

Claudio L Afonso

List of Publications by Year in descending order

Source: [//exaly.com/author-pdf/334640/publications.pdf](https://exaly.com/author-pdf/334640/publications.pdf)

Version: 2024-02-01

128
papers

7,112
citations

65103

42
h-index

63582

80
g-index

132
all docs

132
docs citations

132
times ranked

4918
citing authors

#	ARTICLE	IF	CITATIONS
1	Taxonomy of the order Mononegavirales: update 2016. <i>Archives of Virology</i> , 2016, 161, 2351-2360.	1.9	416
2	Newcastle disease: Evolution of genotypes and the related diagnostic challenges. <i>Infection, Genetics and Evolution</i> , 2010, 10, 26-35.	2.3	337
3	Genetic diversity of avian paramyxovirus type 1: Proposal for a unified nomenclature and classification system of Newcastle disease virus genotypes. <i>Infection, Genetics and Evolution</i> , 2012, 12, 1770-1779.	2.3	329
4	The Genome of Fowlpox Virus. <i>Journal of Virology</i> , 2000, 74, 3815-3831.	3.5	297
5	Newcastle disease vaccines – A solved problem or a continuous challenge?. <i>Veterinary Microbiology</i> , 2017, 206, 126-136.	1.9	251
6	Immune responses of poultry to Newcastle disease virus. <i>Developmental and Comparative Immunology</i> , 2013, 41, 447-453.	2.3	250
7	Updated unified phylogenetic classification system and revised nomenclature for Newcastle disease virus. <i>Infection, Genetics and Evolution</i> , 2019, 74, 103917.	2.3	248
8	Antigenic differences among Newcastle disease virus strains of different genotypes used in vaccine formulation affect viral shedding after a virulent challenge. <i>Vaccine</i> , 2007, 25, 7238-7246.	4.0	236
9	Temporal, geographic, and host distribution of avian paramyxovirus 1 (Newcastle disease virus). <i>Infection, Genetics and Evolution</i> , 2016, 39, 22-34.	2.3	223
10	Phylogenetic Diversity among Low-Virulence Newcastle Disease Viruses from Waterfowl and Shorebirds and Comparison of Genotype Distributions to Those of Poultry-Origin Isolates. <i>Journal of Virology</i> , 2007, 81, 12641-12653.	3.5	204
11	The Genome of <i>Melanoplus sanguinipes</i> Entomopoxvirus. <i>Journal of Virology</i> , 1999, 73, 533-552.	3.5	191
12	African Swine Fever Virus Multigene Family 360 and 530 Genes Affect Host Interferon Response. <i>Journal of Virology</i> , 2004, 78, 1858-1864.	3.5	170
13	Identification of new sub-genotypes of virulent Newcastle disease virus with potential panzootic features. <i>Infection, Genetics and Evolution</i> , 2015, 29, 216-229.	2.3	166
14	Genome Sequence of a Baculovirus Pathogenic for <i>Culex nigripalpus</i> . <i>Journal of Virology</i> , 2001, 75, 11157-11165.	3.5	157
15	Effects of Newcastle disease virus vaccine antibodies on the shedding and transmission of challenge viruses. <i>Developmental and Comparative Immunology</i> , 2013, 41, 505-513.	2.3	152
16	Evolutionary dynamics of Newcastle disease virus. <i>Virology</i> , 2009, 391, 64-72.	2.5	146
17	Virulent Newcastle disease virus elicits a strong innate immune response in chickens. <i>Journal of General Virology</i> , 2011, 92, 931-939.	2.9	127
18	The Genome of Turkey Herpesvirus. <i>Journal of Virology</i> , 2001, 75, 971-978.	3.5	119

