Amanda FernÃ¡ndez-RodrÃ-guez

List of Publications by Year
in descending order

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79	1,032	516710 16	526287 27
papers	citations	h-index	g-index

81	81	81	2045
all docs	docs citations	times ranked	citing authors

#	ARTICLE	IF	CITATIONS
1	Similar humoral immune responses against the SARS-CoV-2 spike protein in HIV and non-HIV individuals after COVID-19. Journal of Infection, 2022, 84, 418-467.	3.3	7
2	Blood microbiome is associated with changes in portal hypertension after successful direct-acting antiviral therapy in patients with HCV-related cirrhosis. Journal of Antimicrobial Chemotherapy, 2022, 77, 719-726.	3.0	7
3	HCV eradication with DAAs differently affects HIV males and females: A whole miRNA sequencing characterization. Biomedicine and Pharmacotherapy, 2022, 145, 112405.	5.6	3
4	Metabolomic changes after DAAs therapy are related to the improvement of cirrhosis and inflammation in HIV/HCV-coinfected patients. Biomedicine and Pharmacotherapy, 2022, 147, 112623.	5.6	6
5	Plasma miRNA profile at COVID-19 onset predicts severity status and mortality. Emerging Microbes and Infections, 2022, 11, 676-688.	6.5	44
6	Dynamics of HIV Reservoir and HIV-1 Viral Splicing in HCV-Exposed Individuals after Elimination with DAAs or Spontaneous Clearance. Journal of Clinical Medicine, 2022, 11, 3579.	2.4	2
7	OLFM4 polymorphisms predict septic shock survival after major surgery. European Journal of Clinical Investigation, 2021, 51, e13416.	3.4	3
8	TRPM5 rs886277 Polymorphism Predicts Hepatic Fibrosis Progression in Non-Cirrhotic HCV-Infected Patients. Journal of Clinical Medicine, 2021, 10, 483.	2.4	1
9	HCV eradication with IFN-based therapy does not completely restore gene expression in PBMCs from HIV/HCV-coinfected patients. Journal of Biomedical Science, 2021, 28, 23.	7.0	6
10	HCV Cure With Direct-Acting Antivirals Improves Liver and Immunological Markers in HIV/HCV-Coinfected Patients. Frontiers in Immunology, 2021, 12, 723196.	4.8	14
11	Age-Adjusted Endothelial Activation and Stress Index for Coronavirus Disease 2019 at Admission Is a Reliable Predictor for 28-Day Mortality in Hospitalized Patients With Coronavirus Disease 2019. Frontiers in Medicine, 2021, 8, 736028.	2.6	4
12	Are Reduced Levels of Coagulation Proteins Upon Admission Linked to COVID-19 Severity and Mortality?. Frontiers in Medicine, 2021, 8, 718053.	2.6	7
13	IL-1R1 rs6755229 polymorphism is related to death in patients undergoing major surgery who develop septic shock: a retrospective study. Infectious Diseases, 2021, , 1-4.	2.8	0
14	CEACAM7 polymorphisms predict genetic predisposition to mortality in post-surgical septic shock patients. Journal of Microbiology, Immunology and Infection, 2021, , .	3.1	0
15	Different HCV Exposure Drives Specific miRNA Profile in PBMCs of HIV Patients. Biomedicines, 2021, 9, 1627.	3.2	2
16	Near normalization of peripheral blood markers in HIV-infected patients on long-term suppressive antiretroviral therapy: a caseâ€“control study. Aids, 2020, 34, 1891-1897.	2.2	4
17	Hepatitis C Virus Influences HIV-1 Viral Splicing in Coinfected Patients. Journal of Clinical Medicine, 2020, 9, 2091.	2.4	3
18	Comparison of methods and characterization of small RNAs from plasma extracellular vesicles of HIV/HCV coinfected patients. Scientific Reports, 2020, 10, 11140.	3.3	22

