Alessio Lorusso

List of Publications by Year in descending order

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Version: 2024-02-01

116 papers 3,620 citations

34 h-index 54 g-index

121 all docs

121 docs citations

times ranked

121

4504 citing authors

#	Article	IF	CITATIONS
1	Novel human coronavirus (SARS-CoV-2): A lesson from animal coronaviruses. Veterinary Microbiology, 2020, 244, 108693.	1.9	298
2	Recombinant Canine Coronaviruses Related to Transmissible Gastroenteritis Virus of Swine Are Circulating in Dogs. Journal of Virology, 2009, 83, 1532-1537.	3.4	123
3	Genetic and antigenic characterization of H1 influenza viruses from United States swine from 2008. Journal of General Virology, 2011, 92, 919-930.	2.9	123
4	Detection and quantification of Anaplasma marginale DNA in blood samples of cattle by real-time PCR. Veterinary Microbiology, 2007, 124, 107-114.	1.9	114
5	Usutu virus infections in humans: a retrospective analysis in the municipality of Modena, Italy. Clinical Microbiology and Infection, 2017, 23, 33-37.	6.0	112
6	Genetic analysis of canine parvovirus type 2c. Virology, 2009, 385, 5-10.	2.4	108
7	A minor groove binder probe real-time PCR assay for discrimination between type 2-based vaccines and field strains of canine parvovirus. Journal of Virological Methods, 2006, 136, 65-70.	2.1	101
8	Novel putative Bluetongue virus in healthy goats from Sardinia, Italy. Infection, Genetics and Evolution, 2017, 51, 108-117.	2.3	89
9	Infection sustained by lineage B.1.1.7 of SARS-CoV-2 is characterised by longer persistence and higher viral RNA loads in nasopharyngeal swabs. International Journal of Infectious Diseases, 2021, 105, 753-755.	3.3	89
10	Occurrence of severe gastroenteritis in pups after canine parvovirus vaccine administration: A clinical and laboratory diagnostic dilemma. Vaccine, 2007, 25, 1161-1166.	3.8	87
11	Modifications in the Polymerase Genes of a Swine-Like Triple-Reassortant Influenza Virus To Generate Live Attenuated Vaccines against 2009 Pandemic H1N1 Viruses. Journal of Virology, 2011, 85, 456-469.	3.4	85
12	Efficacy of inactivated swine influenza virus vaccines against the 2009 A/H1N1 influenza virus in pigs. Vaccine, 2010, 28, 2782-2787.	3.8	82
13	Infectious canine hepatitis: An "old―disease reemerging in Italy. Research in Veterinary Science, 2007, 83, 269-273.	1.9	75
14	Gain, Preservation, and Loss of a Group 1a Coronavirus Accessory Glycoprotein. Journal of Virology, 2008, 82, 10312-10317.	3.4	73
15	Arctic Lineage-Canine Distemper Virus as a Cause of Death in Apennine Wolves (Canis lupus) in Italy. PLoS ONE, 2014, 9, e82356.	2.5	68
16	Vaccination with NS1-truncated H3N2 swine influenza virus primes T cells and confers cross-protection against an H1N1 heterosubtypic challenge in pigs. Vaccine, 2012, 30, 280-288.	3.8	61
17	Live attenuated influenza A virus vaccine protects against A(H1N1)pdm09 heterologous challenge without vaccine associated enhanced respiratory disease. Virology, 2014, 471-473, 93-104.	2.4	60
18	One after the other: A novel Bluetongue virus strain related to Toggenburg virus detected in the Piedmont region (North-western Italy), extends the panel of novel atypical BTV strains. Transboundary and Emerging Diseases, 2018, 65, 370-374.	3.0	57

