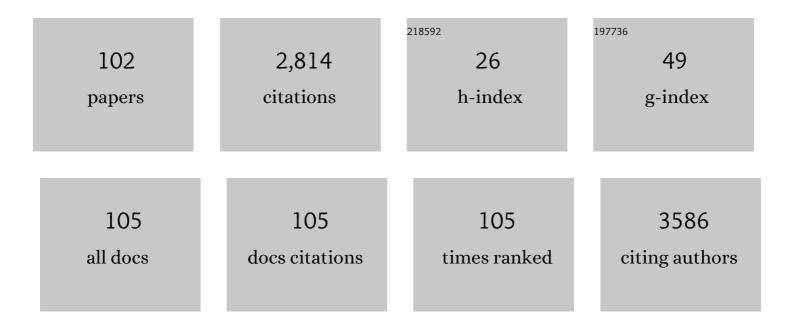
Anna Rita Ciccaglione

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	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
2	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
3	Hepatitis E and blood donation safety in selected European countries: a shift to screening?. Eurosurveillance, 2017, 22, .	3.9	144
4	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
5	Hepatitis E virus infection in Europe: surveillance and descriptive epidemiology of confirmed cases, 2005 to 2015. Eurosurveillance, 2017, 22, .	3.9	127
6	Correlation between virus genotype and chronicity rate in acute hepatitis C. Journal of Hepatology, 1998, 28, 939-944.	1.8	107
7	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
8	High prevalence of anti-hepatitis E virus antibodies among blood donors in central Italy, February to March 2014. Eurosurveillance, 2016, 21, .	3.9	68
9	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
10	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
11	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
12	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
13	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
14	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
15	Hepatitis E. Vox Sanguinis, 2016, 110, 93-103.	0.7	48
16	Molecular characterization of human adenoviruses in urban wastewaters using next generation and Sanger sequencing. Water Research, 2017, 121, 240-247.	5.3	48
17	A nationwide retrospective study on prevalence of hepatitis E virus infection in Italian blood donors. Blood Transfusion, 2018, 16, 413-421.	0.3	45
18	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

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19	Update on epidemiology of HCV in Italy: focus on the Calabria Region. BMC Infectious Diseases, 2014, 14, S2.	1.3	40
20	Naturally occurring mutations associated with resistance to HCV NS5B polymerase and NS3 protease inhibitors in treatment-naÃ ⁻ ve patients with chronic hepatitis C. Virology Journal, 2015, 12, 186.	1.4	38
21	Pointâ€ofâ€Care 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
22	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
23	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
24	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
25	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
26	Hepatitis E in Italy: 5 years of national epidemiological, virological and environmental surveillance, 2012 to 2016. Eurosurveillance, 2018, 23, .	3.9	28
27	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
28	Activation of endoplasmic reticulum stress response by hepatitis C virus proteins. Archives of Virology, 2005, 150, 1339-1356.	0.9	26
29	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
30	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
31	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
32	Hepatitis C Virus E1 Protein Induces Modification of Membrane Permeability inE. coliCells. Virology, 1998, 250, 1-8.	1.1	24
33	The transmembrane domain of hepatitis C virus E1 glycoprotein induces cell death. Virus Research, 2004, 104, 1-9.	1.1	23
34	An integrated approach identifies IFN-regulated microRNAs and targeted mRNAs modulated by different HCV replicon clones. BMC Genomics, 2011, 12, 485.	1.2	23
35	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
36	Hepatitis E virus genotypes and subgenotypes causing acute hepatitis, Bulgaria, 2013–2015. PLoS ONE, 2018, 13, e0198045.	1.1	22

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37	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
38	Evolutionary dynamics of HBVâ€Ð1 genotype epidemic in Turkey. Journal of Medical Virology, 2014, 86, 109-116.	2.5	20
39	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
40	Qualitative and Quantitative Assessment of Hepatitis A Virus in Wastewaters in Tunisia. Food and Environmental Virology, 2014, 6, 246-252.	1.5	19
41	Mutagenesis of hepatitis C virus E1 protein affects its membrane-permeabilizing activity. Journal of General Virology, 2001, 82, 2243-2250.	1.3	19
42	Expression of HCV E1 Protein in Baculovirus-Infected Cells: Effects on Cell Viability and Apoptosis Induction. Intervirology, 2003, 46, 121-126.	1.2	18
43	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
44	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
45	Phylogeny and phylodinamic of Hepatitis C in Italy. BMC Infectious Diseases, 2012, 12, S5.	1.3	17
46	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
47	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
48	Hepatitis a virus genotypes and strains from an endemic area of Europe, Bulgaria 2012–2014. BMC Infectious Diseases, 2017, 17, 497.	1.3	16
49	Recurrence of WHV Integration in theb3nLocus in Woodchuck Hepatocellular Carcinoma. Virology, 1995, 214, 229-234.	1.1	15
50	Microarray analysis identifies a common set of cellular genes modulated by different HCV replicon clones. BMC Genomics, 2008, 9, 309.	1.2	15
51	Genotyping HCV isolates from Italy by type-specific PCR assay in the core region. Research in Virology, 1998, 149, 209-218.	0.7	14
52	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
53	Hepatitis E virus genotypes 1 and 3 in wastewater samples in Tunisia. Archives of Virology, 2015, 160, 183-189.	0.9	14
54	Evaluation of rapid tests for diagnosis of acute hepatitis E. Journal of Clinical Virology, 2016, 78, 4-8.	1.6	14

