Francisco Rodriguez-frias

List of Publications by Year in Descending Order

Source: https://exaly.com/author-pdf/1944542/francisco-rodriguez-frias-publications-by-year.pdf

Version: 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

108
papers2,292
citations30
h-index42
g-index123
ext. papers2,730
ext. citations5.5
avg, IF4.39
L-index

#	Paper	IF	Citations
108	Recovery of serum testosterone levels is an accurate predictor of survival from COVID-19 in male patients <i>BMC Medicine</i> , 2022 , 20, 129	11.4	1
107	Next-Generation Sequencing for Confronting Virus Pandemics Viruses, 2022, 14,	6.2	3
106	Influence of Type 2 Diabetes in the Association of PNPLA3 rs738409 and TM6SF2 rs58542926 Polymorphisms in NASH Advanced Liver Fibrosis. <i>Biomedicines</i> , 2022 , 10, 1015	4.8	1
105	Indirect determination of biochemistry reference intervals using outpatient data <i>PLoS ONE</i> , 2022 , 17, e0268522	3.7	1
104	Hepatitis B Virus Variants with Multiple Insertions and/or Deletions in the X Open Reading Frame 3? End: Common Members of Viral Quasispecies in Chronic Hepatitis B Patients. <i>Biomedicines</i> , 2022 , 10, 1194	4.8	1
103	Cross-sectional evaluation of circulating hepatitis B virus RNA and DNA: Different quasispecies?. <i>World Journal of Gastroenterology</i> , 2021 , 27, 7144-7158	5.6	0
102	Letter to the Editor: standardization of HBV RNA assay for the different phases of chronic hepatitis B is essential. <i>Hepatology</i> , 2021 ,	11.2	1
101	Resistance-associated substitutions after sofosbuvir/velpatasvir/voxilaprevir triple therapy failure. Journal of Viral Hepatitis, 2021 , 28, 1319-1324	3.4	2
100	Utility of Transient Elastography for the Screening of Liver Disease in Patients with Alpha1-Antitrypsin Deficiency. <i>Journal of Clinical Medicine</i> , 2021 , 10,	5.1	3
99	Next-generation sequencing for the diagnosis of hepatitis B: current status and future prospects. <i>Expert Review of Molecular Diagnostics</i> , 2021 , 21, 381-396	3.8	3
98	One-quarter of chronic hepatitis D patients reach HDV-RNA decline or undetectability during the natural course of the disease. <i>Alimentary Pharmacology and Therapeutics</i> , 2021 , 54, 462-469	6.1	4
97	Etiologies and Features of Acute Viral Hepatitis in Spain. <i>Clinical Gastroenterology and Hepatology</i> , 2021 , 19, 1030-1037	6.9	2
96	Viral and immune factors associated with successful treatment withdrawal in HBeAg-negative chronic hepatitis B patients. <i>Journal of Hepatology</i> , 2021 , 74, 1064-1074	13.4	15
95	Reflex viral load testing in dried blood spots generated by plasma separation card allows the screening and diagnosis of chronic viral hepatitis. <i>Journal of Virological Methods</i> , 2021 , 289, 114039	2.6	5
94	Host-dependent editing of SARS-CoV-2 in COVID-19 patients. <i>Emerging Microbes and Infections</i> , 2021 , 10, 1777-1789	18.9	4
93	Sophisticated viral quasispecies with a genotype-related pattern of mutations in the hepatitis B X gene of HBeAg-ve chronically infected patients. <i>Scientific Reports</i> , 2021 , 11, 4215	4.9	1
92	Utility of the Cobas Plasma Separation Card as a Sample Collection Device for Serological and Virological Diagnosis of Hepatitis C Virus Infection. <i>Diagnostics</i> , 2021 , 11,	3.8	1

(2018-2021)