#	ARTICLE	IF	CITATIONS
19	Characterization of Class I Newcastle Disease Virus Isolates from Hong Kong Live Bird Markets and Detection Using Real-Time Reverse Transcription-PCR. <i>Journal of Clinical Microbiology</i> , 2007, 45, 1310-1314.	4.4	119
20	Evidence for a New Avian Paramyxovirus Serotype 10 Detected in Rockhopper Penguins from the Falkland Islands. <i>Journal of Virology</i> , 2010, 84, 11496-11504.	3.5	118
21	The Genome of Swinepox Virus. <i>Journal of Virology</i> , 2002, 76, 783-790.	3.5	95
22	Biological and Phylogenetic Characterization of Pigeon Paramyxovirus Serotype 1 Circulating in Wild North American Pigeons and Doves. <i>Journal of Clinical Microbiology</i> , 2008, 46, 3303-3310.	4.4	89
23	Highly Divergent Virulent Isolates of Newcastle Disease Virus from the Dominican Republic Are Members of a New Genotype That May Have Evolved Unnoticed for Over 2 Decades. <i>Journal of Clinical Microbiology</i> , 2013, 51, 508-517.	4.4	88
24	The Genome of Camelpox Virus. <i>Virology</i> , 2002, 295, 1-9.	2.5	86
25	A robust and cost-effective approach to sequence and analyze complete genomes of small RNA viruses. <i>Virology Journal</i> , 2017, 14, 72.	3.6	78
26	Complete Genome and Clinicopathological Characterization of a Virulent Newcastle Disease Virus Isolate from South America. <i>Journal of Clinical Microbiology</i> , 2012, 50, 378-387.	4.4	77
27	Phylogenetic and Biological Characterization of Newcastle Disease Virus Isolates from Pakistan. <i>Journal of Clinical Microbiology</i> , 2010, 48, 1892-1894.	4.4	73
28	Detection of a Broad Range of Class I and II Newcastle Disease Viruses Using a Multiplex Real-Time Reverse Transcription Polymerase Chain Reaction Assay. <i>Journal of Veterinary Diagnostic Investigation</i> , 2008, 20, 414-425.	1.5	70
29	Characterization of Live LaSota Vaccine Strain-Induced Protection in Chickens upon Early Challenge with a Virulent Newcastle Disease Virus of Heterologous Genotype. <i>Avian Diseases</i> , 2012, 56, 464-470.	1.2	67
30	Biological and Phylogenetic Characterization of a Genotype VII Newcastle Disease Virus from Venezuela: Efficacy of Field Vaccination. <i>Journal of Clinical Microbiology</i> , 2012, 50, 1204-1208.	4.4	64
31	Molecular Epidemiology of Newcastle Disease in Mexico and the Potential Spillover of Viruses from Poultry into Wild Bird Species. <i>Applied and Environmental Microbiology</i> , 2013, 79, 4985-4992.	3.2	63
32	Presence of Virulent Newcastle Disease Virus in Vaccinated Chickens in Farms in Pakistan. <i>Journal of Clinical Microbiology</i> , 2015, 53, 1715-1718.	4.4	63
33	Genome of Crocodilepox Virus. <i>Journal of Virology</i> , 2006, 80, 4978-4991.	3.5	62
34	Characterization of Newcastle Disease Viruses Isolated from Cormorant and Gull Species in the United States in 2010. <i>Avian Diseases</i> , 2012, 56, 128-133.	1.2	59
35	African swine fever virus NL gene is not required for virus virulence.. <i>Journal of General Virology</i> , 1998, 79, 2543-2547.	2.9	56
36	Genetic diversity and mutation of avian paramyxovirus serotype 1 (Newcastle disease virus) in wild birds and evidence for intercontinental spread. <i>Archives of Virology</i> , 2013, 158, 2495-2503.	1.9	55

