## Mattia Cecchinato

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Technological quality, mineral profile, and sensory attributes of broiler chicken breasts affected by White Striping and Wooden Breast myopathies. Poultry Science, 2016, 95, 2707-2714.	1.5	107
2	Effect of "Wooden Breast" appearance on poultry meat quality, histological traits, and lesions characterization. Czech Journal of Animal Science, 2017, 62, 51-57.	0.5	85
3	Porcine circovirus type 2 (PCV2) evolution before and after the vaccination introduction: A large scale epidemiological study. Scientific Reports, 2016, 6, 39458.	1.6	70
4	Demonstration of loss of attenuation and extended field persistence of a live avian metapneumovirus vaccine. Vaccine, 2006, 24, 6476-6482.	1.7	68
5	Avian metapneumovirus (AMPV) attachment protein involvement in probable virus evolution concurrent with mass live vaccine introduction. Veterinary Microbiology, 2010, 146, 24-34.	0.8	54
6	First report of wild boar susceptibility to Porcine circovirus type 3: High prevalence in the Colli Euganei Regional Park (Italy) in the absence of clinical signs. Transboundary and Emerging Diseases, 2018, 65, 957-962.	1.3	52
7	Think globally, act locally: Phylodynamic reconstruction of infectious bronchitis virus (IBV) QX genotype (Gl-19 lineage) reveals different population dynamics and spreading patterns when evaluated on different epidemiological scales. PLoS ONE, 2017, 12, e0184401.	1.1	51
8	A novel variant of the infectious bronchitis virus resulting from recombination events in Italy and Spain. Avian Pathology, 2017, 46, 28-35.	0.8	46
9	Field avian Metapneumovirus evolution avoiding vaccine induced immunity. Vaccine, 2010, 28, 916-921.	1.7	45
10	A molecular epidemiology study based on VP2 gene sequences reveals that a new genotype of infectious bursal disease virus is dominantly prevalent in Italy. Avian Pathology, 2016, 45, 458-464.	0.8	41
11	Infectious Bronchitis Virus Evolution, Diagnosis and Control. Veterinary Sciences, 2020, 7, 79.	0.6	41
12	Evolution of infectious bronchitis virus in the field after homologous vaccination introduction. Veterinary Research, 2019, 50, 92.	1.1	40
13	Effect of different vaccination strategies on IBV QX population dynamics and clinical outbreaks. Vaccine, 2016, 34, 5670-5676.	1.7	38
14	Development and validation of direct PCR and quantitative PCR assays for the rapid, sensitive, and economical detection of porcine circovirus 3. Journal of Veterinary Diagnostic Investigation, 2018, 30, 538-544.	0.5	37
15	In vitro antiviral activity of chestnut and quebracho woods extracts against avian reovirus and metapneumovirus. Research in Veterinary Science, 2009, 87, 482-487.	0.9	36
16	Canine parvovirus type 2 (CPV-2) and Feline panleukopenia virus (FPV) codon bias analysis reveals a progressive adaptation to the new niche after the host jump. Molecular Phylogenetics and Evolution, 2017, 114, 82-92.	1.2	34
17	A turkey rhinotracheitis outbreak caused by the environmental spread of a vaccine-derived avian metapneumovirus. Avian Pathology, 2011, 40, 525-530.	0.8	33
18	Continued use of IBV 793B vaccine needs reassessment after its withdrawal led to the genotype's disappearance. Vaccine, 2014, 32, 6765-6767.	1.7	33

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19	West Nile virus circulation in Veneto region in 2008–2009. Epidemiology and Infection, 2011, 139, 818-825.	1.0	30
20	Development of a real-time RT-PCR assay for the simultaneous identification, quantitation and differentiation of avian metapneumovirus subtypes A and B. Avian Pathology, 2013, 42, 283-289.	0.8	27
21	Molecular characterization of the meq gene of Marek's disease viruses detected in unvaccinated backyard chickens reveals the circulation of low- and high-virulence strains. Poultry Science, 2019, 98, 3130-3137.	1.5	26
22	Molecular investigation of a full-length genome of a Q1-like IBV strain isolated in Italy in 2013. Virus Research, 2015, 210, 77-80.	1.1	23