91	A novel model of care for simplified testing of HBV in African communities during the COVID-19 pandemic in Spain. <i>Scientific Reports</i> , 2021 , 11, 17063	4.9	1
90	Trends in Diagnosis of Alpha-1 Antitrypsin Deficiency Between 2015 and 2019 in a Reference Laboratory. <i>International Journal of COPD</i> , 2020 , 15, 2421-2431	3	1
89	Conservation and variability of hepatitis B core at different chronic hepatitis stages. <i>World Journal of Gastroenterology</i> , 2020 , 26, 2584-2598	5.6	4
88	Deep-sequencing reveals broad subtype-specific HCV resistance mutations associated with treatment failure. <i>Antiviral Research</i> , 2020 , 174, 104694	10.8	20
87	Naturally occurring SARS-CoV-2 gene deletions close to the spike S1/S2 cleavage site in the viral quasispecies of COVID19 patients. <i>Emerging Microbes and Infections</i> , 2020 , 9, 1900-1911	18.9	36
86	Deep-sequencing study of HCV G4a resistance-associated substitutions in Egyptian patients failing DAA treatment. <i>Infection and Drug Resistance</i> , 2019 , 12, 2799-2807	4.2	5
85	Whole-genome characterization and resistance-associated substitutions in a new HCV genotype 1 subtype. <i>Infection and Drug Resistance</i> , 2019 , 12, 947-955	4.2	2
84	Characterization of hepatitis B virus X gene quasispecies complexity in mono-infection and hepatitis delta virus superinfection. <i>World Journal of Gastroenterology</i> , 2019 , 25, 1566-1579	5.6	3
83	Significant Improvement in Diagnosis of Hepatitis C Virus Infection by a One-Step Strategy in a Central Laboratory: an Optimal Tool for Hepatitis C Elimination?. <i>Journal of Clinical Microbiology</i> , 2019 , 58,	9.7	6
82	Viral Load Measurements in Individuals with Hepatitis C Virus Infection: on the European Association for the Study of the Liver Recommendations on Treatment of Hepatitis C 2018. <i>Journal of Clinical Microbiology</i> , 2019 , 58,	9.7	3
81	Amino Acid Substitutions within HLA-B*27-Restricted T Cell Epitopes Prevent Recognition by Hepatitis Delta Virus-Specific CD8 T Cells. <i>Journal of Virology</i> , 2018 , 92,	6.6	14
80	Quantitative characterization of hepatitis delta virus genome edition by next-generation sequencing. <i>Virus Research</i> , 2018 , 243, 52-59	6.4	8
79	Hepatitis B virus: The challenge of an ancient virus with multiple faces and a remarkable replication strategy. <i>Antiviral Research</i> , 2018 , 158, 34-44	10.8	21
78	Real world evaluation of a novel lateral flow assay (AlphaKit QuickScreen) for the detection of alpha-1-antitrypsin deficiency. <i>Respiratory Research</i> , 2018 , 19, 151	7.3	11
77	Analysis of hepatitis B virus preS1 variability and prevalence of the rs2296651 polymorphism in a Spanish population. <i>World Journal of Gastroenterology</i> , 2018 , 24, 680-692	5.6	8
76	New hepatitis C virus genotype 1 subtype naturally harbouring resistance-associated mutations to NS5A inhibitors. <i>Journal of General Virology</i> , 2018 , 99, 97-102	4.9	6
75	Characterization of Novel Missense Variants of SERPINA1 Gene Causing Alpha-1 Antitrypsin Deficiency. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2018 , 58, 706-716	5.7	16
74	Detection of hyper-conserved regions in hepatitis B virus X gene potentially useful for gene therapy. <i>World Journal of Gastroenterology</i> , 2018 , 24, 2095-2107	5.6	10