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19	Analysis of bluetongue serotype 3 spread in Tunisia and discovery of a novel strain related to the bluetongue virus isolated from a commercial sheep pox vaccine. Infection, Genetics and Evolution, 2018, 59, 63-71.	2.3	56
20	Circovirus in domestic and wild carnivores: An important opportunistic agent?. Virology, 2016, 490, 69-74.	2.4	55
21	Bluetongue Serotype 2 and 9 Modified Live Vaccine Viruses as Causative Agents of Abortion in Livestock: A Retrospective Analysis in Italy. Transboundary and Emerging Diseases, 2014, 61, 69-74.	3.0	53
22	Novel coronavirus (SARS-CoV-2) epidemic: a veterinary perspective. Veterinaria Italiana, 2020, 56, 5-10.	0.5	53
23	Serum Neutralization Assay Can Efficiently Replace Plaque Reduction Neutralization Test for Detection and Quantitation of West Nile Virus Antibodies in Human and Animal Serum Samples. Vaccine Journal, 2014, 21, 1460-1462.	3.1	48
24	Lethal distemper in badgers (Meles meles) following epidemic in dogs and wolves. Infection, Genetics and Evolution, 2016, 46, 130-137.	2.3	48
25	SARS-CoV-2 replicates in respiratory ex vivo organ cultures of domestic ruminant species. Veterinary Microbiology, 2021, 252, 108933.	1.9	48
26	Experimental infection of dogs with a novel strain of canine coronavirus causing systemic disease and lymphopenia. Veterinary Microbiology, 2008, 128, 253-260.	1.9	47
27	Duplex Real-Time Polymerase Chain Reaction for Simultaneous Detection and Quantification of $\langle i \rangle$ Anaplasma Marginale $\langle i \rangle$ and $\langle i \rangle$ Anaplasma Centrale $\langle i \rangle$. Journal of Veterinary Diagnostic Investigation, 2008, 20, 606-611.	1.1	46
28	A "One-Health―approach for diagnosis and molecular characterization of SARS-CoV-2 in Italy. One Health, 2020, 10, 100135.	3.4	46
29	Molecular characterization of a canine respiratory coronavirus strain detected in Italy. Virus Research, 2009, 141, 96-100.	2.2	45
30	Contemporary Epidemiology of North American Lineage Triple Reassortant Influenza A Viruses in Pigs. Current Topics in Microbiology and Immunology, 2011, 370, 113-131.	1.1	45
31	Genome characterization of feline morbillivirus from Italy. Journal of Virological Methods, 2016, 234, 160-163.	2.1	45
32	Bluetongue virus serotypes 1 and 4 in Sardinia during autumn 2012: New incursions or re-infection with old strains?. Infection, Genetics and Evolution, 2013, 19, 81-87.	2.3	43
33	Genomic Epidemiology of the First Wave of SARS-CoV-2 in Italy. Viruses, 2020, 12, 1438.	3.3	39
34	Biological and genetic analysis of a bovine-like coronavirus isolated from water buffalo (Bubalus) Tj ETQq0 0 0 r	gBT/Qverl	ock ₃ 10 Tf 50 1
35	One-step real-time RT-PCR for pandemic influenza A virus (H1N1) 2009 matrix gene detection in swine samples. Journal of Virological Methods, 2010, 164, 83-87.	2.1	36
36	Proficiency Testing of Virus Diagnostics Based on Bioinformatics Analysis of Simulated <i>In Silico</i> High-Throughput Sequencing Data Sets. Journal of Clinical Microbiology, 2019, 57, .	3.9	34

