

Roberto Bruni

List of Publications by Year in descending order

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Version: 2024-02-01

71
papers

1,303
citations

361045

20
h-index

433756

31
g-index

72
all docs

72
docs citations

72
times ranked

1861
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. <i>Eurosurveillance</i> , 2022, 27, .	3.9	7
2	Phylogenetic and Molecular Analyses of More Prevalent HCV1b Subtype in the Calabria Region, Southern Italy. <i>Journal of Clinical Medicine</i> , 2021, 10, 1655.	1.0	3
3	Developing and Piloting a Standardized European Protocol for Hepatitis C Prevalence Surveys in the General Population (2016â€“2019). <i>Frontiers in Public Health</i> , 2021, 9, 568524.	1.3	1
4	Hepatitis E Outbreak in the Central Part of Italy Sustained by Multiple HEV Genotype 3 Strains, Juneâ€“December 2019. <i>Viruses</i> , 2021, 13, 1159.	1.5	14
5	Hepatitis A Virus Strains Circulating in the Campania Region (2015â€“2018) Assessed through Bivalve Biomonitoring and Environmental Surveillance. <i>Viruses</i> , 2021, 13, 16.	1.5	14
6	Changing epidemiology of hepatitis C in Italy: a population-based survey in a historically high endemic area. <i>Minerva Medica</i> , 2021, , .	0.3	0
7	Phylogenetic analysis and epidemiological history of Hepatitis E virus 3f and 3c in swine and wild boar, Italy. <i>Heliyon</i> , 2020, 6, e05110.	1.4	10
8	Nine-Year Nationwide Environmental Surveillance of Hepatitis E Virus in Urban Wastewaters in Italy (2011â€“2019). <i>International Journal of Environmental Research and Public Health</i> , 2020, 17, 2059.	1.2	27
9	Sensitivity of hepatitis C virus rapid tests in detecting antibodies in general population. <i>Panminerva Medica</i> , 2020, 62, 125-130.	0.2	1
10	Retrospective analysis of acute HBV infections occurred in 1978â€“79 and 1994â€“95 in North-East Italy: increasing prevalence of BCP/pre-core mutants in sub-genotype D3. <i>BMC Infectious Diseases</i> , 2020, 20, 78.	1.3	3
11	Incidence of hepatitis E virus infection among blood donors in a high endemic area of Central Italy. <i>Journal of Viral Hepatitis</i> , 2019, 26, 506-512.	1.0	22
12	Human hepatitis E virus circulation in Bulgaria: Deep Bayesian phylogenetic analysis for viral spread control in the country. <i>Journal of Medical Virology</i> , 2019, 91, 132-138.	2.5	8
13	Identification of human papillomavirus type 16 variants circulating in the Calabria region by sequencing and phylogenetic analysis of HPV16 from cervical smears. <i>Infection, Genetics and Evolution</i> , 2019, 68, 185-193.	1.0	10
14	HEVnet: a One Health, collaborative, interdisciplinary network and sequence data repository for enhanced hepatitis E virus molecular typing, characterisation and epidemiological investigations. <i>Eurosurveillance</i> , 2019, 24, .	3.9	53
15	Improving preparedness to respond to cross-border hepatitis A outbreaks in the European Union/European Economic Area: towards comparable sequencing of hepatitis A virus. <i>Eurosurveillance</i> , 2019, 24, .	3.9	10
16	Hepatitis E virus infection prevalence among men who have sex with men involved in a hepatitis A virus outbreak in Italy. <i>Blood Transfusion</i> , 2019, 17, 428-432.	0.3	1
17	Hepatitis A outbreak affecting men who have sex with men (MSM) in South Italy. <i>New Microbiologica</i> , 2019, 42, 181-183.	0.1	5
18	The genetic diversity of hepatitis A genotype I in Bulgaria. <i>Medicine (United States)</i> , 2018, 97, e9632.	0.4	9