#	ARTICLE	IF	CITATIONS
37	Presence of Vaccine-Derived Newcastle Disease Viruses in Wild Birds. <i>PLoS ONE</i> , 2016, 11, e0162484.	2.5	55
38	Epidemiology, control, and prevention of Newcastle disease in endemic regions: Latin America. <i>Tropical Animal Health and Production</i> , 2019, 51, 1033-1048.	1.4	53
39	Rapid, multiplexed, whole genome and plasmid sequencing of foodborne pathogens using long-read nanopore technology. <i>Scientific Reports</i> , 2019, 9, 16350.	3.4	51
40	Effect of Probe-Site Mismatches on Detection of Virulent Newcastle Disease Viruses Using a Fusion-Gene Real-Time Reverse Transcription Polymerase Chain Reaction Test. <i>Journal of Veterinary Diagnostic Investigation</i> , 2006, 18, 519-528.	1.5	48
41	Expression of interferon gamma by a highly virulent strain of Newcastle disease virus decreases its pathogenicity in chickens. <i>Microbial Pathogenesis</i> , 2013, 61-62, 73-83.	2.9	47
42	Phylogenetic assessment reveals continuous evolution and circulation of pigeon-derived virulent avian avulaviruses 1 in Eastern Europe, Asia, and Africa. <i>BMC Veterinary Research</i> , 2017, 13, 291.	2.0	45
43	Repeated isolation of virulent Newcastle disease viruses in poultry and captive non-poultry avian species in Pakistan from 2011 to 2016. <i>Preventive Veterinary Medicine</i> , 2017, 142, 1-6.	1.9	43
44	Global phylodynamic analysis of avian paramyxovirus-1 provides evidence of inter-host transmission and intercontinental spatial diffusion. <i>BMC Evolutionary Biology</i> , 2019, 19, 108.	3.1	42
45	Newcastle Disease Viruses Causing Recent Outbreaks Worldwide Show Unexpectedly High Genetic Similarity to Historical Virulent Isolates from the 1940s. <i>Journal of Clinical Microbiology</i> , 2016, 54, 1228-1235.	4.4	40
46	Development of an improved vaccine evaluation protocol to compare the efficacy of Newcastle disease vaccines. <i>Biologicals</i> , 2015, 43, 136-145.	1.4	39
47	Evolutionary Changes Affecting Rapid Identification of 2008 Newcastle Disease Viruses Isolated from Double-Crested Cormorants. <i>Journal of Clinical Microbiology</i> , 2010, 48, 2440-2448.	4.4	38
48	Genetic characterization and pathogenesis of the first H9N2 low pathogenic avian influenza viruses isolated from chickens in Kenyan live bird markets. <i>Infection, Genetics and Evolution</i> , 2020, 78, 104074.	2.3	35
49	Biological and Phylogenetic Characterization of Virulent Newcastle Disease Virus Circulating in Mexico. <i>Avian Diseases</i> , 2008, 52, 472-479.	1.2	34
50	Experimental co-infections of domestic ducks with a virulent Newcastle disease virus and low or highly pathogenic avian influenza viruses. <i>Veterinary Microbiology</i> , 2015, 177, 7-17.	1.9	33
51	Virulent Newcastle disease viruses from chicken origin are more pathogenic and transmissible to chickens than viruses normally maintained in wild birds. <i>Veterinary Microbiology</i> , 2019, 235, 25-34.	1.9	33
52	Whole-genome sequencing of genotype VI Newcastle disease viruses from formalin-fixed paraffin-embedded tissues from wild pigeons reveals continuous evolution and previously unrecognized genetic diversity in the U.S.. <i>Virology Journal</i> , 2018, 15, 9.	3.6	32
53	Pathogenicity and transmission of virulent Newcastle disease virus from the 2018–2019 California outbreak and related viruses in young and adult chickens. <i>Virology</i> , 2019, 531, 203-218.	2.5	31
54	Newcastle disease virus fusion and haemagglutinin-neuraminidase proteins contribute to its macrophage host range. <i>Journal of General Virology</i> , 2013, 94, 1189-1194.	2.9	30

