Anna Rita Ciccaglione

List of Publications by Year in descending order

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102 papers 2,814 citations

218592 26 h-index 197736 49 g-index

105 all docs 105 docs citations

105 times ranked 3586 citing authors

#	Article	IF	Citations
1	Prevalence and risk factors for hepatitis E virus infection in blood donors: a nationwide survey in Italy, 2017 to 2019. Eurosurveillance, 2022, 27, .	3.9	7
2	An Evaluation of Hepatitis E Virus Molecular Typing Methods. Clinical Chemistry, 2021, 68, 181-191.	1.5	5
3	Time and Mode of Epidemic HCV-2 Subtypes Spreading in Europe: Phylodynamics in Italy and Albania. Diagnostics, 2021, 11, 327.	1.3	2
4	Phylogenetic and Molecular Analyses of More Prevalent HCV1b Subtype in the Calabria Region, Southern Italy. Journal of Clinical Medicine, 2021, 10, 1655.	1.0	3
5	Developing and Piloting a Standardized European Protocol for Hepatitis C Prevalence Surveys in the General Population (2016–2019). Frontiers in Public Health, 2021, 9, 568524.	1.3	1
6	Hepatitis E Outbreak in the Central Part of Italy Sustained by Multiple HEV Genotype 3 Strains, June–December 2019. Viruses, 2021, 13, 1159.	1.5	14
7	Immunogenicity of Viral Vaccines in the Italian Military. Biomedicines, 2021, 9, 87.	1.4	5
8	Hepatitis A Virus Strains Circulating in the Campania Region (2015–2018) Assessed through Bivalve Biomonitoring and Environmental Surveillance. Viruses, 2021, 13, 16.	1.5	14
9	Phylogenetic analysis and epidemiological history of Hepatitis E virus 3f and 3c in swine and wild boar, Italy. Heliyon, 2020, 6, e05110.	1.4	10
10	Nine-Year Nationwide Environmental Surveillance of Hepatitis E Virus in Urban Wastewaters in Italy (2011–2019). International Journal of Environmental Research and Public Health, 2020, 17, 2059.	1.2	27
11	Sensitivity of hepatitis C virus rapid tests in detecting antibodies in general population. Panminerva Medica, 2020, 62, 125-130.	0.2	1
12	Retrospective analysis of acute HBV infections occurred in 1978–79 and 1994–95 in North-East Italy: increasing prevalence of BCP/pre-core mutants in sub-genotype D3. BMC Infectious Diseases, 2020, 20, 78.	1.3	3
13	Changing epidemiology of acute liver failure in Italy: a single-center experience over 25 years. Minerva Medica, 2020, 111, 330-336.	0.3	1
14	Incidence of hepatitis E virus infection among blood donors in a high endemic area of Central Italy. Journal of Viral Hepatitis, 2019, 26, 506-512.	1.0	22
15	Standardising surveillance of hepatitis E virus infection in the EU/EEA: A review of national practices and suggestions for the way forward. Journal of Clinical Virology, 2019, 120, 63-67.	1.6	14
16	Human hepatitis E virus circulation in Bulgaria: Deep Bayesian phylogenetic analysis for viral spread control in the country. Journal of Medical Virology, 2019, 91, 132-138.	2.5	8
17	Identification of human papillomavirus type 16 variants circulating in the Calabria region by sequencing and phylogenetic analysis of HPV16 from cervical smears. Infection, Genetics and Evolution, 2019, 68, 185-193.	1.0	10
18	HEVnet: a One Health, collaborative, interdisciplinary network and sequence data repository for enhanced hepatitis E virus molecular typing, characterisation and epidemiological investigations. Eurosurveillance, 2019, 24, .	3.9	53