73	HDV evolution-will viral resistance be an issue in HDV infection?. <i>Current Opinion in Virology</i> , 2018 , 32, 100-107	7.5	1
72	Pipeline for specific subtype amplification and drug resistance detection in hepatitis C virus. <i>BMC Infectious Diseases</i> , 2018 , 18, 446	4	19
71	Baseline hepatitis C virus resistance-associated substitutions present at frequencies lower than 15% may be clinically significant. <i>Infection and Drug Resistance</i> , 2018 , 11, 2207-2210	4.2	20
70	Spanish Registry of Patients With Alpha-1 Antitrypsin Deficiency: Database Evaluation and Population Analysis. <i>Archivos De Bronconeumologia</i> , 2017 , 53, 13-18	0.7	20
69	Spanish Registry of Patients With Alpha-1 Antitrypsin Deficiency: Database Evaluation and Population Analysis. <i>Archivos De Bronconeumologia</i> , 2017 , 53, 13-18	0.7	2
68	Selection of the highly replicative and partially multidrug resistant rtS78T HBV polymerase mutation during TDF-ETV combination therapy. <i>Journal of Hepatology</i> , 2017 , 67, 246-254	13.4	41
67	A new HDV mouse model identifies mitochondrial antiviral signaling protein (MAVS) as a key player in IFN-Induction. <i>Journal of Hepatology</i> , 2017 , 67, 669-679	13.4	31
66	Genetic diagnosis of 🛘 -antitrypsin deficiency using DNA from buccal swab and serum samples. <i>Clinical Chemistry and Laboratory Medicine</i> , 2017 , 55, 1276-1283	5.9	5
65	Deep sequencing in the management of hepatitis virus infections. <i>Virus Research</i> , 2017 , 239, 115-125	6.4	14
64	Assessment of a Novel Automatic Real-Time PCR Assay on the Cobas 4800 Analyzer as a Screening Platform for Hepatitis C Virus Genotyping in Clinical Practice: Comparison with Massive Sequencing. <i>Journal of Clinical Microbiology</i> , 2017 , 55, 504-509	9.7	15
63	Gene and miRNA expression profiles in PBMCs from patients with severe and mild emphysema and PiZZ alpha1-antitrypsin deficiency. <i>International Journal of COPD</i> , 2017 , 12, 3381-3390	3	19
62	Laboratory Diagnosis by Genotyping. <i>Methods in Molecular Biology</i> , 2017 , 1639, 45-60	1.4	
61	Red blood cell transfusion-transmitted acute hepatitis E in an immunocompetent subject in Europe: a case report. <i>Transfusion</i> , 2017 , 57, 244-247	2.9	30
60	Sofosbuvir and daclatasvir in mono- and HIV-coinfected patients with recurrent hepatitis C after liver transplant. <i>Annals of Hepatology</i> , 2017 , 16, 86-93	3.1	18
59	Rapid detection of Mmalton II-antitrypsin deficiency allele by real-time PCR and melting curves in whole blood, serum and dried blood spot samples. <i>Clinical Chemistry and Laboratory Medicine</i> , 2016 , 54, 241-8	5.9	13
58	Evidence of an Exponential Decay Pattern of the Hepatitis Delta Virus Evolution Rate and Fluctuations in Quasispecies Complexity in Long-Term Studies of Chronic Delta Infection. <i>PLoS ONE</i> , 2016 , 11, e0158557	3.7	14
57	New real-time-PCR method to identify single point mutations in hepatitis C virus. <i>World Journal of Gastroenterology</i> , 2016 , 22, 9604-9612	5.6	5
56	Application of a diagnostic algorithm for the rare deficient variant Mmalton of alpha-1-antitrypsin deficiency: a new approach. <i>International Journal of COPD</i> , 2016 , 11, 2535-2541	3	9

55	Viral quasispecies complexity measures. Virology, 2016, 493, 227-37	3.6	59
54	Tenofovir discontinuation after long-term viral suppression in HBeAg negative chronic hepatitis B. Can HBsAg levels be useful?. <i>Journal of Clinical Virology</i> , 2015 , 68, 61-8	14.5	13
53	Phylogenetic demonstration of hepatitis E infection transmitted by pork meat ingestion. <i>Journal of Clinical Gastroenterology</i> , 2015 , 49, 165-8	3	68
52	Immunogenicity of hepatitis B vaccine in patients with inflammatory bowel disease and the benefits of revaccination. <i>Journal of Gastroenterology and Hepatology (Australia)</i> , 2015 , 30, 92-8	4	22
51	Indications for active case searches and intravenous alpha-1 antitrypsin treatment for patients with alpha-1 antitrypsin deficiency chronic pulmonary obstructive disease: an update. <i>Archivos De Bronconeumologia</i> , 2015 , 51, 185-92	0.7	48
50	High-resolution hepatitis C virus subtyping using NS5B deep sequencing and phylogeny, an alternative to current methods. <i>Journal of Clinical Microbiology</i> , 2015 , 53, 219-26	9.7	64
49	Complex Genotype Mixtures Analyzed by Deep Sequencing in Two Different Regions of Hepatitis B Virus. <i>PLoS ONE</i> , 2015 , 10, e0144816	3.7	8
48	Indications for Active Case Searches and Intravenous Alpha-1 Antitrypsin Treatment for Patients With Alpha-1 Antitrypsin Deficiency Chronic Pulmonary Obstructive Disease: An Update. <i>Archivos De Bronconeumologia</i> , 2015 , 51, 185-192	0.7	3
47	Quasispecies dynamics in hepatitis C liver transplant recipients receiving grafts from hepatitis C virus infected donors. <i>Journal of General Virology</i> , 2015 , 96, 3493-3498	4.9	8
46	Inference with viral quasispecies diversity indices: clonal and NGS approaches. <i>Bioinformatics</i> , 2014 , 30, 1104-1111	7.2	46
45	Clinical application of estimating hepatitis B virus quasispecies complexity by massive sequencing: correlation between natural evolution and on-treatment evolution. <i>PLoS ONE</i> , 2014 , 9, e112306	3.7	24
44	Relevance of a full-length genomic RNA standard and a thermal-shock step for optimal hepatitis delta virus quantification. <i>Journal of Clinical Microbiology</i> , 2014 , 52, 3334-8	9.7	11
43	Identification of host and viral factors involved in a dissimilar resolution of a hepatitis C virus infection. <i>Liver International</i> , 2014 , 34, 896-906	7.9	9
42	Severe alpha-1 antitrypsin deficiency in composite heterozygotes inheriting a new splicing mutation QOMadrid. <i>Respiratory Research</i> , 2014 , 15, 125	7:3	31
41	Cirrhosis, liver transplantation and HIV infection are risk factors associated with hepatitis E virus infection. <i>PLoS ONE</i> , 2014 , 9, e103028	3.7	33
40	A comparative study of ultra-deep pyrosequencing and cloning to quantitatively analyze the viral quasispecies using hepatitis B virus infection as a model. <i>Antiviral Research</i> , 2013 , 98, 273-83	10.8	45
39	Ultra-deep pyrosequencing (UDPS) data treatment to study amplicon HCV minor variants. <i>PLoS ONE</i> , 2013 , 8, e83361	3.7	49
38	Molecular epidemiology and putative origin of hepatitis C virus in random volunteers from Argentina. <i>World Journal of Gastroenterology</i> , 2013 , 19, 5813-27	5.6	15