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19	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. <i>Eurosurveillance</i> , 2018, 23, .	3.9	128
20	Hepatitis E in Italy: 5 years of national epidemiological, virological and environmental surveillance, 2012 to 2016. <i>Eurosurveillance</i> , 2018, 23, .	3.9	28
21	Hepatitis E virus genotypes and subgenotypes causing acute hepatitis, Bulgaria, 2013–2015. <i>PLoS ONE</i> , 2018, 13, e0198045.	1.1	22
22	Following a patient with prolonged response against hepatitis E virus. <i>Panminerva Medica</i> , 2018, 60, 232-234.	0.2	1
23	A nationwide retrospective study on prevalence of hepatitis E virus infection in Italian blood donors. <i>Blood Transfusion</i> , 2018, 16, 413-421.	0.3	45
24	Hepatitis A virus strains circulating during 1997-2015 in Campania, a Southern Italy region with periodic outbreaks. <i>Journal of Medical Virology</i> , 2017, 89, 1931-1936.	2.5	14
25	Hepatitis E Virus (Genotype 3) in Slurry Samples from Swine Farming Activities in Italy. <i>Food and Environmental Virology</i> , 2017, 9, 219-229.	1.5	16
26	Evolutionary dynamics of HBV- δ 7 subgenotype in Tunisia. <i>Journal of Medical Virology</i> , 2017, 89, 469-475.	2.5	5
27	Hepatitis a virus genotypes and strains from an endemic area of Europe, Bulgaria 2012–2014. <i>BMC Infectious Diseases</i> , 2017, 17, 497.	1.3	16
28	A large prolonged outbreak of hepatitis A associated with consumption of frozen berries, Italy, 2013–14. <i>Journal of Medical Microbiology</i> , 2017, 66, 342-349.	0.7	41
29	Antiviral treatment of HBV positive pregnant women: an additional tool to reduce perinatal transmission. <i>Pathogens and Global Health</i> , 2016, 110, 275-276.	1.0	2
30	Naturally Occurring Surface Antigen Variants of Hepatitis B Virus in Tunisian Patients. <i>Intervirology</i> , 2016, 59, 36-47.	1.2	12
31	Correlates of infection and molecular characterization of blood-borne HIV, HCV, and HBV infections in HIV-1 infected inmates in Italy. <i>Medicine (United States)</i> , 2016, 95, e5257.	0.4	10
32	Evaluation of rapid tests for diagnosis of acute hepatitis E. <i>Journal of Clinical Virology</i> , 2016, 78, 4-8.	1.6	14
33	Key Role of Sequencing to Trace Hepatitis A Viruses Circulating in Italy During a Large Multi-Country European Foodborne Outbreak in 2013. <i>PLoS ONE</i> , 2016, 11, e0149642.	1.1	31
34	High prevalence of anti-hepatitis E virus antibodies among blood donors in central Italy, February to March 2014. <i>Eurosurveillance</i> , 2016, 21, .	3.9	68
35	Naturally occurring mutations associated with resistance to HCV NS5B polymerase and NS3 protease inhibitors in treatment-naïve patients with chronic hepatitis C. <i>Virology Journal</i> , 2015, 12, 186.	1.4	38
36	Migration pattern of hepatitis A virus genotype IA in North-Central Tunisia. <i>Virology Journal</i> , 2015, 12, 17.	1.4	3

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37	Hepatitis A and E Viruses in Wastewaters, in River Waters, and in Bivalve Molluscs in Italy. <i>Food and Environmental Virology</i> , 2015, 7, 316-324.	1.5	66
38	Molecular epidemiology and phylogenetic analysis of Hepatitis B virus in a group of migrants in Italy. <i>BMC Infectious Diseases</i> , 2015, 15, 287.	1.3	12
39	Hepatitis E virus genotypes 1 and 3 in wastewater samples in Tunisia. <i>Archives of Virology</i> , 2015, 160, 183-189.	0.9	14
40	microRNA levels in paraffin-embedded indolent B-cell non-Hodgkin lymphoma tissues from patients chronically infected with hepatitis B or C virus. <i>BMC Infectious Diseases</i> , 2014, 14, S6.	1.3	14
41	Evolutionary dynamics of HBV Δ 1 genotype epidemic in Turkey. <i>Journal of Medical Virology</i> , 2014, 86, 109-116.	2.5	20
42	Back to the origin of HCV 2c subtype and spreading to the Calabria region (Southern Italy) over the last two centuries: A phylogenetic study. <i>Infection, Genetics and Evolution</i> , 2014, 26, 352-358.	1.0	21
43	Qualitative and Quantitative Assessment of Hepatitis A Virus in Wastewaters in Tunisia. <i>Food and Environmental Virology</i> , 2014, 6, 246-252.	1.5	19
44	Surveillance of hepatitis A virus in urban sewages and comparison with cases notified in the course of an outbreak, Italy 2013. <i>BMC Infectious Diseases</i> , 2014, 14, 419.	1.3	66
45	Molecular characterisation of human hepatitis E virus from Italy: comparative analysis of five reverse transcription-PCR assays. <i>Virology Journal</i> , 2014, 11, 72.	1.4	25
46	Hepatitis A outbreak in Italy, 2013: a matched case-control study. <i>Eurosurveillance</i> , 2014, 19, .	3.9	11
47	IFN γ Regulates Blimp-1 Expression via miR-23a and miR-125b in Both Monocytes-Derived DC and pDC. <i>PLoS ONE</i> , 2013, 8, e72833.	1.1	26
48	Ongoing outbreak of hepatitis A in Italy: preliminary report as of 31 May 2013. <i>Eurosurveillance</i> , 2013, 18, 20518.	3.9	9
49	Hepatitis C virus genotype 4d in Southern Italy: Reconstruction of its origin and spread by a phylodynamic analysis. <i>Journal of Medical Virology</i> , 2012, 84, 1613-1619.	2.5	29
50	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. <i>BMC Research Notes</i> , 2012, 5, 297.	0.6	19
51	An integrated approach identifies IFN-regulated microRNAs and targeted mRNAs modulated by different HCV replicon clones. <i>BMC Genomics</i> , 2011, 12, 485.	1.2	23
52	A computational approach to identify point mutations associated with occult hepatitis B: significant mutations affect coding regions but not regulative elements of HBV. <i>Virology Journal</i> , 2011, 8, 394.	1.4	9
53	Reconstruction of the evolutionary dynamics of the hepatitis C virus 1b epidemic in Turkey. <i>Infection, Genetics and Evolution</i> , 2011, 11, 863-868.	1.0	17
54	Improving HIV-2 Detection by a Combination of Serological and Nucleic Acid Amplification Test Assays. <i>Journal of Clinical Microbiology</i> , 2010, 48, 2902-2908.	1.8	7