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19	Improving preparedness to respond to cross-border hepatitis A outbreaks in the European Union/European Economic Area: towards comparable sequencing of hepatitis A virus. Eurosurveillance, 2019, 24, .	3.9	10
20	Hepatitis A outbreak affecting men who have sex with men (MSM) in South Italy. New Microbiologica, 2019, 42, 181-183.	0.1	5
21	The genetic diversity of hepatitis A genotype I in Bulgaria. Medicine (United States), 2018, 97, e9632.	0.4	9
22	Genetic Diversity Among Genogroup II Noroviruses and Progressive Emergence of GII.17 in Wastewaters in Italy (2011–2016) Revealed by Next-Generation and Sanger Sequencing. Food and Environmental Virology, 2018, 10, 141-150.	1.5	29
23	Hepatitis A outbreak disproportionately affecting men who have sex with men (MSM) in the European Union and European Economic Area, June 2016 to May 2017. Eurosurveillance, 2018, 23, .	3.9	128
24	Hepatitis E in Italy: 5 years of national epidemiological, virological and environmental surveillance, 2012 to 2016. Eurosurveillance, 2018, 23, .	3.9	28
25	Hepatitis E virus genotypes and subgenotypes causing acute hepatitis, Bulgaria, 2013–2015. PLoS ONE, 2018, 13, e0198045.	1.1	22
26	Following a patient with prolonged response against hepatitis E virus. Panminerva Medica, 2018, 60, 232-234.	0.2	1
27	A nationwide retrospective study on prevalence of hepatitis E virus infection in Italian blood donors. Blood Transfusion, 2018, 16, 413-421.	0.3	45
28	Methodological approach towards a Gap Assessment of the Serbian microbiology system in the function of surveillance in line with EU standards and acquis. Annali Dell'Istituto Superiore Di Sanita, 2018, 54, 324-331.	0.2	0
29	Molecular characterization of human adenoviruses in urban wastewaters using next generation and Sanger sequencing. Water Research, 2017, 121, 240-247.	5.3	48
30	Hepatitis A virus strains circulating during 1997-2015 in Campania, a Southern Italy region with periodic outbreaks. Journal of Medical Virology, 2017, 89, 1931-1936.	2.5	14
31	Hepatitis E Virus (Genotype 3) in Slurry Samples from Swine Farming Activities in Italy. Food and Environmental Virology, 2017, 9, 219-229.	1.5	16
32	Evolutionary dynamics of HBVâ€D7 subgenotype in Tunisia. Journal of Medical Virology, 2017, 89, 469-475.	2.5	5
33	Hepatitis a virus genotypes and strains from an endemic area of Europe, Bulgaria 2012–2014. BMC Infectious Diseases, 2017, 17, 497.	1.3	16
34	Hepatitis E and blood donation safety in selected European countries: a shift to screening?. Eurosurveillance, 2017, 22, .	3.9	144
35	A large prolonged outbreak of hepatitis A associated with consumption of frozen berries, Italy, 2013–14. Journal of Medical Microbiology, 2017, 66, 342-349.	0.7	41
36	Hepatitis E virus infection in Europe: surveillance and descriptive epidemiology of confirmed cases, 2005 to 2015. Eurosurveillance, 2017, 22, .	3.9	127

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37	Performance of rapid diagnostic tests for the detection of antibodies to hepatitis C virus in whole blood collected on dried blood spots. Journal of Viral Hepatitis, 2016, 23, 399-401.	1.0	26
38	Antiviral treatment of HBV positive pregnant women: an additional tool to reduce perinatal transmission. Pathogens and Global Health, 2016, 110, 275-276.	1.0	2
39	Hepatitis E. Vox Sanguinis, 2016, 110, 93-103.	0.7	48
40	Naturally Occurring Surface Antigen Variants of Hepatitis B Virus in Tunisian Patients. Intervirology, 2016, 59, 36-47.	1.2	12
41	Correlates of infection and molecular characterization of blood-borne HIV, HCV, and HBV infections in HIV-1 infected inmates in Italy. Medicine (United States), 2016, 95, e5257.	0.4	10
42	Hepatitis E virus: Assessment of the epidemiological situation in humans in Europe, 2014/15. Journal of Clinical Virology, 2016, 82, 9-16.	1.6	168
43	Evaluation of rapid tests for diagnosis of acute hepatitis E. Journal of Clinical Virology, 2016, 78, 4-8.	1.6	14
44	Key Role of Sequencing to Trace Hepatitis A Viruses Circulating in Italy During a Large Multi-Country European Foodborne Outbreak in 2013. PLoS ONE, 2016, 11, e0149642.	1.1	31
45	High prevalence of anti-hepatitis E virus antibodies among blood donors in central Italy, February to March 2014. Eurosurveillance, 2016, 21, .	3.9	68
46	A family cluster of hepatitis A virus due to an uncommon IA strain circulating in Campania (southern) Tj ETQq0 hepatitis A?. Infezioni in Medicina, 2016, 24, 230-3.	0 0 rgBT /0 0.7	Overlock 10 Tf 3
47	Naturally occurring mutations associated with resistance to HCV NS5B polymerase and NS3 protease inhibitors in treatment-na $ ilde{A}$ ve patients with chronic hepatitis C. Virology Journal, 2015, 12, 186.	1.4	38
48	Woodchuck hepatitis virus core gene deletions and proliferative responses of peripheral blood mononuclear cells stimulated by an immunodominant epitope: a viral immune escape in the woodchuck model of chronic hepatitis B?. Archives of Virology, 2015, 160, 1065-1073.	0.9	1
49	Migration pattern of hepatitis A virus genotype IA in North-Central Tunisia. Virology Journal, 2015, 12, 17.	1.4	3
50	Evidence for the presence of autochthonous (locally acquired) cases of acute hepatitis E virus infections in Italy since the 80s. European Journal of Internal Medicine, 2015, 26, 348-350.	1.0	13
51	Hepatitis A and E Viruses in Wastewaters, in River Waters, and in Bivalve Molluscs in Italy. Food and Environmental Virology, 2015, 7, 316-324.	1.5	66
52	Pointâ€of are Screening, Prevalence, and Risk Factors for Hepatitis B Infection Among 3,728 Mainly Undocumented Migrants From Nonâ€EU Countries in Northern Italy. Journal of Travel Medicine, 2015, 22, 78-86.	1.4	38
53	Molecular epidemiology and phylogenetic analysis of Hepatitis B virus in a group of migrants in Italy. BMC Infectious Diseases, 2015, 15, 287.	1.3	12
54	Hepatitis E virus genotypes 1 and 3 in wastewater samples in Tunisia. Archives of Virology, 2015, 160, 183-189.	0.9	14