#	ARTICLE	IF	CITATIONS
55	International Biological Engagement Programs Facilitate Newcastle Disease Epidemiological Studies. <i>Frontiers in Public Health</i> , 2015, 3, 235.	2.8	30
56	Expression of chicken interleukin-2 by a highly virulent strain of Newcastle disease virus leads to decreased systemic viral load but does not significantly affect mortality in chickens. <i>Virology Journal</i> , 2015, 12, 122.	3.6	30
57	Genomic comparison of Newcastle disease viruses isolated in Nigeria between 2002 and 2015 reveals circulation of highly diverse genotypes and spillover into wild birds. <i>Archives of Virology</i> , 2019, 164, 2031-2047.	1.9	30
58	Wild Bird Surveillance for Avian Paramyxoviruses in the Azov-Black Sea Region of Ukraine (2006 to) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 <i>Microbiology</i> , 2014, 80, 5427-5438.	3.2	28
59	Neuropathogenic Capacity of Lentogenic, Mesogenic, and Velogenic Newcastle Disease Virus Strains in Day-Old Chickens. <i>Veterinary Pathology</i> , 2016, 53, 53-64.	2.0	27
60	H9N2 low pathogenic avian influenza in Pakistan (2012â€“2015). <i>Veterinary Record Open</i> , 2016, 3, e000171.	1.0	26
61	Detection of Fowlpox virus carrying distinct genome segments of Reticuloendotheliosis virus. <i>Virus Research</i> , 2019, 260, 53-59.	2.3	26
62	Rapid virulence prediction and identification of Newcastle disease virus genotypes using third-generation sequencing. <i>Virology Journal</i> , 2018, 15, 179.	3.6	25
63	Virulence during Newcastle Disease Viruses Cross Species Adaptation. <i>Viruses</i> , 2021, 13, 110.	3.4	25
64	Primary chicken tracheal cell culture system for the study of infection with avian respiratory viruses. <i>Avian Pathology</i> , 2008, 37, 25-31.	2.0	24
65	Separate Evolution of Virulent Newcastle Disease Viruses from Mexico and Central America. <i>Journal of Clinical Microbiology</i> , 2014, 52, 1382-1390.	4.4	23
66	Repeated isolation of virulent Newcastle disease viruses of sub-genotype VIIId from backyard chickens in Bulgaria and Ukraine between 2002 and 2013. <i>Archives of Virology</i> , 2016, 161, 3345-3353.	1.9	23
67	Not So Fast on Recombination Analysis of Newcastle Disease Virus. <i>Journal of Virology</i> , 2008, 82, 9303-9303.	3.5	22
68	Previous infection with virulent strains of Newcastle disease virus reduces highly pathogenic avian influenza virus replication, disease, and mortality in chickens. <i>Veterinary Research</i> , 2015, 46, 97.	3.1	22
69	Complete Genome Sequence of an Avian Paramyxovirus Representative of Putative New Serotype 13. <i>Genome Announcements</i> , 2016, 4, .	0.8	22
70	Development, characterization and optimization of a new suspension chicken-induced pluripotent cell line for the production of Newcastle disease vaccine. <i>Biologicals</i> , 2016, 44, 24-32.	1.4	22
71	Effects of Chicken Interferon Gamma on Newcastle Disease Virus Vaccine Immunogenicity. <i>PLoS ONE</i> , 2016, 11, e0159153.	2.5	22
72	Rapid evolution of Mexican H7N3 highly pathogenic avian influenza viruses in poultry. <i>PLoS ONE</i> , 2019, 14, e0222457.	2.5	21

