018, 12, e0006326.	3.0	10
30	Identification and characterization of novel mosquito-borne (Kammavanpettai virus) and tick-borne (Wad Medani) reoviruses isolated in India. <i>Journal of General Virology</i> , 2018, 99, 991-1000.	2.9	9
31	Genome Sequences of Crimean-Congo Hemorrhagic Fever Virus Strains Isolated in South Africa, Namibia, and Turkey. <i>Genome Announcements</i> , 2017, 5, .	0.8	7
32	Lassa Virus Replicon Particle Vaccine Protects Strain 13/N Guinea Pigs Against Challenge With Geographically and Genetically Diverse Viral Strains. <i>Journal of Infectious Diseases</i> , 2022, 226, 1545-1550.	4.0	7