23	Molecular epidemiology of infectious bronchitis virus and avian metapneumovirus in Greece. Poultry Science, 2019, 98, 5374-5384.	1.5	22
24	Avian Metapneumovirus subtype B around Europe: a phylodynamic reconstruction. Veterinary Research, 2020, 51, 88.	1.1	22
25	The effects of control measures on the economic burden associated with epidemics of avian influenza in Italy. Poultry Science, 2010, 89, 1115-1121.	1.5	21
26	The analysis of genome composition and codon bias reveals distinctive patterns between avian and mammalian circoviruses which suggest a potential recombinant origin for Porcine circovirus 3. PLoS ONE, 2018, 13, e0199950.	1.1	21
27	Italian Field Survey Reveals a High Diffusion of Avian Metapneumovirus Subtype B in Layers and Weaknesses in the Vaccination Strategy Applied. Avian Diseases, 2012, 56, 720-724.	0.4	20
28	Observation of high recombination occurrence of Porcine Reproductive and Respiratory Syndrome Virus in field condition. Virus Research, 2014, 194, 159-166.	1.1	19
29	Molecular insight into Italian canine parvovirus heterogeneity and comparison with the worldwide scenario. Infection, Genetics and Evolution, 2018, 66, 171-179.	1.0	19
30	Vaccine or field strains: the jigsaw pattern of infectious bronchitis virus molecular epidemiology in Poland. Poultry Science, 2019, 98, 6388-6392.	1.5	19
31	Free to Circulate: An Update on the Epidemiological Dynamics of Porcine Circovirus 2 (PCV-2) in Italy Reveals the Role of Local Spreading, Wild Populations, and Foreign Countries. Pathogens, 2020, 9, 221.	1.2	19
32	An Assessment of the Level of Protection Against Colibacillosis Conferred by Several Autogenous and/or Commercial Vaccination Programs in Conventional Pullets upon Experimental Challenge. Veterinary Sciences, 2020, 7, 80.	0.6	19
33	Avian Metapneumovirus circulation in Italian broiler farms. Poultry Science, 2018, 97, 503-509.	1.5	18
34	Avian metapneumoviruses expressing Infectious Bronchitis virus genes are stable and induce protection. Vaccine, 2013, 31, 2565-2571.	1.7	17
35	Molecular characterization of whole genome sequence of infectious bronchitis virus 624I genotype confirms the close relationship with Q1 genotype. Transboundary and Emerging Diseases, 2019, 66, 207-216.	1.3	17
36	Rapid detection of subtype B avian metapneumoviruses using RT-PCR restriction endonuclease digestion indicates field circulation of vaccine-derived viruses in older turkeys. Avian Pathology, 2014, 43, 51-56.	0.8	16

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37	Genome sequence analysis of a distinctive Italian infectious bursal disease virus. Poultry Science, 2017, 96, 4370-4377.	1.5	16
38	Avian Pneumovirus infection in turkey and broiler farms in Italy: a virological, molecular and serological field survey. Italian Journal of Animal Science, 2004, 3, 287-292.	0.8	15
39	First Report of Avian Metapneumovirus Subtype B Field Strain in a Romanian Broiler Flock During an Outbreak of Respiratory Disease. Avian Diseases, 2017, 61, 250.	0.4	15
40	Bovine Coronavirus: Variability, Evolution, and Dispersal Patterns of a No Longer Neglected Betacoronavirus. Viruses, 2020, 12, 1285.	1.5	15
41	Phylodynamic analysis and evaluation of the balance between anthropic and environmental factors affecting IBV spreading among Italian poultry farms. Scientific Reports, 2020, 10, 7289.	1.6	15
42	<i>Gammacoronavirus</i> and <i>Deltacoronavirus</i> in Quail. Avian Diseases, 2016, 60, 656-661.	0.4	14
43	Gamma and Deltacoronaviruses in quail and pheasants from Northern Italy. Poultry Science, 2017, 96, 717-722.	1.5	14
44	Vaccination reduced the incidence of outbreaks of low pathogenicity avian influenza in northern Italy. Vaccine, 2009, 27, 3655-3661.	1.7	13
45	Phylodynamic analysis of porcine reproductive and respiratory syndrome virus (PRRSV) in Italy: Action of selective pressures and interactions between different clades. Infection, Genetics and Evolution, 2015, 31, 149-157.	1.0	13