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37	Possible Human-to-Dog Transmission of SARS-CoV-2, Italy, 2020. Emerging Infectious Diseases, 2021, 27, 1981-1984.	4.3	34
38	Restored PB1-F2 in the 2009 Pandemic H1N1 Influenza Virus Has Minimal Effects in Swine. Journal of Virology, 2012, 86, 5523-5532.	3.4	33
39	First report of feline morbillivirus in Europe. Veterinaria Italiana, 2015, 51, 235-7.	0.5	33
40	Emergence and Spread of SARS-CoV-2 Lineages B.1.1.7 and P.1 in Italy. Viruses, 2021, 13, 794.	3.3	32
41	A new member of the Pteropine Orthoreovirus species isolated from fruit bats imported to Italy. Infection, Genetics and Evolution, 2015, 30, 55-58.	2.3	31
42	Tissue distribution of the antigenic variants of canine parvovirus type 2 in dogs. Veterinary Microbiology, 2007, 121, 39-44.	1.9	30
43	Longâ€term persistence of neutralizing SARSâ€CoVâ€2 antibodies in pets. Transboundary and Emerging Diseases, 2022, 69, 3073-3076.	3.0	30
44	Further circulation of West Nile and Usutu viruses in wild birds in Italy. Infection, Genetics and Evolution, 2015, 32, 292-297.	2.3	29
45	A novel Bluetongue virus serotype 3 strain in Tunisia, November 2016. Transboundary and Emerging Diseases, 2017, 64, 709-715.	3.0	29
46	Transplacental transmission of field and rescued strains of BTV-2 and BTV-8 in experimentally infected sheep. Veterinary Research, 2013, 44, 75.	3.0	27
47	Western Bluetongue virusÂserotype 3 in Sardinia, diagnosis and characterization. Transboundary and Emerging Diseases, 2019, 66, 1426-1431.	3.0	25
48	Absence of 2009 Pandemic H1N1 Influenza A Virus in Fresh Pork. PLoS ONE, 2009, 4, e8367.	2.5	23
49	West Nile virus lineage 2 in Sardinian wild birds in 2012: a further threat to public health. Epidemiology and Infection, 2013, 141, 2313-2316.	2.1	22
50	SARS-CoV-2 RNA Persistence in Naso-Pharyngeal Swabs. Microorganisms, 2020, 8, 1124.	3.6	22
51	Diagnosis and characterization of canine distemper virus through sequencing by MinION nanopore technology. Scientific Reports, 2019, 9, 1714.	3.3	21
52	Multiple detection and spread of novel strains of the SARS-CoV-2 B.1.177 (B.1.177.75) lineage that test negative by a commercially available nucleocapsid gene real-time RT-PCR. Emerging Microbes and Infections, 2021, 10, 1148-1155.	6.5	21
53	Bluetongue virus serotype 3 in Western Sicily, November 2017. Veterinaria Italiana, 2017, 53, 273-275.	0.5	20
54	Caprine herpesvirus 1 vaccine with the LTK63 mutant as a mucosal adjuvant induces strong protection against genital infection in goats. Vaccine, 2007, 25, 7927-7930.	3.8	19

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55	Molecular epidemiology of bluetongue virus serotype 1 circulating in Italy and its connection with northern Africa. Infection, Genetics and Evolution, 2014, 28, 144-149.	2.3	19
56	Epidemiology, pathological aspects and genome heterogeneity of feline morbillivirus in Italy. Veterinary Microbiology, 2020, 240, 108484.	1.9	19
57	Reptile-associated Salmonellosis in Man, Italy. Emerging Infectious Diseases, 2006, 12, 358-329.	4.3	18
58	A real-time RT-PCR assay for molecular identification and quantitation of feline morbillivirus RNA from biological specimens. Journal of Virological Methods, 2018, 258, 24-28.	2.1	18
59	Influenza A Virus PB1-F2 Protein Expression Is Regulated in a Strain-Specific Manner by Sequences Located Downstream of the PB1-F2 Initiation Codon. Journal of Virology, 2013, 87, 10687-10699.	3.4	16
60	Bluetongue virus surveillance in the Islamic Republic of Mauritania: Is serotype 26 circulating among cattle and dromedaries?. Infection, Genetics and Evolution, 2016, 40, 109-112.	2.3	16
61	Genome Sequence of Canine Adenovirus Type 1 Isolated from a Wolf ($<$ i>Canis lupus $<$ $ $ i> $>$) in Southern Italy. Genome Announcements, 2017, 5, .	0.8	15
62	Whole genome sequence analysis of the arctic-lineage strain responsible for distemper in Italian wolves and dogs through a fast and robust next generation sequencing protocol. Journal of Virological Methods, 2014, 202, 64-68.	2.1	14
63	Experimental infection of rock pigeons (<i>Columba livia</i>) with three West Nile virus lineage 1 strains isolated in Italy between 2009 and 2012. Epidemiology and Infection, 2016, 144, 1301-1311.	2.1	14
64	Experimental Usutu Virus Infection in Domestic Canaries Serinus canaria. Viruses, 2020, 12, 164.	3.3	14
65	A real-time PCR assay for detection and quantification of Mycoplasma agalactiae DNA. Journal of Applied Microbiology, 2007, 103, 918-923.	3.1	13
66	Comparative virulence of wild-type H1N1pdm09 influenza A isolates in swine. Veterinary Microbiology, 2015, 176, 40-49.	1.9	13
67	Identification and genetic characterization of bovine enterovirus by combination of two next generation sequencing platforms. Journal of Virological Methods, 2018, 260, 21-25.	2.1	13
68	Early Renal Involvement in Cats with Natural Feline Morbillivirus Infection. Animals, 2020, 10, 828.	2.3	13
69	Isolation and genome sequences of two Feline Morbillivirus genotype 1 strains from Italy. Veterinaria Italiana, $2019,55,179-182.$	0.5	12
70	New species of the genus Culicoides (Diptera Ceratopogonidae) for Tunisia, with detection of Bluetongue viruses in vectors. Veterinaria Italiana, 2017, 53, 357-366.	0.5	12
71	SARS-CoV-2 surveillance in Italy through phylogenomic inferences based on Hamming distances derived from pan-SNPs, -MNPs and -InDels. BMC Genomics, 2021, 22, 782.	2.8	12
72	Canine distemper and endangered wildlife: Is it time for mandatory vaccination of dogs?. Vaccine, 2015, 33, 6519.	3.8	11

