Sonia VÃ¡zquez-MorÃ³n

List of Publications by Year
in descending order

42 papers	1,649 citations	430874 18 h-index	289244 40 g-index
42 all docs	42 docs citations	42 times ranked	2799 citing authors

#	Article	IF	Citations
1	Emergence of Progressive Mutations in SARS-CoV-2 From a Hematologic Patient With Prolonged Viral Replication. Frontiers in Microbiology, 2022, 13, 826883.	3.5	7
2	HCV screening based on dried blood samples and linkage to care in people who use drugs: A prospective study. International Journal of Drug Policy, 2021, 92, 103134.	3.3	11
3	HIV screening and retention in care in people who use drugs in Madrid, Spain: a prospective study. Infectious Diseases of Poverty, 2021, 10, 111.	3.7	1
4	Detection of active hepatitis C in a single visit and linkage to care among marginalized people using a mobile unit in Madrid, Spain. International Journal of Drug Policy, 2021, 96, 103424.	3.3	10
5	Gender-based vulnerability in women who inject drugs in a harm reduction setting. PLoS ONE, 2020, 15, e0230886.	2.5	20
6	Rapid decrease in titer and breadth of neutralizing anti-HCV antibodies in HIV/HCV-coinfected patients who achieved SVR. Scientific Reports, 2019, 9, 12163.	3.3	2
7	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.	3.3	35
8	Prevalence of hepatitis E infection in HIV/HCV-coinfected patients in Spain (2012â€“2014). Scientific Reports, 2019, 9, 1143.	3.3	8
9	PNPLA3 rs738409 polymorphism is associated with liver fibrosis progression in patients with chronic hepatitis C: A repeated measures study. Journal of Clinical Virology, 2018, 103, 71-74.	3.1	10
10	Taxonomy of the order Mononegavirales: update 2018. Archives of Virology, 2018, 163, 2283-2294.	2.1	153
11	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.	3.3	34
12	The IL7RA rs6897932 polymorphism is associated with progression of liver fibrosis in patients with chronic hepatitis C: Repeated measurements design. PLoS ONE, 2018, 13, e0197115.	2.5	10
13	New Adenovirus Groups in Western Palaearctic Bats. Viruses, 2018, 10, 443.	3.3	18
14	First cases of European bat lyssavirus type 1 in Iberian serotine bats: Implications for the molecular epidemiology of bat rabies in Europe. PLoS Neglected Tropical Diseases, 2018, 12, e0006290.	3.0	8
15	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.	3.3	22
16	Low frequency of NS5A relevant resistance-associated substitutions to Elbasvir among hepatitis C virus genotype 1a in Spain: a cross-sectional study. Scientific Reports, 2017, 7, 2892.	3.3	8
17	CXCL9â€11 polymorphisms are associated with liver fibrosis in patients with chronic hepatitis C: a crossâ€sectional study. Clinical and Translational Medicine, 2017, 6, 26.	4.0	13
18	Identification of Novel Betaherpesviruses in Iberian Bats Reveals Parallel Evolution. PLoS ONE, 2016, 11, e0169153.	2.5	25

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19	Optimal vitamin D plasma levels are associated with lower bacterial DNA translocation in HIV/hepatitis c virus coinfected patients. Aids, 2016, 30, 1069-1074.	2.2	7
20	IL15 polymorphism is associated with advanced fibrosis, inflammationâ€related biomarkers and virological response in human immunodeficiency virus/hepatitis C virus coinfection. Liver International, 2016, 36, 1258-1266.	3.9	5
21	Impact of patatin-like phospholipase domain-containing 3 gene polymorphism (rs738409) on severity of liver disease in HIV/hepatitis C virus-coinfected patients. Aids, 2016, 30, 465-470.	2.2	12
22	NS3 Resistance-Associated Variants (RAVs) in Patients Infected with HCV Genotype 1a in Spain. PLoS ONE, 2016, 11, e0163197.	2.5	16
23	Relationship between ITPA polymorphisms and hemolytic anemia in HCV-infected patients after ribavirin-based therapy: a meta-analysis. Journal of Translational Medicine, 2015, 13, 320.	4.4	19
24	Association between IL7R polymorphisms and severe liver disease in HIV/HCV coinfected patients: a cross-sectional study. Journal of Translational Medicine, 2015, 13, 206.	4.4	10
25	Complete Genomic Sequence of European Bat Lyssavirus 1, Isolated from Eptesicus isabellinus in Spain. Genome Announcements, 2015, 3, .	0.8	5
26	Association between IL7RA polymorphisms and the successful therapy against HCV in HIV/HCV-coinfected patients. European Journal of Clinical Microbiology and Infectious Diseases, 2015, 34, 385-393.	2.9	4
27	FTOrs9939609 polymorphism is associated with metabolic disturbances and response to HCV therapy in HIV/HCV-coinfected patients. BMC Medicine, 2014, 12, 198.	5.5	4
28	FTO rs9939609 polymorphism is associated with metabolic disturbances and response to HCV therapy in HIV/HCV-coinfected patients. BMC Medicine, 2014, 12, 198.	5.5	7
29	Viral infections of the central nervous system in Spain: A prospective study. Journal of Medical Virology, 2013, 85, 554-562.	5.0	132
30	Bat Rabies Surveillance in Europe. Zoonoses and Public Health, 2013, 60, 22-34.	2.2	116
31	Molecular Epidemiology of Bat Lyssaviruses in Europe. Zoonoses and Public Health, 2013, 60, 35-45.	2.2	45
32	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.	2.1	19
33	Detection of rhabdovirus viral RNA in oropharyngeal swabs and ectoparasites of Spanish bats. Journal of General Virology, 2013, 94, 69-75.	2.9	42
34	Novel Lyssavirus in Bat, Spain. Emerging Infectious Diseases, 2013, 19, 793-795.	4.3	132
35	A Step Forward in Molecular Diagnostics of Lyssaviruses â€“ Results of a Ring Trial among European Laboratories. PLoS ONE, 2013, 8, e58372.	2.5	16
36	Phylogeny of European Bat Lyssavirus 1 inEptesicus isabellinusBats, Spain. Emerging Infectious Diseases, 2011, 17, 520-523.	4.3	22

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37	Detection of alpha and betacoronaviruses in multiple Iberian bat species. Archives of Virology, 2011, 156, 1883-1890.	2.1	82
38	Phylogeny of European Bat Lyssavirus 1 in Eptesicus isabellinus Bats, Spain. Emerging Infectious Diseases, 2011, 17, 520-3.	4.3	10
39	Discovery of an Ebolavirus-Like Filovirus in Europe. PLoS Pathogens, 2011, 7, e1002304.	4.7	340
40	Phylodynamics and Human-Mediated Dispersal of a Zoonotic Virus. PLoS Pathogens, 2010, 6, e1001166.	4.7	124
41	Endemic Circulation of European Bat Lyssavirus Type 1 in Serotine Bats, Spain. Emerging Infectious Diseases, 2008, 14, 1263-1266.	4.3	46
42	RT-PCR for detection of all seven genotypes of Lyssavirus genus. Journal of Virological Methods, 2006, 135, 281-287.	2.1	39