

Sonia Vázquez-Morán

List of Publications by Year in descending order

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42
papers

1,649
citations

430442

18
h-index

288905

40
g-index

42
all docs

42
docs citations

42
times ranked

2799
citing authors

#	ARTICLE	IF	CITATIONS
1	Discovery of an Ebolavirus-Like Filovirus in Europe. PLoS Pathogens, 2011, 7, e1002304.	2.1	340
2	Taxonomy of the order Mononegavirales: update 2018. Archives of Virology, 2018, 163, 2283-2294.	0.9	153
3	Viral infections of the central nervous system in Spain: A prospective study. Journal of Medical Virology, 2013, 85, 554-562.	2.5	132
4	Novel Lyssavirus in Bat, Spain. Emerging Infectious Diseases, 2013, 19, 793-795.	2.0	132
5	Phylodynamics and Human-Mediated Dispersal of a Zoonotic Virus. PLoS Pathogens, 2010, 6, e1001166.	2.1	124
6	Bat Rabies Surveillance in Europe. Zoonoses and Public Health, 2013, 60, 22-34.	0.9	116
7	Detection of alpha and betacoronaviruses in multiple Iberian bat species. Archives of Virology, 2011, 156, 1883-1890.	0.9	82
8	Endemic Circulation of European Bat Lyssavirus Type 1 in Serotine Bats, Spain. Emerging Infectious Diseases, 2008, 14, 1263-1266.	2.0	46
9	Molecular Epidemiology of Bat Lyssaviruses in Europe. Zoonoses and Public Health, 2013, 60, 35-45.	0.9	45
10	Detection of rhabdovirus viral RNA in oropharyngeal swabs and ectoparasites of Spanish bats. Journal of General Virology, 2013, 94, 69-75.	1.3	42
11	RT-PCR for detection of all seven genotypes of Lyssavirus genus. Journal of Virological Methods, 2006, 135, 281-287.	1.0	39
12	Evaluation of the diagnostic accuracy of laboratory-based screening for hepatitis C in dried blood spot samples: A systematic review and meta-analysis. Scientific Reports, 2019, 9, 7316.	1.6	35
13	Evaluation of dried blood spot samples for screening of hepatitis C and human immunodeficiency virus in a real-world setting. Scientific Reports, 2018, 8, 1858.	1.6	34
14	Identification of Novel Betaherpesviruses in Iberian Bats Reveals Parallel Evolution. PLoS ONE, 2016, 11, e0169153.	1.1	25
15	Phylogeny of European Bat Lyssavirus 1 in <i>Eptesicus isabellinus</i> Bats, Spain. Emerging Infectious Diseases, 2011, 17, 520-523.	2.0	22
16	Mx1, OAS1 and OAS2 polymorphisms are associated with the severity of liver disease in HIV/HCV-coinfected patients: A cross-sectional study. Scientific Reports, 2017, 7, 41516.	1.6	22
17	Gender-based vulnerability in women who inject drugs in a harm reduction setting. PLoS ONE, 2020, 15, e0230886.	1.1	20
18	Comparative assay of fluorescent antibody test results among twelve European National Reference Laboratories using various anti-rabies conjugates. Journal of Virological Methods, 2013, 191, 88-94.	1.0	19