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73	Exploiting serological data to understand the epidemiology of bluetongue virus serotypes circulating in Libya. Veterinary Medicine and Science, 2019, 5, 79-86.	1.6	11
74	A COVID-19 Hotspot Area: Activities and Epidemiological Findings. Microorganisms, 2020, 8, 1711.	3.6	10
75	Old diseases for new nightmares: distemper strikes back in Italy. Veterinaria Italiana, 2014, 50, 151-4.	0.5	10
76	Competitive enzyme-linked immunosorbent assay using baculovirus-expressed VP7 for detection of epizootic haemorrhagic disease virus (EHDV) antibodies. Journal of Virological Methods, 2017, 248, 212-216.	2.1	9
77	Outbreak of porcine epidemic diarrhoea virus (<scp>PEDV</scp>) in Abruzzi region, centralâ€ltaly. Veterinary Medicine and Science, 2018, 4, 73-79.	1.6	9
78	Specific capture and whole-genome phylogeography of Dolphin morbillivirus. Scientific Reports, 2020, 10, 20831.	3.3	9
79	Neutralization of SARS-CoV-2 Variants by Serum from BNT162b2 Vaccine Recipients. Viruses, 2021, 13, 2011.	3.3	9
80	Complete Genome Sequence of Bluetongue Virus Serotype 1 Circulating in Italy, Obtained through a Fast Next-Generation Sequencing Protocol. Genome Announcements, 2014, 2, .	0.8	8
81	Usutu Virus Infection of Embryonated Chicken Eggs and a Chicken Embryo-Derived Primary Cell Line. Viruses, 2020, 12, 531.	3.3	8
82	Feline Morbillivirus in Southern Italy: Epidemiology, Clinico-Pathological Features and Phylogenetic Analysis in Cats. Viruses, 2021, 13, 1449.	3.3	8
83	Epizootic haemorrhagic disease virus circulation in Tunisia. Veterinaria Italiana, 2018, 54, 87-90.	0.5	8
84	Innocuity of a commercial live attenuated vaccine for epizootic hemorrhagic disease virus serotype 2 in late-term pregnant cows. Vaccine, 2016, 34, 1430-1435.	3.8	7
85	Antigenic relationship among zoonotic flaviviruses from Italy. Infection, Genetics and Evolution, 2019, 68, 91-97.	2.3	7
86	Detection of Astrovirus in a Cow with Neurological Signs by Nanopore Technology, Italy. Viruses, 2020, 12, 530.	3.3	7
87	Bluetongue Serotype 3 in Israel 2013–2018: Clinical Manifestations of the Disease and Molecular Characterization of Israeli Strains. Frontiers in Veterinary Science, 2020, 7, 112.	2.2	7
88	Novel SARS-CoV-2 Variants in Italy: The Role of Veterinary Public Health Institutes. Viruses, 2021, 13, 549.	3.3	7
89	Feline Morbillivirus Infection in Domestic Cats: What Have We Learned So Far?. Viruses, 2021, 13, 683.	3.3	7
90	Epidemiological Significance of SARS-CoV-2 RNA Dynamic in Naso-Pharyngeal Swabs. Microorganisms, 2021, 9, 1264.	3.6	7

