

Carmen F Manso

List of Publications by Year in descending order

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Version: 2024-02-01

20
papers

293
citations

933447

10
h-index

940533

16
g-index

21
all docs

21
docs citations

21
times ranked

558
citing authors

#	ARTICLE	IF	CITATIONS
1	Real world SOF/VEL/VOX retreatment outcomes and viral resistance analysis for HCV patients with prior failure to DAA therapy. <i>Journal of Viral Hepatitis</i> , 2021, 28, 1256-1264.	2.0	16
2	Clinical evaluation of a Hepatitis C Virus whole-genome sequencing pipeline for genotyping and resistance testing. <i>Clinical Microbiology and Infection</i> , 2021, , .	6.0	1
3	Technical Validation of a Hepatitis C Virus Whole Genome Sequencing Assay for Detection of Genotype and Antiviral Resistance in the Clinical Pathway. <i>Frontiers in Microbiology</i> , 2020, 11, 576572.	3.5	13
4	Surveillance of HIV-1 transmitted integrase strand transfer inhibitor resistance in the UK. <i>Journal of Antimicrobial Chemotherapy</i> , 2020, 75, 3311-3318.	3.0	15
5	Enhanced Detection of DNA Viruses in the Cerebrospinal Fluid of Encephalitis Patients Using Metagenomic Next-Generation Sequencing. <i>Frontiers in Microbiology</i> , 2020, 11, 1879.	3.5	10
6	Real-world retreatment of HCV-infected patients with prior failure to direct acting antiviral therapy using sofosbuvir, velpatasvir and voxilaprevir. <i>Journal of Hepatology</i> , 2020, 73, S336.	3.7	3
7	LBP-34-Effect of Resistance Associated Substitutions on Retreatment of HCV infected patients with prior failure to Direct Acting Antiviral Therapy. <i>Journal of Hepatology</i> , 2019, 70, e158-e159.	3.7	0
8	Interpreting Viral Deep Sequencing Data with GLUE. <i>Viruses</i> , 2019, 11, 323.	3.3	29
9	Clinical evaluation of herPHEgen [®] : a herpesvirus genotype-to-phenotype antiviral resistance database and testing service. <i>Access Microbiology</i> , 2019, 1, .	0.5	0
10	The Effect of Nucleic Acid Extraction Platforms and Sample Storage on the Integrity of Viral RNA for Use in Whole Genome Sequencing. <i>Journal of Molecular Diagnostics</i> , 2017, 19, 303-312.	2.8	15
11	Efficient and unbiased metagenomic recovery of RNA virus genomes from human plasma samples. <i>Scientific Reports</i> , 2017, 7, 4173.	3.3	31
12	Molecular epidemiology of norovirus from patients with acute gastroenteritis in northwestern Spain. <i>Epidemiology and Infection</i> , 2015, 143, 316-324.	2.1	10
13	Efficiency of hepatitis A virus removal in six sewage treatment plants from central Tunisia. <i>Applied Microbiology and Biotechnology</i> , 2015, 99, 10759-10769.	3.6	20
14	Bioaccumulation and Removal Dynamics of Murine Norovirus in Manila Clams (<i>Venerupis</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 222 Td		
15	Detection and Characterization of Hepatitis A Virus and Norovirus in Mussels from Galicia (NW) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 222 Td		
16	Role of norovirus in acute gastroenteritis in the Northwest of Spain during 2010â€“2011. <i>Journal of Medical Virology</i> , 2013, 85, 2009-2015.	5.0	10
17	Norovirus, hepatitis A virus and enterovirus presence in shellfish from high quality harvesting areas in Portugal. <i>Food Microbiology</i> , 2011, 28, 936-941.	4.2	48
18	Genotyping of hepatitis A virus detected in bivalve shellfish in Galicia (NW Spain). <i>Water Science and Technology</i> , 2010, 61, 15-24.	2.5	16

#	ARTICLE	IF	CITATIONS
19	Imported Mollusks and Dissemination of Human Enteric Viruses. Emerging Infectious Diseases, 2010, 16, 1036-1038.	4.3	25
20	Detection and quantification of Hepatitis A and E virus in wastewater from Tunisia. Frontiers in Marine Science, 0, 1, .	2.5	0