#	ARTICLE	IF	CITATIONS
73	Genome-wide analysis reveals class and gene specific codon usage adaptation in avian paramyxoviruses 1. <i>Infection, Genetics and Evolution</i> , 2017, 50, 28-37.	2.3	19
74	A retrospective study of Newcastle disease in Kenya. <i>Tropical Animal Health and Production</i> , 2020, 52, 699-710.	1.4	19
75	Pathogenesis of New Strains of Newcastle Disease Virus From Israel and Pakistan. <i>Veterinary Pathology</i> , 2016, 53, 792-796.	2.0	18
76	Assessment of contemporary genetic diversity and inter-taxa/inter-region exchange of avian paramyxovirus serotype 1 in wild birds sampled in North America. <i>Virology Journal</i> , 2017, 14, 43.	3.6	18
77	Genome Sequence Variations of Infectious Bronchitis Virus Serotypes From Commercial Chickens in Mexico. <i>Frontiers in Veterinary Science</i> , 0, 9, .	2.3	17
78	Complete Genome Sequence of a Genotype XVII Newcastle Disease Virus, Isolated from an Apparently Healthy Domestic Duck in Nigeria. <i>Genome Announcements</i> , 2016, 4, .	0.8	16
79	MinION sequencing to genotype US strains of infectious laryngotracheitis virus. <i>Avian Pathology</i> , 2019, 48, 255-269.	2.0	16
80	Surveillance and Genetic Characterization of Virulent Newcastle Disease Virus Subgenotype V.3 in Indigenous Chickens from Backyard Poultry Farms and Live Bird Markets in Kenya. <i>Viruses</i> , 2021, 13, 103.	3.4	16
81	Presence of Newcastle disease viruses of sub-genotypes Vc and Vln in backyard chickens and in apparently healthy wild birds from Mexico in 2017. <i>Virus Genes</i> , 2019, 55, 479-489.	1.8	15
82	First Complete Genome Sequence of Currently Circulating Infectious Bronchitis Virus Strain DMV/1639 of the GI-17 Lineage. <i>Microbiology Resource Announcements</i> , 2019, 8, .	2.0	14
83	Pathology and Distribution of Velogenic Viscerotropic Newcastle Disease Virus in the Reproductive System of Vaccinated and Unvaccinated Laying Hens (<i>Gallus gallus domesticus</i>) by Immunohistochemical Labelling. <i>Journal of Comparative Pathology</i> , 2018, 159, 36-48.	0.3	13
84	Reverse Genetics of Newcastle Disease Virus. <i>Methods in Molecular Biology</i> , 2017, 1602, 141-158.	0.0	12
85	Avian Paramyxovirus Serotype 1 (Newcastle Disease Virus), Avian Influenza Virus, and <i>Salmonella</i> spp. in Mute Swans (<i>Cygnus olor</i>) in the Great Lakes Region and Atlantic Coast of the United States. <i>Avian Diseases</i> , 2014, 58, 129-136.	1.2	11
86	Complete Genome Sequence of a Recent Panzootic Virulent Newcastle Disease Virus from Pakistan. <i>Genome Announcements</i> , 2015, 3, .	0.8	11
87	Identification of Avian Paramyxovirus Serotype-1 in Wild Birds in the USA. <i>Journal of Wildlife Diseases</i> , 2016, 52, 657.	0.8	11
88	Natural Infections With Pigeon Paramyxovirus Serotype 1: Pathologic Changes in Eurasian Collared-Doves (<i>Streptopelia decaocto</i>) and Rock Pigeons (<i>Columba livia</i>) in the United States. <i>Veterinary Pathology</i> , 2017, 54, 695-703.	2.0	11
89	First Complete Genome Sequence of a Subgenotype Vd Newcastle Disease Virus Isolate. <i>Microbiology Resource Announcements</i> , 2019, 8, .	2.0	11
90	Derivation of chicken induced pluripotent stem cells tolerant to Newcastle disease virus-induced lysis through multiple rounds of infection. <i>Virology Journal</i> , 2016, 13, 205.	3.6	10