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19	MTHFR rs1801133 Polymorphism Is Associated With Liver Fibrosis Progression in Chronic Hepatitis C: A Retrospective Study. Frontiers in Medicine, 2020, 7, 582666.	2.6	4
20	Telomere Length Increase in HIV/HCV-Coinfected Patients with Cirrhosis after HCV Eradication with Direct-Acting Antivirals. Journal of Clinical Medicine, 2020, 9, 2407.	2.4	5
21	IFNL3 rs12980275 Polymorphism Predicts Septic Shock-Related Death in Patients Undergoing Major Surgery: A Retrospective Study. Frontiers in Medicine, 2020, 7, 186.	2.6	1
22	Plasma metabolomic fingerprint of advanced cirrhosis stages among HIV/HCVâ€coinfected and HCVâ€monoinfected patients. Liver International, 2020, 40, 2215-2227.	3.9	11
23	Brief Report: CYP27B1 rs10877012 T Allele Was Linked to Non-AIDS Progression in ART-NaÃ¯ve HIV-Infected Patients: A Retrospective Study. Journal of Acquired Immune Deficiency Syndromes (1999), 2020, 85, 659-664.	2.1	2
24	DBP rs16846876 and rs12512631 polymorphisms are associated with progression to AIDS naÃ¯ve HIV-infected patients: a retrospective study. Journal of Biomedical Science, 2019, 26, 83.	7.0	2
25	MicroRNA Profile of HCV Spontaneous Clarified Individuals, Denotes Previous HCV Infection. Journal of Clinical Medicine, 2019, 8, 849.	2.4	11
26	VDR rs2228570 Polymorphism Is Related to Non-Progression to AIDS in Antiretroviral Therapy NaÃ¯ve HIV-Infected Patients. Journal of Clinical Medicine, 2019, 8, 311.	2.4	9
27	TNFAIP3, TNIP1, and MyD88 Polymorphisms Predict Septic-Shock-Related Death in Patients Who Underwent Major Surgery. Journal of Clinical Medicine, 2019, 8, 283.	2.4	5
28	Impact of DARC rs12075 Variants on Liver Fibrosis Progression in Patients with Chronic Hepatitis C: A Retrospective Study. Biomolecules, 2019, 9, 143.	4.0	7
29	HCV-coinfection is related to an increased HIV-1 reservoir size in cART-treated HIV patients: a cross-sectional study. Scientific Reports, 2019, 9, 5606.	3.3	22
30	Genetic variants upstream of TNFAIP3 in the 6q23 region are associated with liver disease severity in HIV/HCV-coinfected patients: A cross-sectional study. Infection, Genetics and Evolution, 2019, 67, 112-120.	2.3	2
31	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
32	Association of CD14 rs2569190 polymorphism with mortality in shock septic patients who underwent major cardiac or abdominal surgery: A retrospective study. Scientific Reports, 2018, 8, 2698.	3.3	7
33	The Myeloid-Epithelial-Reproductive Tyrosine Kinase (MERTK) rs4374383 Polymorphism Predicts Progression of Liver Fibrosis in Hepatitis C Virus-Infected Patients: A Longitudinal Study. Journal of Clinical Medicine, 2018, 7, 473.	2.4	15
34	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
35	Vitamin D in Human Immunodeficiency Virus Infection: Influence on Immunity and Disease. Frontiers in Immunology, 2018, 9, 458.	4.8	110
36	IL-6 rs1800795 polymorphism is associated with septic shock-related death in patients who underwent major surgery: a preliminary retrospective study. Annals of Intensive Care, 2017, 7, 22.	4.6	12