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55	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
56	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
57	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
58	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
59	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
60	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
61	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
62	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
63	Naturally Occurring Surface Antigen Variants of Hepatitis B Virus in Tunisian Patients. Intervirology, 2016, 59, 36-47.	1.2	12
64	Hepatitis A outbreak in Italy, 2013: a matched case–control study. Eurosurveillance, 2014, 19, .	3.9	11
65	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
66	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
67	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
68	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
69	Expression and membrane association of hepatitis C virus envelope 1 protein. Virus Genes, 2000, 21, 223-226.	0.7	9
70	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
71	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
72	The genetic diversity of hepatitis A genotype I in Bulgaria. Medicine (United States), 2018, 97, e9632.	0.4	9

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73	Ongoing outbreak of hepatitis A in Italy: preliminary report as of 31 May 2013. Eurosurveillance, 2013, 18, 20518.	3.9	9
74	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
75	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
76	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
77	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
78	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
79	Evolutionary dynamics of HBVâ€D7 subgenotype in Tunisia. Journal of Medical Virology, 2017, 89, 469-475.	2.5	5
80	An Evaluation of Hepatitis E Virus Molecular Typing Methods. Clinical Chemistry, 2021, 68, 181-191.	1.5	5
81	Immunogenicity of Viral Vaccines in the Italian Military. Biomedicines, 2021, 9, 87.	1.4	5
82	Hepatitis A outbreak affecting men who have sex with men (MSM) in South Italy. New Microbiologica, 2019, 42, 181-183.	0.1	5
83	Persistence of HCV–RNA in a Blood Donor with Negative Antibody Assays. Vox Sanguinis, 1999, 76, 192-193.	0.7	4
84	Migration pattern of hepatitis A virus genotype IA in North-Central Tunisia. Virology Journal, 2015, 12, 17.	1.4	3
85	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
86	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
87	A family cluster of hepatitis A virus due to an uncommon IA strain circulating in Campania (southern) Tj ETQq1 hepatitis A?. Infezioni in Medicina, 2016, 24, 230-3.	1 0.784314 0.7	rgBT /Overlo 3
88	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
89	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
90	Time and Mode of Epidemic HCV-2 Subtypes Spreading in Europe: Phylodynamics in Italy and Albania. Diagnostics, 2021, 11, 327.	1.3	2

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91	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
92	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
93	Following a patient with prolonged response against hepatitis E virus. Panminerva Medica, 2018, 60, 232-234.	0.2	1
94	Sensitivity of hepatitis C virus rapid tests in detecting antibodies in general population. Panminerva Medica, 2020, 62, 125-130.	0.2	1
95	Changing epidemiology of acute liver failure in Italy: a single-center experience over 25 years. Minerva Medica, 2020, 111, 330-336.	0.3	1
96	The expression of the transmembrane domains of HCV E1 protein induce cell death. Journal of Hepatology, 2002, 36, 5.	1.8	0
97	Expression of HCV E1 transmembrane region in eucaryotic systems: Effect on cell viability. Journal of Hepatology, 2003, 38, 117-118.	1.8	0
98	285 HCV expression in a tetracycline-regulated cell line activates endoplasmic reticulum stress-mediated apoptosis. Journal of Hepatology, 2004, 40, 88.	1.8	0
99	[428] MICROARRAY ANALYSIS OF LIVER CELLS CONTAINING A FULL-LENGHT HEPATITIS C VIRUS REPLICON. Journal of Hepatology, 2007, 46, S164.	1.8	0
100	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	0
101	Molecular analysis of hepatitis C virus E1 channel protein. Journal of Hepatology, 2001, 34, 115-116.	1.8	0
102	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