#	ARTICLE	IF	CITATIONS
91	Genetic stability of a Newcastle disease virus vectored infectious laryngotracheitis virus vaccine after serial passages in chicken embryos. <i>Vaccine</i> , 2020, 38, 925-932.	4.0	10
92	Single-Nucleotide Polymorphism Analysis to Select Conserved Regions for an Improved Real-Time Reverse Transcriptionâ€“PCR Test Specific for Newcastle Disease Virus. <i>Avian Diseases</i> , 2019, 63, 625.	1.2	9
93	Sequencing of avian influenza virus genomes following random amplification. <i>BioTechniques</i> , 2007, 43, 188-192.	1.8	8
94	Repeated Challenge with Virulent Newcastle Disease Virus Does Not Decrease the Efficacy of Vaccines. <i>Avian Diseases</i> , 2017, 61, 245-249.	1.2	8
95	Detection and Genome Sequence Analysis of Avian Metapneumovirus Subtype A Viruses Circulating in Commercial Chicken Flocks in Mexico. <i>Veterinary Sciences</i> , 2022, 9, 579.	1.7	8
96	Unique Variants of Avian Coronaviruses from Indigenous Chickens in Kenya. <i>Viruses</i> , 2023, 15, 264.	3.4	8
97	Effect of Infection with a Mesogenic Strain of Newcastle Disease Virus on Infection with Highly Pathogenic Avian Influenza Virus in Chickens. <i>Avian Diseases</i> , 2016, 60, 269-278.	1.2	7
98	Complete Genome Sequences of Four Avian Paramyxoviruses of Serotype 10 Isolated from Rockhopper Penguins on the Falkland Islands. <i>Genome Announcements</i> , 2017, 5, .	0.8	7
99	First Genome Sequence of Newcastle Disease Virus of Genotype VIIi from Jordan. <i>Microbiology Resource Announcements</i> , 2018, 7, .	2.0	7
100	Complete Genome Sequence of Genotype VI Newcastle Disease Viruses Isolated from Pigeons in Pakistan. <i>Genome Announcements</i> , 2016, 4, .	0.8	6
101	Risk factors for the transmission of infectious diseases agents at the wild birds -commercial birds interface. a pilot study in the region of the altos de Jalisco, Mexico. <i>Bulletin De L'Academie Veterinaire De France</i> , 2017, 170, 142-150.	0.1	6
102	Complete Genome Sequence of a Virulent Newcastle Disease Virus Strain Isolated from a Clinically Healthy Duck (<i>Anas platyrhynchos domesticus</i>) in Pakistan. <i>Genome Announcements</i> , 2016, 4, .	0.8	5
103	Identification and Complete Genome Sequence Analysis of a Genotype XIV Newcastle Disease Virus from Nigeria. <i>Genome Announcements</i> , 2016, 4, .	0.8	5
104	Draft Genome Sequences of Five Novel <i>Ochrobactrum</i> spp. Isolated from Different Avian Hosts in Nigeria. <i>Genome Announcements</i> , 2018, 6, .	0.8	5
105	Enhanced phylogenetic resolution of Newcastle disease outbreaks using complete viral genome sequences from formalin-fixed paraffin-embedded tissue samples. <i>Virus Genes</i> , 2019, 55, 502-512.	1.8	5
106	Genomic comparison of the complete coding and intergenic regions of the VG/GA Newcastle disease virus and its respirotropic clone 5. <i>Virus Genes</i> , 2008, 37, 161-167.	1.8	4
107	Complete Genome Sequences of New Emerging Newcastle Disease Virus Strains Isolated from China. <i>Genome Announcements</i> , 2013, 1, .	0.8	4
108	A Novel Recombinant Newcastle Disease Vaccine Improves Post- In Ovo Vaccination Survival with Sustained Protection against Virulent Challenge. <i>Vaccines</i> , 2021, 9, 953.	4.5	4