37	Quasispecies structure, cornerstone of hepatitis B virus infection: mass sequencing approach. <i>World Journal of Gastroenterology</i> , 2013 , 19, 6995-7023	5.6	31
36	Rare alpha-1-antitrypsin variants: are they really so rare?. <i>Therapeutic Advances in Respiratory Disease</i> , 2012 , 6, 79-85	4.9	48
35	Ultra-deep pyrosequencing detects conserved genomic sites and quantifies linkage of drug-resistant amino acid changes in the hepatitis B virus genome. <i>PLoS ONE</i> , 2012 , 7, e37874	3.7	48
34	II-Antitrypsin in fibromyalgia: results of a randomized, placebo-controlled, double-blind and crossover pilot trial. <i>Musculoskeletal Care</i> , 2012 , 10, 178-83	1.6	4
33	Twelve-week posttreatment follow-up to predict sustained virologic response for recurrent hepatitis C infection in liver recipients. <i>Transplantation</i> , 2012 , 93, 450-3	1.8	5
32	Quasispecies dynamics in main core epitopes of hepatitis B virus by ultra-deep-pyrosequencing. World Journal of Gastroenterology, 2012 , 18, 6096-105	5.6	14
31	Ultra-deep pyrosequencing analysis of the hepatitis B virus preCore region and main catalytic motif of the viral polymerase in the same viral genome. <i>Nucleic Acids Research</i> , 2011 , 39, 8457-71	20.1	40
30	HBV core region variability: effect of antiviral treatments on main epitopic regions. <i>Antiviral Therapy</i> , 2011 , 16, 37-49	1.6	11
29	Infrequent detection of hepatitis E virus RNA in pregnant women with hepatitis E virus antibodies in Spain. <i>Liver International</i> , 2010 , 30, 1549-51	7.9	3
28	Quantitative longitudinal evaluations of hepatitis delta virus RNA and hepatitis B virus DNA shows a dynamic, complex replicative profile in chronic hepatitis B and D. <i>Journal of Hepatology</i> , 2010 , 52, 658	-64·4	118
27	Response-guided therapy for chronic hepatitis C virus infection in patients coinfected with HIV: a pilot trial. <i>Clinical Infectious Diseases</i> , 2009 , 48, 1152-9	11.6	19
26	Use of the novel INNO-LiPA line probe assay for detection of hepatitis B virus variants that confer resistance to entecavir therapy. <i>Journal of Clinical Microbiology</i> , 2009 , 47, 485-8	9.7	9
25	Successful use of entecavir for a severe case of reactivation of hepatitis B virus following polychemotherapy containing rituximab. <i>Journal of Hepatology</i> , 2009 , 51, 1091-6	13.4	32
24	The dynamic DNA methylomes of double-stranded DNA viruses associated with human cancer. <i>Genome Research</i> , 2009 , 19, 438-51	9.7	201
23	Analysis of hepatitis B genotype changes in chronic hepatitis B infection: Influence of antiviral therapy. <i>Journal of Hepatology</i> , 2008 , 49, 695-701	13.4	33
22	Redetection of HBV lamivudine-resistant mutations in a patient under entecavir therapy, who had been treated sequentially with nucleos(t)ide analogues. <i>Journal of Medical Virology</i> , 2007 , 79, 1671-3	19.7	3
21	Community-based seroepidemiological survey of hepatitis E virus infection in Catalonia, Spain. <i>Vaccine Journal</i> , 2006 , 13, 1328-32		62
20	Hepatitis B virus genome variability and disease progression: the impact of pre-core mutants and HBV genotypes. <i>Journal of Clinical Virology</i> , 2005 , 34 Suppl 1, S79-82	14.5	38