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55	Large and prolonged food-borne multistate hepatitis A outbreak in Europe associated with consumption of frozen berries, 2013 to 2014. Eurosurveillance, 2015, 20, 21192.	3.9	101
56	microRNA levels in paraffin-embedded indolent B-cell non-Hodgkin lymphoma tissues from patients chronically infected with hepatitis B or C virus. BMC Infectious Diseases, 2014, 14, S6.	1.3	14
57	Update on epidemiology of HCV in Italy: focus on the Calabria Region. BMC Infectious Diseases, 2014, 14, S2.	1.3	40
58	Evolutionary dynamics of HBVâ€D1 genotype epidemic in Turkey. Journal of Medical Virology, 2014, 86, 109-116.	2.5	20
59	Back to the origin of HCV 2c subtype and spreading to the Calabria region (Southern Italy) over the last two centuries: A phylogenetic study. Infection, Genetics and Evolution, 2014, 26, 352-358.	1.0	21
60	Qualitative and Quantitative Assessment of Hepatitis A Virus in Wastewaters in Tunisia. Food and Environmental Virology, 2014, 6, 246-252.	1.5	19
61	Surveillance of hepatitis A virus in urban sewages and comparison with cases notified in the course of an outbreak, Italy 2013. BMC Infectious Diseases, 2014, 14, 419.	1.3	66
62	Molecular characterisation of human hepatitis E virus from Italy: comparative analysis of five reverse transcription-PCR assays. Virology Journal, 2014, 11, 72.	1.4	25
63	Hepatitis A outbreak in Italy, 2013: a matched case–control study. Eurosurveillance, 2014, 19, .	3.9	11
64	Hepatitis C virus infection in an endemic area of Southern Italy 14 years later: Evidence for a vanishing infection. Digestive and Liver Disease, 2013, 45, 403-407.	0.4	58
65	IFN-α Regulates Blimp-1 Expression via miR-23a and miR-125b in Both Monocytes-Derived DC and pDC. PLoS ONE, 2013, 8, e72833.	1.1	26
66	Ongoing outbreak of hepatitis A in Italy: preliminary report as of 31 May 2013. Eurosurveillance, 2013, 18, 20518.	3.9	9
67	929 HEPATITIS C VIRUS INFECTION IN AN ENDEMIC SOUTHERN AREA OF ITALY 14 YEARS LATER: EVIDENCE FOR A VANISHING INFECTION. Journal of Hepatology, 2012, 56, S362.	1.8	O
68	Hepatitis C virus genotype 4d in Southern Italy: Reconstruction of its origin and spread by a phylodynamic analysis. Journal of Medical Virology, 2012, 84, 1613-1619.	2.5	29
69	Diagnosis of HEV infection by serological and real-time PCR assays: a study on acute non-A-C hepatitis collected from 2004 to 2010 in Italy. BMC Research Notes, 2012, 5, 297.	0.6	19
70	Phylogeny and phylodinamic of Hepatitis C in Italy. BMC Infectious Diseases, 2012, 12, S5.	1.3	17
71	Janus-faced liposomes enhance antimicrobial innate immune response in <i>Mycobacterium tuberculosis</i> infection. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, E1360-8.	3.3	60
72	Strong CD8+ T cell antigenicity and immunogenicity of large foreign proteins incorporated in HIV-1 VLPs able to induce a Nef-dependent activation/maturation of dendritic cells. Vaccine, 2011, 29, 3465-3475.	1.7	17