#	ARTICLE	IF	CITATIONS
109	Evaluation of chickens infected with a recombinant virulent NDV clone expressing chicken IL4. <i>Microbial Pathogenesis</i> , 2021, 159, 105116.	2.9	4
110	Experimental Infection and Transmission of Newcastle Disease Vaccine Virus in Four Wild Passerines. <i>Avian Diseases</i> , 2019, 63, 389.	1.2	4
111	Novel Recombinant Newcastle Disease Virus-Based In Ovo Vaccines Bypass Maternal Immunity to Provide Full Protection from Early Virulent Challenge. <i>Vaccines</i> , 2021, 9, 1189.	4.5	4
112	Newcastle Disease Virus. , 0, , .		3
113	Complete Genome Sequence of <i>Avian Coronavirus</i> Strain GA08 (GI-27 Lineage). <i>Microbiology Resource Announcements</i> , 2020, 9, .	2.0	3
114	Near-Complete Genome Sequences of Five Siciniviruses from North America. <i>Microbiology Resource Announcements</i> , 2021, 10, .	2.0	3
115	Draft Genome Sequences of Three <i>Ochrobactrum</i> spp. Isolated from Different Avian Hosts in Pakistan. <i>Genome Announcements</i> , 2018, 6, .	0.8	2
116	A 25-Year-Old Sample Contributes the Complete Genome Sequence of Avian Coronavirus Vaccine Strain ArkDPI, Reisolated from Commercial Broilers in the United States. <i>Microbiology Resource Announcements</i> , 2020, 9, .	2.0	2
117	Complete Genome Sequences of 11 Newcastle Disease Virus Isolates of Subgenotype VII.2 from Indonesia. <i>Microbiology Resource Announcements</i> , 2020, 9, .	2.0	2
118	A 24-Year-Old Sample Contributes the Complete Genome Sequence of Fowl Aviadenovirus D from the United States. <i>Microbiology Resource Announcements</i> , 2021, 10, .	2.0	2
119	Complete Genome Sequences of Avian Metapneumovirus Subtype B Vaccine Strains from Brazil. <i>Microbiology Resource Announcements</i> , 2023, 12, .	2.0	2
120	Next-Generation Sequencing for the Detection of Microbial Agents in Avian Clinical Samples. <i>Veterinary Sciences</i> , 2023, 10, 690.	1.7	2
121	Reply to "May Newly Defined Subgenotypes Va and Vb of Newcastle Disease Virus in Poultry Be Considered Two Different Genotypes?" <i>Journal of Clinical Microbiology</i> , 2016, 54, 2205-2206.	4.4	1
122	Intracellular fixation buffer inactivates Newcastle disease virus in chicken allantoic fluid, macrophages and splenocytes. <i>Journal of Virological Methods</i> , 2018, 251, 1-6.	2.1	1
123	Whole-Genome Sequence of <i>Avian coronavirus</i> from a 15-Year-Old Sample Confirms Evidence of GA08-like Strain Circulation 4 Years Prior to Its First Reported Outbreak. <i>Microbiology Resource Announcements</i> , 2021, 10, .	2.0	1
124	Complete Coding Sequences of Three Chicken Parvovirus Isolates from the United States. <i>Microbiology Resource Announcements</i> , 2020, 9, .	2.0	1
125	Complete genome sequence of seven virulent Newcastle disease virus isolates of sub-genotype XIII.1.1 from Tanzania. <i>Microbiology Resource Announcements</i> , 2023, 12, .	2.0	1
126	Complete Genome Sequence of an Avian Orthoavulavirus 13 Strain Detected in Ukraine. <i>Microbiology Resource Announcements</i> , 2023, 12, .	2.0	0

#	ARTICLE	IF	CITATIONS
127	Non-Targeted RNA Sequencing: Towards the Development of Universal Clinical Diagnosis Methods for Human and Veterinary Infectious Diseases. <i>Veterinary Sciences</i> , 2024, 11, 239.	1.7	0
128	Sequencing of historic samples provides complete coding sequences of chicken calicivirus from the United States. <i>Microbiology Resource Announcements</i> , 0, , .	2.0	0