(1991-2004)

19	Usefulness of dried blood samples for quantification and molecular characterization of HBV-DNA. <i>Hepatology</i> , 2004 , 40, 133-9	11.2	37
18	Hepatitis C virus Core Antigen as a predictor of non-response in genotype 1 chronic hepatitis C patients treated with peginterferon alpha-2b plus ribavirin. <i>Journal of Hepatology</i> , 2004 , 40, 527-32	13.4	22
17	Sporadic cases of acute autochthonous hepatitis E in Spain. <i>Journal of Hepatology</i> , 2004 , 41, 126-31	13.4	68
16	Emphysema due to alpha-antitrypsin deficiency: familial study of the YBARCELONA variant. <i>Chest</i> , 2003 , 124, 404-6	5.3	15
15	Applicability of real-time PCR combined with melting curve analysis for the study of hepatitis B virus genome mutations. <i>Hepatology</i> , 2003 , 37, 478-9; author reply 479-80	11.2	1
14	Is hepatitis B virus subtype testing useful in predicting virological response and resistance to lamivudine?. <i>Journal of Hepatology</i> , 2002 , 36, 445-6	13.4	55
13	Characterization of the new alpha-1-antitrypsin-deficient PI M-type allele, PI M(vall dhebron) (Pro(369)>Ser). <i>Human Heredity</i> , 2000 , 50, 320-1	1.1	11
12	Rapid screening of the Cys282Tyr and His63Asp mutations in the HFE gene. <i>Journal of Hepatology</i> , 2000 , 32, 362-3	13.4	3
11	Genetic alterations in the S gene of hepatitis B virus in patients with acute hepatitis B, chronic hepatitis B and hepatitis B liver cirrhosis before and after liver transplantation. <i>Liver International</i> , 1999 , 19, 177-82	7.9	39
10	Rapid detection of lamivudine-resistant hepatitis B virus polymerase gene variants. <i>Journal of Virological Methods</i> , 1999 , 83, 181-7	2.6	43
9	Transient emergence of hepatitis B variants in a patient with chronic hepatitis B resistant to lamivudine. <i>Journal of Hepatology</i> , 1998 , 28, 510-3	13.4	31
8	The value of quantitative detection of HBV-DNA amplified by PCR in the study of hepatitis B infection. <i>Journal of Hepatology</i> , 1996 , 24, 680-5	13.4	38
7	Hepatitis E virus infection in acute hepatitis in Spain. Journal of Virological Methods, 1995, 55, 49-54	2.6	31
6	Etiology of acute sporadic hepatitis in Spain: the role of hepatitis C and E viruses. <i>Journal of Hepatology</i> , 1994 , 20, 589-92	13.4	21
5	Detection of hepatitis B precore mutants by the fluorescent linear polymerase chain reaction sequencing method. <i>Journal of Hepatology</i> , 1994 , 21, 934-9	13.4	6
4	Treatment of chronic type D hepatitis and concomitant human immunodeficiency infection with alpha-interferon. <i>Journal of Hepatology</i> , 1992 , 14, 412-3	13.4	7
3	Detection of hepatitis B virus DNA in human serum samples: use of digoxigenin-labeled oligonucleotides as modified primers for the polymerase chain reaction. <i>Analytical Biochemistry</i> , 1992 , 206, 36-42	3.1	17
2	Clinical significance of two forms of IgM antibody to hepatitis delta virus. <i>Hepatology</i> , 1991 , 14, 25-8	11.2	13

Chronic delta hepatitis: detection of hepatitis delta virus antigen in serum by immunoblot and correlation with other markers of delta viral replication. *Hepatology*, **1989**, 10, 907-10

11.2 26