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37	ADAR1 polymorphisms are related to severity of liver fibrosis in HIV/HCV-coinfected patients. Scientific Reports, 2017, 7, 12918.	3.3	7
38	IL7RApolymorphisms are not associated with AIDS progression. European Journal of Clinical Investigation, 2017, 47, 719-727.	3.4	3
39	ILâ€1B rs16944 polymorphism is related to septic shock and death. European Journal of Clinical Investigation, 2017, 47, 53-62.	3.4	17
40	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
41	Genetic Polymorphisms Associated with Liver Disease Progression in HIV/HCV-Coinfected Patients. AIDS Reviews, 2017, 19, 3-15.	1.0	14
42	Relationship of TRIM5 and TRIM22 polymorphisms with liver disease and HCV clearance after antiviral therapy in HIV/HCV coinfected patients. Journal of Translational Medicine, 2016, 14, 257.	4.4	20
43	Short Communication: CXCL12 rs1029153 Polymorphism Is Associated with the Sustained Virological Response in HIV/Hepatitis C Virus-Coinfected Patients on Hepatitis C Virus Therapy. AIDS Research and Human Retroviruses, 2016, 32, 226-231.	1.1	0
44	Reply. Hepatology, 2015, 62, 1643-1643.	7.3	2
45	Single Nucleotide Polymorphisms of CXCL9-11 Chemokines Are Associated With Liver Fibrosis in HIV/HCV-Coinfected Patients. Journal of Acquired Immune Deficiency Syndromes (1999), 2015, 68, 386-395.	2.1	11
46	Toll-like receptor 8 (TLR8) polymorphisms are associated with non-progression of chronic hepatitis C in HIV/HCV coinfected patients. Infection, Genetics and Evolution, 2015, 36, 339-344.	2.3	6
47	Mitochondrial DNA haplogroups are associated with severe sepsis and mortality in patients who underwent major surgery. Journal of Infection, 2015, 70, 20-29.	3.3	17
48	rs7903146 Polymorphism atTranscription Factor 7 Like 2Gene Is Associated with Total Cholesterol and Lipoprotein Profile in HIV/Hepatitis C Virus-Coinfected Patients. AIDS Research and Human Retroviruses, 2015, 31, 326-334.	1.1	5
49	Relationship between European Mitochondrial Haplogroups and Chronic Renal Allograft Rejection in Patients with Kidney Transplant. International Journal of Medical Sciences, 2014, 11, 1129-1132.	2.5	3
50	Association of adiponectin (<scp>ADIPOQ</scp>) rs2241766 polymorphism and dyslipidemia in <scp>HIV</scp>/<scp>HCV</scp>â€coinfected patients. European Journal of Clinical Investigation, 2014, 44, 453-462.	3.4	12
51	PPARÎ³2 Pro12Ala Polymorphism Is Associated With Sustained Virological Response in HIV/HCV-Coinfected Patients Under HCV Therapy. Journal of Acquired Immune Deficiency Syndromes (1999), 2014, 67, 113-119.	2.1	5
52	SLC30A8 rs13266634 polymorphism is related to a favorable cardiometabolic lipid profile in HIV/hepatitis C virus-coinfected patients. Aids, 2014, 28, 1325-1332.	2.2	9
53	<scp>IL</scp>28<scp>RA</scp> polymorphism (rs10903035) is associated with insulin resistance in <scp>HIV</scp>/<scp>HCV</scp>â€coinfected patients. Journal of Viral Hepatitis, 2014, 21, 189-197.	2.0	5
54	Relationship of vitamin D status with advanced liver fibrosis and response to hepatitis C virus therapy: A meta-analysis. Hepatology, 2014, 60, 1541-1550.	7.3	68

