

Ariadna Rando-Segura

List of Publications by Year in descending order

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Version: 2024-02-01

41
papers

485
citations

759190

12
h-index

794568

19
g-index

41
all docs

41
docs citations

41
times ranked

738
citing authors

#	ARTICLE	IF	CITATIONS
1	Letter to the editor: Standardization of HBVâ€RNA assay for the different phases of chronic hepatitis B is essential. <i>Hepatology</i> , 2022, 75, 501-502.	7.3	2
2	A year living with SARS-CoV-2: an epidemiological overview of viral lineage circulation by whole-genome sequencing in Barcelona city (Catalonia, Spain). <i>Emerging Microbes and Infections</i> , 2022, 11, 172-181.	6.5	17
3	Inspecting the Ribozyme Region of Hepatitis Delta Virus Genotype 1: Conservation and Variability. <i>Viruses</i> , 2022, 14, 215.	3.3	4
4	Standardized Hepatitis B Virus RNA Quantification in Untreated and Treated Chronic Patients: a Promising Marker of Infection Follow-Up. <i>Microbiology Spectrum</i> , 2022, 10, e0214921.	3.0	2
5	ACE Score Identifies HBeAg-negative Inactive Carriers at a Single-point Evaluation, Regardless of HBV Genotype. <i>Journal of Clinical and Translational Hepatology</i> , 2022, 000, 000-000.	1.4	1
6	Costâ€effectiveness analysis of an active search to retrieve <scp>HCV</scp> patients lost to followâ€up (<scp>RELINKâ€C</scp> strategy) and the impact of <scp>COVID</scp>â€19. <i>Journal of Viral Hepatitis</i> , 2022, 29, 579-583.	2.0	7
7	Hepatitis B Virus Variants with Multiple Insertions and/or Deletions in the X Open Reading Frame 3â€2 End: Common Members of Viral Quasispecies in Chronic Hepatitis B Patients. <i>Biomedicines</i> , 2022, 10, 1194.	3.2	1
8	Characteristics of 24 SARS-CoV-2-Sequenced Reinfection Cases in a Tertiary Hospital in Spain. <i>Frontiers in Microbiology</i> , 2022, 13, .	3.5	1
9	Detailed stratified GWAS analysis for severe COVID-19 in four European populations. <i>Human Molecular Genetics</i> , 2022, 31, 3945-3966.	2.9	46
10	Etiologies and Features of Acute Viral Hepatitis in Spain. <i>Clinical Gastroenterology and Hepatology</i> , 2021, 19, 1030-1037.	4.4	12
11	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.	2.1	17
12	Host-dependent editing of SARS-CoV-2 in COVID-19 patients. <i>Emerging Microbes and Infections</i> , 2021, 10, 1777-1789.	6.5	13
13	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.	3.3	3
14	Utility of the CobasÂ® Plasma Separation Card as a Sample Collection Device for Serological and Virological Diagnosis of Hepatitis C Virus Infection. <i>Diagnostics</i> , 2021, 11, 473.	2.6	7
15	Resistanceâ€associated substitutions after sofosbuvir/velpatasvir/voxilaprevir triple therapy failure. <i>Journal of Viral Hepatitis</i> , 2021, 28, 1319-1324.	2.0	7
16	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.	3.1	6
17	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.	3.3	11
18	Barriers to linkage to care in hepatitis C patients with substance use disorders and dual diagnoses, despite centralized management. <i>Therapeutic Advances in Gastroenterology</i> , 2021, 14, 175628482110165.	3.2	0

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19	Molecular characterization of rpoB gene mutations in isolates from tuberculosis patients in Cubal, Republic of Angola. <i>BMC Infectious Diseases</i> , 2021, 21, 1056.	2.9	9
20	Study of Quasispecies Complexity and Liver Damage Progression after Liver Transplantation in Hepatitis C Virus Infected Patients. <i>Genes</i> , 2021, 12, 1731.	2.4	0
21	Cross-sectional evaluation of circulating hepatitis B virus RNA and DNA: Different quasispecies?. <i>World Journal of Gastroenterology</i> , 2021, 27, 7144-7158.	3.3	1
22	Assessing Liver Fibrosis Using the FIB4 Index in the Community Setting. <i>Diagnostics</i> , 2021, 11, 2236.	2.6	16
23	Microorganisms as Shapers of Human Civilization, from Pandemics to Even Our Genomes: Villains or Friends? A Historical Approach. <i>Microorganisms</i> , 2021, 9, 2518.	3.6	6
24	Prevalence of <i>Mycoplasma genitalium</i> and macrolide resistance among asymptomatic people visiting a point of care service for rapid STI screening: a cross-sectional study. <i>Sexually Transmitted Infections</i> , 2020, 96, 300-305.	1.9	19
25	Deep-sequencing reveals broad subtype-specific HCV resistance mutations associated with treatment failure. <i>Antiviral Research</i> , 2020, 174, 104694.	4.1	39
26	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.	6.5	57
27	Unexpected long-lasting anti-HEV IgM positivity: Is HEV antigen a better serological marker for hepatitis E infection diagnosis?. <i>Journal of Viral Hepatitis</i> , 2020, 27, 747-753.	2.0	26
28	Conservation and variability of hepatitis B core at different chronic hepatitis stages. <i>World Journal of Gastroenterology</i> , 2020, 26, 2584-2598.	3.3	8
29	The challenge of the laboratory diagnosis in a confirmed congenital Zika virus syndrome in utero. <i>Medicine (United States)</i> , 2019, 98, e15532.	1.0	9
30	<p>Deep-sequencing study of HCV G4a resistance-associated substitutions in Egyptian patients failing DAA treatment</p>. <i>Infection and Drug Resistance</i> , 2019, Volume 12, 2799-2807.	2.7	5
31	Screening for Zika virus infection in 1057 potentially exposed pregnant women, Catalonia (northeastern Spain). <i>Travel Medicine and Infectious Disease</i> , 2019, 29, 69-71.	3.0	7
32	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, .	3.9	21
33	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, .	3.9	6
34	Retrospective Study of Cystic Echinococcosis in a Recent Cohort of a Referral Center for Liver Surgery. <i>Journal of Gastrointestinal Surgery</i> , 2019, 23, 1148-1156.	1.7	19
35	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.	3.3	7
36	Treatment Outcomes and Adverse Events from a Standardized Multidrug-Resistant Tuberculosis Regimen in a Rural Setting in Angola. <i>American Journal of Tropical Medicine and Hygiene</i> , 2019, 101, 502-509.	1.4	8

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37	Evaluation of the novel DiaSorin LIAISON Â® Campylobacter assay for the rapid detection of Campylobacter spp.. Enfermedades Infecciosas Y MicrobiologÃa ClÃnica, 2018, 36, 293-295.	0.5	3
38	Zika virus dynamics in body fluids and risk of sexual transmission in a nonâ€ndemic area. Tropical Medicine and International Health, 2018, 23, 92-100.	2.3	26
39	Detection of hyper-conserved regions in hepatitis B virus X gene potentially useful for gene therapy. World Journal of Gastroenterology, 2018, 24, 2095-2107.	3.3	17
40	Evaluation of Seegene Allplex Respiratory Panel 1 kit for the detection of influenza virus and human respiratory syncytial virus. Journal of Clinical Virology, 2018, 105, 31-34.	3.1	13
41	Drug Resistance of <i>Mycobacterium tuberculosis</i> Complex in a Rural Setting, Angola. Emerging Infectious Diseases, 2018, 24, 569-572.	4.3	6