

Chiara Chiapponi

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1824818/publications.pdf>

Version: 2024-02-01

42
papers

1,526
citations

361413

20
h-index

315739

38
g-index

42
all docs

42
docs citations

42
times ranked

1996
citing authors

#	ARTICLE	IF	CITATIONS
1	Taxonomy of the order Mononegavirales: update 2019. Archives of Virology, 2019, 164, 1967-1980.	2.1	224
2	Molecular Epidemiology and Evolution of Influenza Viruses Circulating within European Swine between 2009 and 2013. Journal of Virology, 2015, 89, 9920-9931.	3.4	148
3	Replication, Pathogenesis and Transmission of Pandemic (H1N1) 2009 Virus in Non-Immune Pigs. PLoS ONE, 2010, 5, e9068.	2.5	144
4	Novel H1N2 swine influenza reassortant strain in pigs derived from the pandemic H1N1/2009 virus. Veterinary Microbiology, 2011, 149, 472-477.	1.9	115
5	European Surveillance Network for Influenza in Pigs: Surveillance Programs, Diagnostic Tools and Swine Influenza Virus Subtypes Identified in 14 European Countries from 2010 to 2013. PLoS ONE, 2014, 9, e115815.	2.5	107
6	Detection of Influenza D Virus among Swine and Cattle, Italy. Emerging Infectious Diseases, 2016, 22, 352-354.	4.3	90
7	Virological Surveillance and Preliminary Antigenic Characterization of Influenza Viruses in Pigs in Five European Countries from 2006 to 2008. Zoonoses and Public Health, 2011, 58, 93-101.	2.2	73
8	Influenza D in Italy: towards a better understanding of an emerging viral infection in swine. Scientific Reports, 2017, 7, 11660.	3.3	59
9	Co-circulation of two Usutu virus strains in Northern Italy between 2009 and 2014. Infection, Genetics and Evolution, 2017, 51, 255-262.	2.3	39
10	Rapid detection and subtyping of European swine influenza viruses in porcine clinical samples by haemagglutinin- and neuraminidase-specific tetra- and triplex real-time RT-PCR. Influenza and Other Respiratory Viruses, 2016, 10, 504-517.	3.4	37
11	Detection and Characterization of a Novel Reassortant Mammalian Orthoreovirus in Bats in Europe. Viruses, 2015, 7, 5844-5854.	3.3	35
12	Genomic characterization of H1N2 swine influenza viruses in Italy. Veterinary Microbiology, 2012, 156, 265-276.	1.9	30
13	First identification of mammalian orthoreovirus type 3 in diarrheic pigs in Europe. Virology Journal, 2016, 13, 139.	3.4	30
14	Isolation and Genomic Sequence of Hepatitis A Virus from Mixed Frozen Berries in Italy. Food and Environmental Virology, 2014, 6, 202-206.	3.4	29
15	Development and evaluation of a new Real-Time RT-PCR assay for detection of proposed influenza D virus. Journal of Virological Methods, 2017, 243, 31-34.	2.1	29
16	Genetic analysis of human and swine influenza A viruses isolated in Northern Italy during 2010-2015. Zoonoses and Public Health, 2018, 65, 114-123.	2.2	29
17	Comparison of the usefulness of the CACO-2 cell line with standard substrates for isolation of swine influenza A viruses. Journal of Virological Methods, 2010, 163, 162-165.	2.1	27
18	Multiplex RT-PCR assay for differentiating European swine influenza virus subtypes H1N1, H1N2 and H3N2. Journal of Virological Methods, 2012, 184, 117-120.	2.1	26