46	Canine parvovirus (CPV) phylogeny is associated with disease severity. Scientific Reports, 2019, 9, 11266.	1.6	13
47	Detection and molecular characterization of a new genotype of infectious bursal disease virus in Portugal. Avian Pathology, 2022, 51, 97-105.	0.8	13
48	Low Pathogenicity Avian Influenza in Italy During 2007 and 2008: Epidemiology and Control. Avian Diseases, 2010, 54, 323-328.	0.4	12
49	Epidemiology and Control of Low Pathogenicity Avian Influenza Infections in Rural Poultry in Italy. Avian Diseases, 2011, 55, 13-20.	0.4	12
50	Impact of coccidiosis control program and feeding plan on white striping prevalence and severity degree on broiler breast fillets evaluated at three growing ages. Poultry Science, 2015, 94, 2114-2123.	1.5	12
51	Evaluation of 793/B-like and Mass-like vaccine strain kinetics in experimental and field conditions by real-time RT-PCR quantification. Poultry Science, 2018, 97, 303-312.	1.5	12
52	GI-16 lineage (624/I or Q1), there and back again: The history of one of the major threats for poultry farming of our era. PLoS ONE, 2018, 13, e0203513.	1.1	12
53	First evidence of avian metapneumovirus subtype A infection in turkeys in Egypt. Tropical Animal Health and Production, 2014, 46, 1093-1097.	0.5	11
54	Effect of genome composition and codon bias on infectious bronchitis virus evolution and adaptation to target tissues. BMC Genomics, 2021, 22, 244.	1.2	11

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55	Outbreak of <i>Eimeria kofoidi</i> and <i>E. legionensis</i> coccidiosis in red-legged partridges <i>(Alectoris rufa)</i> . Italian Journal of Animal Science, 2006, 5, 318-320.	0.8	10
56	A Sensitive, Reproducible, and Economic Real-Time Reverse Transcription PCR Detecting Avian Metapneumovirus Subtypes A and B. Avian Diseases, 2014, 58, 216-222.	0.4	10
57	First Identification and Molecular Characterization of Avian metapneumovirus Subtype B from Chickens in Greece. Avian Diseases, 2017, 61, 409.	0.4	10
58	Phylogeny and evolution of infectious bursal disease virus circulating in Turkish broiler flocks. Poultry Science, 2019, 98, 1976-1984.	1.5	10
59	Comparative in vivo pathogenicity study of an ITA genotype isolate (G6) of infectious bursal disease virus. Transboundary and Emerging Diseases, 2020, 67, 1025-1031.	1.3	10
60	What is new on molecular characteristics of Avian metapneumovirus strains circulating in Europe?. Transboundary and Emerging Diseases, 2021, 68, 1314-1322.	1.3	10
61	Porcine Gammaherpesviruses in Italian Commercial Swine Population: Frequent but Harmless. Pathogens, 2021, 10, 47.	1.2	10
62	Genetic Insights into Feline Parvovirus: Evaluation of Viral Evolutionary Patterns and Association between Phylogeny and Clinical Variables. Viruses, 2021, 13, 1033.	1.5	9
63	First Molecular Characterization of Avian Metapneumovirus (aMPV) in Turkish Broiler Flocks. Avian Diseases, 2018, 62, 425.	0.4	9
64	Molecular Characterization of Avian Metapneumovirus from Guinea Fowls (Numida meleagridis). Pakistan Veterinary Journal, 2018, 38, 419-423.	0.5	9
65	First detection of avian metapneumovirus subtype C Eurasian lineage in a Eurasian wigeon ( <i>Mareca) Tj ETQq1 viral epidemiology. Avian Pathology, 2022, 51, 283-290.</i>	1 0.7843 0.8	14 rgBT /Ove 9
66	Identification of IBV QX vaccine markers : Should vaccine acceptance by authorities require similar identifications for all live IBV vaccines?. Vaccine, 2017, 35, 5531-5534.	1.7	8
67	Molecular Epidemiology and Genotyping of Infectious Bronchitis Virus and Avian Metapneumovirus in Backyard and Commercial Chickens in Jimma Zone, Southwestern Ethiopia. Veterinary Sciences, 2020, 7, 187.	0.6	8
68	Phylodynamic and phylogeographic reconstruction of porcine reproductive and respiratory syndrome virus (PRRSV) in Europe: Patterns and determinants. Transboundary and Emerging Diseases, 2022, 69, .	1.3	8