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19	Relationship between ITPA polymorphisms and hemolytic anemia in HCV-infected patients after ribavirin-based therapy: a meta-analysis. <i>Journal of Translational Medicine</i> , 2015, 13, 320.	1.8	19
20	New Adenovirus Groups in Western Palaearctic Bats. <i>Viruses</i> , 2018, 10, 443.	1.5	18
21	A Step Forward in Molecular Diagnostics of Lyssaviruses – Results of a Ring Trial among European Laboratories. <i>PLoS ONE</i> , 2013, 8, e58372.	1.1	16
22	NS3 Resistance-Associated Variants (RAVs) in Patients Infected with HCV Genotype 1a in Spain. <i>PLoS ONE</i> , 2016, 11, e0163197.	1.1	16
23	CXCL9 polymorphisms are associated with liver fibrosis in patients with chronic hepatitis C: a cross-sectional study. <i>Clinical and Translational Medicine</i> , 2017, 6, 26.	1.7	13
24	Impact of patatin-like phospholipase domain-containing 3 gene polymorphism (rs738409) on severity of liver disease in HIV/hepatitis C virus-coinfected patients. <i>Aids</i> , 2016, 30, 465-470.	1.0	12
25	HCV screening based on dried blood samples and linkage to care in people who use drugs: A prospective study. <i>International Journal of Drug Policy</i> , 2021, 92, 103134.	1.6	11
26	Phylogeny of European Bat Lyssavirus 1 in <i>Eptesicus isabellinus</i> Bats, Spain. <i>Emerging Infectious Diseases</i> , 2011, 17, 520-3.	2.0	10
27	Association between IL7R polymorphisms and severe liver disease in HIV/HCV coinfecting patients: a cross-sectional study. <i>Journal of Translational Medicine</i> , 2015, 13, 206.	1.8	10
28	PNPLA3 rs738409 polymorphism is associated with liver fibrosis progression in patients with chronic hepatitis C: A repeated measures study. <i>Journal of Clinical Virology</i> , 2018, 103, 71-74.	1.6	10
29	The IL7RA rs6897932 polymorphism is associated with progression of liver fibrosis in patients with chronic hepatitis C: Repeated measurements design. <i>PLoS ONE</i> , 2018, 13, e0197115.	1.1	10
30	Detection of active hepatitis C in a single visit and linkage to care among marginalized people using a mobile unit in Madrid, Spain. <i>International Journal of Drug Policy</i> , 2021, 96, 103424.	1.6	10
31	Low frequency of NS5A relevant resistance-associated substitutions to Elbasvir among hepatitis C virus genotype 1a in Spain: a cross-sectional study. <i>Scientific Reports</i> , 2017, 7, 2892.	1.6	8
32	First cases of European bat lyssavirus type 1 in Iberian serotine bats: Implications for the molecular epidemiology of bat rabies in Europe. <i>PLoS Neglected Tropical Diseases</i> , 2018, 12, e0006290.	1.3	8
33	Prevalence of hepatitis E infection in HIV/HCV-coinfecting patients in Spain (2012–2014). <i>Scientific Reports</i> , 2019, 9, 1143.	1.6	8
34	Optimal vitamin D plasma levels are associated with lower bacterial DNA translocation in HIV/hepatitis c virus coinfecting patients. <i>Aids</i> , 2016, 30, 1069-1074.	1.0	7
35	FTO rs9939609 polymorphism is associated with metabolic disturbances and response to HCV therapy in HIV/HCV-coinfecting patients. <i>BMC Medicine</i> , 2014, 12, 198.	2.3	7
36	Emergence of Progressive Mutations in SARS-CoV-2 From a Hematologic Patient With Prolonged Viral Replication. <i>Frontiers in Microbiology</i> , 2022, 13, 826883.	1.5	7

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37	Complete Genomic Sequence of European Bat Lyssavirus 1, Isolated from <i>Eptesicus isabellinus</i> in Spain. <i>Genome Announcements</i> , 2015, 3, .	0.8	5
38	<i>IL15</i> polymorphism is associated with advanced fibrosis, inflammation-related biomarkers and virological response in human immunodeficiency virus/hepatitis C virus coinfection. <i>Liver International</i> , 2016, 36, 1258-1266.	1.9	5
39	FTOrs9939609 polymorphism is associated with metabolic disturbances and response to HCV therapy in HIV/HCV-coinfected patients. <i>BMC Medicine</i> , 2014, 12, 198.	2.3	4
40	Association between <i>IL7RA</i> polymorphisms and the successful therapy against HCV in HIV/HCV-coinfected patients. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2015, 34, 385-393.	1.3	4
41	Rapid decrease in titer and breadth of neutralizing anti-HCV antibodies in HIV/HCV-coinfected patients who achieved SVR. <i>Scientific Reports</i> , 2019, 9, 12163.	1.6	2
42	HIV screening and retention in care in people who use drugs in Madrid, Spain: a prospective study. <i>Infectious Diseases of Poverty</i> , 2021, 10, 111.	1.5	1