#	ARTICLE	IF	CITATIONS
19	Detection of a New Genetic Cluster of Influenza D Virus in Italian Cattle. <i>Viruses</i> , 2019, 11, 1110.	3.3	25
20	Isolation of three novel reassortant phleboviruses, Ponticelli I, II, III, and of Toscana virus from field-collected sandflies in Italy. <i>Parasites and Vectors</i> , 2018, 11, 84.	2.5	21
21	MAb-based competitive ELISA for the detection of antibodies against influenza D virus. <i>Transboundary and Emerging Diseases</i> , 2019, 66, 268-276.	3.0	21
22	Isolation of a novel Rhabdovirus from an insectivorous bat (<i>Pipistrellus kuhlii</i>) in Italy. <i>Virology Journal</i> , 2018, 15, 37.	3.4	19
23	Emerging Influenza D virus infection in European livestock as determined in serology studies: Are we underestimating its spread over the continent?. <i>Transboundary and Emerging Diseases</i> , 2021, 68, 1125-1135.	3.0	18
24	Different evolutionary trends of swine H1N2 influenza viruses in Italy compared to European viruses. <i>Veterinary Research</i> , 2013, 44, 112.	3.0	15
25	Detection of antibodies against influenza D virus in swine veterinarians in Italy in 2004. <i>Journal of Medical Virology</i> , 2022, 94, 2855-2859.	5.0	15
26	Co-circulation of eight different phleboviruses in sand flies collected in the Northern Apennine Mountains (Italy). <i>Infection, Genetics and Evolution</i> , 2018, 64, 131-134.	2.3	12
27	Hypugopoxvirus: A Novel Poxvirus Isolated from <i>Hypsugo savii</i> in Italy. <i>Viruses</i> , 2019, 11, 568.	3.3	12
28	Host range of mammalian orthoreovirus type 3 widening to alpine chamois. <i>Veterinary Microbiology</i> , 2019, 230, 72-77.	1.9	12
29	Epidemiological survey of swine influenza A virus in the wild boar population of two Italian provinces. <i>Influenza and Other Respiratory Viruses</i> , 2013, 7, 16-20.	3.4	11
30	Co-Circulation of Phleboviruses and Leishmania Parasites in Sand Flies from a Single Site in Italy Monitored between 2017 and 2020. <i>Viruses</i> , 2021, 13, 1660.	3.3	11
31	Full-Genome Sequence of a Reassortant H1N1 Swine Influenza Virus Isolated from Pigs in Italy. <i>Genome Announcements</i> , 2013, 1, .	0.8	9
32	Surveillance for Adenoviruses in Bats in Italy. <i>Viruses</i> , 2019, 11, 523.	3.3	9
33	First expert elicitation of knowledge on drivers of emergence of influenza D in Europe. <i>Transboundary and Emerging Diseases</i> , 2021, 68, 3349-3359.	3.0	9
34	Identification and Characterization of Swine Influenza Virus H1N1 Variants Generated in Vaccinated and Nonvaccinated, Challenged Pigs. <i>Viruses</i> , 2021, 13, 2087.	3.3	9
35	Genetic Variability among Swine Influenza Viruses in Italy: Data Analysis of the Period 2017–2020. <i>Viruses</i> , 2022, 14, 47.	3.3	8
36	Detection of influenza A virus by RT-PCR and standard methods in experimental infection of Ducks. <i>New Microbiologica</i> , 2005, 28, 31-5.	0.1	6

#	ARTICLE	IF	CITATIONS
37	Swine Influenza A(H3N2) Virus Infection in Immunocompromised Man, Italy, 2014. <i>Emerging Infectious Diseases</i> , 2015, 21, 1189-91.	4.3	3
38	Protein mutations following adaptation of avian influenza viruses in different biological systems. <i>Research in Veterinary Science</i> , 2015, 103, 176-178.	1.9	3
39	Influenza A in Wild Boars: Viral Circulation in the Emilia-Romagna Region (Northern Italy) between 2017 and 2022. <i>Animals</i> , 2022, 12, 1593.	2.3	3
40	Temporal insight into the natural generation of a new reassortant porcine influenza virus in a swine holding. <i>Veterinary Microbiology</i> , 2014, 174, 9-15.	1.9	2
41	Risk assessment for influenza D in Europe. <i>EFSA Supporting Publications</i> , 2020, 17, 1853E.	0.7	2
42	Retrospective Characterization of the 2006â€“2007 Swine Vesicular Disease Epidemic in Northern Italy by Whole Genome Sequence Analysis. <i>Viruses</i> , 2021, 13, 1186.	3.3	0