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55	A computational approach identifies two regions of Hepatitis C Virus E1 protein as interacting domains involved in viral fusion process. <i>BMC Structural Biology</i> , 2009, 9, 48.	2.3	9
56	Microarray analysis identifies a common set of cellular genes modulated by different HCV replicon clones. <i>BMC Genomics</i> , 2008, 9, 309.	1.2	15
57	Lack of WHV integration nearby N-myc2 and in the downstream b3n and win loci in a considerable fraction of liver tumors with activated N-myc2 from naturally infected wild woodchucks. <i>Virology</i> , 2006, 345, 258-269.	1.1	13
58	The win locus involved in activation of the distal N-myc2 gene upon WHV integration in woodchuck liver tumors harbors S/MAR elements. <i>Virology</i> , 2004, 329, 1-10.	1.1	12
59	Immunization of woodchucks with adjuvanted sHDAg (p24): immune response and outcome following challenge. <i>Vaccine</i> , 2004, 22, 457-466.	1.7	14
60	Scaffold attachment region located in a locus targeted by hepadnavirus integration in hepatocellular carcinomas. <i>Cancer Detection and Prevention</i> , 2003, 27, 175-181.	2.1	3
61	Ultrasonography in the study of hepatocellular carcinoma in woodchucks chronically infected with WHV. <i>Laboratory Animals</i> , 2003, 37, 233-240.	0.5	10
62	Cellular gene activation by HBV integration in or close to chromosomal regulative elements: a hypothesis from the WHV/woodchuck model. <i>Journal of Hepatology</i> , 2002, 36, 83-84.	1.8	0
63	In vivo transmission and dynamics of deleted genomes after experimental infection of woodchuck hepatitis B virus in adult animals. <i>Virus Genes</i> , 2002, 25, 147-157.	0.7	7
64	Activation of the N-myc2 Oncogene by Woodchuck Hepatitis Virus Integration in the Linked Downstreamb3nLocus in Woodchuck Hepatocellular Carcinoma. <i>Virology</i> , 1999, 257, 483-490.	1.1	27
65	Hepadnavirus evolution and molecular strategy of adaptation in a new host.. <i>Journal of General Virology</i> , 1999, 80, 617-626.	1.3	15
66	Identification of Scaffold/Matrix Attachment Region in Recurrent Site of Woodchuck Hepatitis Virus Integration. <i>DNA and Cell Biology</i> , 1998, 17, 519-527.	0.9	15
67	Woodchuck hepatitis virus DNA integration in a common chromosomal region of the woodchuck genome in two independent hepatocellular carcinomas. <i>Archives of Virology</i> , 1997, 142, 499-509.	0.9	4
68	Sequence and phylogenetic analysis of the VP1 gene in two cell culture-adapted HAV strains from a unique pathogenic isolate. <i>Virus Genes</i> , 1995, 10, 37-43.	0.7	3
69	A PCR-based strategy for rapid mapping of hepadnavirus integrated sequences in hepatocellular carcinomas. <i>Journal of Virological Methods</i> , 1995, 52, 347-360.	1.0	7
70	Recurrence of WHV Integration in theb3nLocus in Woodchuck Hepatocellular Carcinoma. <i>Virology</i> , 1995, 214, 229-234.	1.1	15
71	<i>Vibrio cholerae</i> in the Horn of Africa: Epidemiology, Plasmids, Tetracycline Resistance Gene Amplification, and Comparison Between O1 and Non-O1 Strains. <i>American Journal of Tropical Medicine and Hygiene</i> , 1995, 53, 351-359.	0.6	32