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91	First influenza D virus full-genome sequence retrieved from livestock in Namibia, Africa. Acta Tropica, 2022, 232, 106482.	2.0	7
92	Bluetongue Virus in Lebanon. Transboundary and Emerging Diseases, 2013, 60, 390-394.	3.0	6
93	Complete Genome Sequence Analysis of a Reassortant Strain of Bluetongue Virus Serotype 16 from Italy. Genome Announcements, 2013, 1, .	0.8	6
94	Whole-Genome Sequence of a Suid Herpesvirus-1 Strain Isolated from the Brain of a Hunting Dog in Italy. Genome Announcements, 2016, 4, .	0.8	6
95	SARS-CoV-2 Pandemic: Not the First, Not the Last. Microorganisms, 2021, 9, 433.	3.6	6
96	Oral and Poster Manuscripts. Influenza and Other Respiratory Viruses, 2011, 5, 54-442.	3.4	5
97	Efficacy of vaccination for bluetongue virus serotype 8 performed shortly before challenge and implications for animal trade. Preventive Veterinary Medicine, 2017, 136, 49-55.	1.9	5
98	Genome Sequences of Three SARS-CoV-2 P.1 Strains Identified from Patients Returning from Brazil to Italy. Microbiology Resource Announcements, 2021, 10, .	0.6	5
99	SARS-CoV-2 Delta VOC in a Paucisymptomatic Dog, Italy. Pathogens, 2022, 11, 514.	2.8	5
100	Reemergence of an atypical bluetongue virus strain in goats, Sardinia, Italy. Research in Veterinary Science, 2022, 151, 36-41.	1.9	5
101	Polymorphisms in the haemagglutinin gene influenced the viral shedding of pandemic 2009 influenza virus in swine. Journal of General Virology, 2014, 95, 2618-2626.	2.9	4
102	Replication kinetics and cellular tropism of emerging reoviruses in sheep and swine respiratory ex vivo organ cultures. Veterinary Microbiology, 2019, 234, 119-127.	1.9	4
103	Molecular typing of Bluetongue virus using the nCounter® analysis system platform. Journal of Virological Methods, 2019, 269, 64-69.	2.1	4
104	Erasing the Invisible Line to Empower the Pandemic Response. Viruses, 2021, 13, 348.	3.3	4
105	Insights into SARS-CoV-2, the Coronavirus Underlying COVID-19: Recent Genomic Data and the Development of Reverse Genetics Systems. Journal of General Virology, 2020, 101, 1021-1024.	2.9	4
106	Transplacental transmission of the Italian Bluetongue virus serotype 2 in sheep. Veterinaria Italiana, 2019, 55, 131-141.	0.5	4
107	Epidemiological and genomic findings of the first documented Italian outbreak of SARS-CoV-2 Alpha variant of concern. Epidemics, 2022, 39, 100578.	3.0	4
108	Comparison of Human-Like H1 (-Cluster) Influenza A Viruses in the Swine Host. Influenza Research and Treatment, 2012, 2012, 1-7.	1.5	3

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109	Genome Sequencing of a Camelpox Vaccine Reveals Close Similarity to Modified Vaccinia virus Ankara (MVA). Viruses, 2020, 12, 786.	3.3	3
110	Development of a Digital RT-PCR Method for Absolute Quantification of Bluetongue Virus in Field Samples. Frontiers in Veterinary Science, 2020, 7, 170.	2.2	3
111	Whole-Genome Sequences of SARS-CoV-2 Lineage B.1.525 Strains (Variant η) Detected from Patients in the Abruzzo Region (Central Italy) during Spring 2021. Microbiology Resource Announcements, 2021, 10, e0061821.	0.6	2
112	The envelope protein of Usutu virus attenuates West Nile virus virulence in immunocompetent mice. Veterinary Microbiology, 2021, 263, 109262.	1.9	2
113	OIEBTLABNET: the web-based network of the OIE Bluetongue Reference Laboratories. Veterinaria Italiana, 2016, 52, 187-193.	0.5	1
114	Bluetongue Disease., 2019,, 305-322.		0
115	Development of a serum neutralization assay to detect Pteropine Orthoreovirus Indonesia/2010 neutralizing antibodies. Veterinaria Italiana, 2018, 54, 161-164.	0.5	0
116	Full Genome Characterization of Respiratory Syncytial Virus Causing a Fatal Infection in an Immunocompromised Patient in Tunisia. Pathogens, 2022, 11, 758.	2.8	O