69	Infectious bronchitis virus gel vaccination: evaluation of Mass-like (B-48) and 793/B-like (1/96) vaccine kinetics after combined administration at 1 day of age. Poultry Science, 2018, 97, 3501-3509.	1.5	7
70	Molecular epidemiology of fowl adenoviruses in Greece. Poultry Science, 2020, 99, 5983-5990.	1.5	7
71	Research Note: Detection of Avian metapneumovirus subgroup C specific antibodies in a mallard flock in Italy. Poultry Science, 2021, 100, 101186.	1.5	7
72	Design, Validation, and Absolute Sensitivity of a Novel Test for the Molecular Detection of Avian Pneumovirus. Journal of Veterinary Diagnostic Investigation, 2004, 16, 582-585.	0.5	6

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73	Inoculation of specific pathogen-free chickens with an infectious bursal disease virus of the ITA genotype (G6) leads to a high and persistent viral load in lymphoid tissues and to a delayed antiviral response. Veterinary Microbiology, 2019, 235, 136-142.	0.8	6
74	Phylodynamic and Recombination Analyses of Avian Infectious Bronchitis GI-23 Reveal a Widespread Recombinant Cluster and New Among-Countries Linkages. Animals, 2021, 11, 3182.	1.0	6
75	Low pathogenic avian influenza in Italy. Veterinary Record, 2008, 162, 64-64.	0.2	5
76	No good vaccination quality without good control: the positive impact of a hatchery vaccination service program. Poultry Science, 2020, 99, 2976-2982.	1.5	5
77	Evaluation of unintended 1/96 infectious bronchitis vaccine transmission in broilers after direct contact with vaccinated ones. Veterinarni Medicina, 2018, 63, 287-291.	0.2	4
78	Two similar commercial live attenuated AMPV vaccines prepared by random passage of the identical field isolate, have unrelated sequences. Vaccine, 2019, 37, 2765-2767.	1.7	4
79	Comparison and validation of different models and variable selection methods for predicting survival after canine parvovirus infection. Veterinary Record, 2020, 187, e76.	0.2	4
80	A comparison of AMPV subtypes A and B full genomes, gene transcripts and proteins led to reverse-genetics systems rescuing both subtypes. Journal of General Virology, 2016, 97, 1324-1332.	1.3	4
81	Investigation of Serotype Prevalence of Escherichia coli Strains Isolated from Layer Poultry in Greece and Interactions with Other Infectious Agents. Veterinary Sciences, 2022, 9, 152.	0.6	4
82	Infectious Bronchitis Hatchery Vaccination: Comparison between Traditional Spray Administration and a Newly Developed Gel Delivery System in Field Conditions. Veterinary Sciences, 2021, 8, 145.	0.6	3
83	Impact of viral features, host jumps and phylogeography on the rapid evolution of Aleutian mink disease virus (AMDV). Scientific Reports, 2021, 11, 16464.	1.6	3
84	Do modified live virus vaccines against bovine viral diarrhea induce fetal cross-protection against HoBi-like Pestivirus?. Veterinary Microbiology, 2021, 260, 109178.	0.8	3
85	Reconciling individual and population levels of porcine reproductive and respiratory syndrome virus evolution. Virologica Sinica, 2017, 32, 342-345.	1.2	2
86	Effect of assay choice, viral concentration and operator interpretation on infectious bronchitis virus detection and characterization. Avian Pathology, 2021, 50, 357-365.	0.8	2
87	Detection and Molecular Characterization of a Novel Species of Circovirus in a Tawny Owl (Strix) Tj ETQq1 1 0	784314 rgB 1.0	T /Overlock
88	Molecular Survey and Identification of Campylobacter spp. in Layer Farms in Central Ethiopia. Tropical Medicine and Infectious Disease, 2022, 7, 31.	0.9	2
89	An Electronic Learning Course on Avian Influenza in Italy (2008). Avian Diseases, 2010, 54, 784-786.	0.4	1
90	Impact of Rotten Eggs on Hatchery Performances: A Multicentric Study. Animals, 2020, 10, 1725.	1.0	1

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91	An Outbreak of Blindness Due to Retinopathy in Nine Flocks of Guinea Fowl. Avian Diseases, 2014, 58, 337-339.	0.4	0
92	Molecular Survey of Viral Poultry Diseases with an Indirect Public Health Significance in Central Ethiopia. Animals, 2021, 11, 3564.	1.0	0