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55	<scp>KIT</scp> and melanoma predisposition in pigs: sequence variants and association analysis. Animal Genetics, 2014, 45, 445-448.	1.7	8
56	PPARÎ³2 Pro12Ala polymorphism was associated with favorable cardiometabolic risk profile in HIV/HCV coinfected patients: a cross-sectional study. Journal of Translational Medicine, 2014, 12, 235.	4.4	11
57	European mitochondrial haplogroups are not associated with hepatitis <scp>C</scp> virus (<scp>HCV</scp>) treatment response in <scp>HIV</scp>/<scp>HCV</scp>â€coinfected patients. HIV Medicine, 2014, 15, 425-430.	2.2	5
58	Meta-analysis: implications of interleukin-28B polymorphisms in spontaneous and treatment-related clearance for patients with hepatitis C. BMC Medicine, 2013, 11, 6.	5.5	80
59	Comment on: â€˜Interleukin-28 polymorphisms on the SVR in the treatment of naÃ¨ve chronic hepatitis C with pegylated interferon-Î± plus ribavirin: A meta-analysisâ€™. Gene, 2013, 522, 121.	2.2	2
60	<scp>IL</scp>28<scp>RA</scp> polymorphism is associated with early hepatitis <scp>C</scp> virus (<scp>HCV</scp>) treatment failure in human immunodeficiency virusâ€/<scp>HCV</scp>â€coinfected patients. Journal of Viral Hepatitis, 2013, 20, 358-366.	2.0	17
61	IL28B polymorphisms are associated with severity ofÂliver disease in human immunodeficiency virus (HIV) patients coinfected with hepatitis C virus. Journal of Infection, 2013, 66, 170-178.	3.3	13
62	HLA-E variants are associated with sustained virological response in HIV/hepatitis C virus-coinfected patients on hepatitis C virus therapy. Aids, 2013, 27, 1231-1238.	2.2	15
63	Prediction of Hepatic Fibrosis in Patients Coinfected With HIV and Hepatitis C Virus Based on Genetic Markers. Journal of Acquired Immune Deficiency Syndromes (1999), 2013, 64, 434-442.	2.1	6
64	European mitochondrial haplogroups are associated with CD4+ T cell recovery in HIV-infected patients on combination antiretroviral therapy. Journal of Antimicrobial Chemotherapy, 2013, 68, 2349-2357.	3.0	17
65	Variability-specific differential gene expression across reproductive stages in sows. Animal, 2013, 7, 378-385.	3.3	1
66	Mitochondrial Haplogroups Are Associated With Clinical Pattern of AIDS Progression in HIV-Infected Patients. Journal of Acquired Immune Deficiency Syndromes (1999), 2013, 63, 178-183.	2.1	21
67	Analysis of IL28B alleles with virologic response patterns and plasma cytokine levels in HIV/HCV-coinfected patients. Aids, 2013, 27, 163-173.	2.2	12
68	Selection of Internal Control Genes for Real-Time Quantitative PCR in Ovary and Uterus of Sows across Pregnancy. PLoS ONE, 2013, 8, e66023.	2.5	26
69	Plasma IL-6 and IL-9 predict the failure of interferon-Â plus ribavirin therapy in HIV/HCV-coinfected patients. Journal of Antimicrobial Chemotherapy, 2012, 67, 1238-1245.	3.0	30
70	Bacterial DNA Translocation and Liver Disease Severity Among HIV-Infected Patients With Chronic Hepatitis C. Journal of Acquired Immune Deficiency Syndromes (1999), 2012, 61, 552-556.	2.1	11
71	High plasma CXCL10 levels are associated with HCV-genotype 1, and higher insulin resistance, fibrosis, and HIV viral load in HIV/HCV coinfected patients. Cytokine, 2012, 57, 25-29.	3.2	20
72	Genetic polymorphisms located in TGFB1, AGTR1, and VEGFA genes are associated to chronic renal allograft dysfunction. Cytokine, 2012, 58, 321-326.	3.2	17

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73	Genetic polymorphisms located in genes related to immune and inflammatory processes are associated with end-stage renal disease: a preliminary study. BMC Medical Genetics, 2012, 13, 58.	2.1	9
74	Sequencing and gene expression of the porcine ITIH SSC13 cluster and its effect on litter size in an IberianÃ—Meishan F2 population. Animal Reproduction Science, 2011, 128, 85-92.	1.5	4
75	Analysis of porcine MUC4 gene as a candidate gene for prolificacy QTL on SSC13 in an Iberian Ã— Meishan F2 population. BMC Genetics, 2011, 12, 93.	2.7	8
76	Differential Gene Expression in Ovaries of Pregnant Pigs with High and Low Prolificacy Levels and Identification of Candidate Genes for Litter Size. Biology of Reproduction, 2011, 84, 299-307.	2.7	31
77	Analysis of candidate genes underlying two epistatic quantitative trait loci on SSC12 affecting litter size in pig. Animal Genetics, 2010, 41, 73-80.	1.7	10
78	Identification of mitochondrial markers for genetic traceability of European wild boars and Iberian and Duroc pigs. Animal, 2009, 3, 1216-1223.	3.3	16
79	QTL detection on porcine chromosome 12 for fattyâ€acid composition and association analyses of thefatty acid synthase, gastric inhibitory polypeptideandacetylâ€coenzyme A carboxylase alphagenes. Animal Genetics, 2007, 38, 639-646.	1.7	40