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73	An integrated approach identifies IFN-regulated microRNAs and targeted mRNAs modulated by different HCV replicon clones. BMC Genomics, 2011, 12, 485.	1.2	23
74	A computational approach to identify point mutations associated with occult hepatitis B: significant mutations affect coding regions but not regulative elements of HBV. Virology Journal, 2011, 8, 394.	1.4	9
75	Reconstruction of the evolutionary dynamics of the hepatitis C virus 1b epidemic in Turkey. Infection, Genetics and Evolution, 2011, 11, 863-868.	1.0	17
76	May Phylogenetic Analysis Support Epidemiological Investigation in Identifying the Source of HIV Infection?. AIDS Research and Human Retroviruses, 2011, 27, 455-457.	0.5	2
77	Improving HIV-2 Detection by a Combination of Serological and Nucleic Acid Amplification Test Assays. Journal of Clinical Microbiology, 2010, 48, 2902-2908.	1.8	7
78	A computational approach identifies two regions of Hepatitis C Virus E1 protein as interacting domains involved in viral fusion process. BMC Structural Biology, 2009, 9, 48.	2.3	9
79	Microarray analysis identifies a common set of cellular genes modulated by different HCV replicon clones. BMC Genomics, 2008, 9, 309.	1.2	15
80	Tâ€cellâ€mediated and antigenâ€dependent differentiation of human monocyte into different dendritic cell subsets: a feedback control of Th1/Th2 responses. FASEB Journal, 2008, 22, 3370-3379.	0.2	12
81	Repression of Interferon Regulatory Factor 1 by Hepatitis C Virus Core Protein Results in Inhibition of Antiviral and Immunomodulatory Genes. Journal of Virology, 2007, 81, 202-214.	1.5	53
82	Activation of the ER stress gene gadd153 by hepatitis C virus sensitizes cells to oxidant injury. Virus Research, 2007, 126, 128-138.	1.1	29
83	[428] MICROARRAY ANALYSIS OF LIVER CELLS CONTAINING A FULL-LENGHT HEPATITIS C VIRUS REPLICON. Journal of Hepatology, 2007, 46, S164.	1.8	0
84	Activation of endoplasmic reticulum stress response by hepatitis C virus proteins. Archives of Virology, 2005, 150, 1339-1356.	0.9	26
85	The transmembrane domain of hepatitis C virus E1 glycoprotein induces cell death. Virus Research, 2004, 104, 1-9.	1.1	23
86	285 HCV expression in a tetracycline-regulated cell line activates endoplasmic reticulum stress-mediated apoptosis. Journal of Hepatology, 2004, 40, 88.	1.8	0
87	Expression of HCV E1 transmembrane region in eucaryotic systems: Effect on cell viability. Journal of Hepatology, 2003, 38, 117-118.	1.8	0
88	Expression of HCV E1 Protein in Baculovirus-Infected Cells: Effects on Cell Viability and Apoptosis Induction. Intervirology, 2003, 46, 121-126.	1.2	18
89	The expression of the transmembrane domains of HCV E1 protein induce cell death. Journal of Hepatology, 2002, 36, 5.	1.8	0
90	Mutagenesis of hepatitis C virus E1 protein affects its membrane-permeabilizing activity. Journal of General Virology, 2001, 82, 2243-2250.	1.3	19

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91	Molecular analysis of hepatitis C virus E1 channel protein. Journal of Hepatology, 2001, 34, 115-116.	1.8	O
92	Expression and membrane association of hepatitis C virus envelope 1 protein. Virus Genes, 2000, 21, 223-226.	0.7	9
93	Persistence of HCV–RNA in a Blood Donor with Negative Antibody Assays. Vox Sanguinis, 1999, 76, 192-193.	0.7	4
94	Hepatitis C Virus E1 Protein Induces Modification of Membrane Permeability in E. coliCells. Virology, 1998, 250, 1-8.	1.1	24
95	Secretion and purification of HCV E1 protein forms as glutathione-S-transferase fusion in the baculovirus insect cell system. Virus Research, 1998, 55, 157-165.	1.1	12
96	Genotyping HCV isolates from Italy by type-specific PCR assay in the core region. Research in Virology, 1998, 149, 209-218.	0.7	14
97	Correlation between virus genotype and chronicity rate in acute hepatitis C. Journal of Hepatology, 1998, 28, 939-944.	1.8	107
98	Recurrence of WHV Integration in theb3nLocus in Woodchuck Hepatocellular Carcinoma. Virology, 1995, 214, 229-234.	1.1	15
99	Inhibition of woodchuck hepatitis virus replication by adenine arabinoside monophosphate coupled to lactosaminated poly-lysine and administered by intramuscular route*1. Hepatology, 1995, 22, 1072-1077.	3.6	16
100	Hepatitis C Virus Infection as a Risk Factor for Hepatocellular Carcinoma in Patients with Cirrhosis. Annals of Internal Medicine, 1992, 116, 97-102.	2.0	328
101	Age- and sex-related study of HBV-DNA in HBsAg asymptomatic children from an endemic area (Cameroon). Annals of Tropical Paediatrics, 1991, 11, 325-329.	1.0	7
102	Prevalence of HBeAg, anti-HBe serological markers and HBV-DNA in asymptomatic carriers in Ethiopia. European Journal of Epidemiology, 1989, 5, 481-485.	2.5	6