

Marina Selleri

List of Publications by Year in descending order

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Version: 2024-02-01

27
papers

519
citations

567281

15
h-index

677142

22
g-index

27
all docs

27
docs citations

27
times ranked

1183
citing authors

#	ARTICLE	IF	CITATIONS
1	Role of HIV-1 matrix protein p17 variants in lymphoma pathogenesis. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 14331-14336.	7.1	58
2	Duration of viral shedding in hospitalized patients infected with pandemic H1N1. BMC Infectious Diseases, 2011, 11, 140.	2.9	47
3	Recombinant interferon- β improves immune response to hepatitis B vaccination in haemodialysis patients: Results of a randomised clinical trial. Vaccine, 2009, 27, 5654-5660.	3.8	39
4	Effective Highly Active Antiretroviral Therapy in Patients With Primary HIV-1 Infection Prevents the Evolution of the Avidity of HIV-1-Specific Antibodies. Journal of Acquired Immune Deficiency Syndromes (1999), 2007, 46, 145-150.	2.1	32
5	Frequency of Detection of Upper Respiratory Tract Viruses in Patients Tested for Pandemic H1N1/09 Viral Infection. Journal of Clinical Microbiology, 2010, 48, 3383-3385.	3.9	30
6	Detection of haemagglutinin D222 polymorphisms in influenza A(H1N1)pdm09-infected patients by ultra-deep pyrosequencing. Clinical Microbiology and Infection, 2013, 19, 668-673.	6.0	29
7	Frequency of detection of respiratory viruses in the lower respiratory tract of hospitalized adults. Journal of Clinical Virology, 2008, 42, 215-220.	3.1	27
8	Fatal Outbreak in Tonkean Macaques Caused by Possibly Novel Orthopoxvirus, Italy, January 2015. Emerging Infectious Diseases, 2017, 23, 1941-1949.	4.3	27
9	Characterization of the patterns of drug-resistance mutations in newly diagnosed HIV-1 infected patients naïve to the antiretroviral drugs. BMC Infectious Diseases, 2009, 9, 111.	2.9	23
10	Possible Compartmentalization of Hepatitis C Viral Replication in the Genital Tract of HIV-1 Coinfected Women. Journal of Infectious Diseases, 2006, 194, 1529-1536.	4.0	21
11	Quasispecies tropism and compartmentalization in gut and peripheral blood during early and chronic phases of HIV-1 infection: possible correlation with immune activation markers. Clinical Microbiology and Infection, 2014, 20, O157-O166.	6.0	20
12	Evaluation of the effects of human leukocyte IFN- γ on the immune response to the HBV vaccine in healthy unvaccinated individuals. Vaccine, 2008, 26, 1038-1049.	3.8	19
13	Extent of HCV NS3 protease variability and resistance-associated mutations assessed by next generation sequencing in HCV monoinfected and HIV/HCV coinfecting patients. Virus Research, 2013, 177, 205-208.	2.2	18
14	Evolutionary pattern of pandemic influenza (H1N1) 2009 virus in the late phases of the 2009 pandemic. PLOS Currents, 2010, 2, RRN1149.	1.4	18
15	Whole Genome Characterization of Orthopoxvirus (OPV) Abatino, a Zoonotic Virus Representing a Putative Novel Clade of Old World Orthopoxviruses. Viruses, 2018, 10, 546.	3.3	17
16	Dynamics and phylogenetic relationships of HIV-1 transmitted drug resistance according to subtype in Italy over the years 2000-2014. Journal of Antimicrobial Chemotherapy, 2017, 72, 2837-2845.	3.0	15
17	Design and clinical application of a molecular method for detection and typing of the influenza A/H1N1pdm virus. Journal of Virological Methods, 2010, 163, 486-488.	2.1	11
18	Markers of human papillomavirus infection and their correlation with cervical dysplasia in human immunodeficiency virus-positive women. Clinical Microbiology and Infection, 2007, 13, 94-97.	6.0	10

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19	Improved detection of human influenza A and B viruses in respiratory tract specimens by hemi-nested PCR. <i>Journal of Virological Methods</i> , 2007, 141, 225-228.	2.1	8
20	Phylogenetic analysis of human coronavirus NL63 circulating in Italy. <i>Journal of Clinical Virology</i> , 2008, 43, 114-119.	3.1	8
21	Hemagglutinin 222 Variants in Pandemic (H1N1) 2009 Virus. <i>Emerging Infectious Diseases</i> , 2011, 17, 749-751.	4.3	8
22	Detection of HIV-1 Matrix Protein p17 Quasispecies Variants in Plasma of Chronic HIV-1 Infected Patients by Ultra-Deep Pyrosequencing. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2014, 66, 332-339.	2.1	8
23	Clinical and virological properties of hepatitis C virus genotype 4 infection in patients treated with different direct-acting antiviral agents. <i>Infection and Drug Resistance</i> , 2018, Volume 11, 2117-2127.	2.7	8
24	HCV NS3 quasispecies in liver and plasma and dynamics of telaprevir-resistant variants in breakthrough patients assessed by UDPS: A case study. <i>Journal of Clinical Virology</i> , 2015, 72, 60-65.	3.1	7
25	Near full length hepatitis C virus genome reconstruction by next generation sequencing based on genotype-independent amplification. <i>Digestive and Liver Disease</i> , 2015, 47, 608-612.	0.9	5
26	Comparison of two real-time RT-PCR-based systems for the detection and typing of the pandemic influenza A virus, 2009. <i>Journal of Virological Methods</i> , 2011, 177, 193-196.	2.1	4
27	In-Depth Analysis of HA and NS1 Genes in A(H1N1)pdm09 Infected Patients. <i>PLoS ONE</i> , 2016, 11, e0155661.	2.5	2