

# Kiril M Dimitrov

## List of Publications by Year in descending order

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46  
papers

1,493  
citations

394421

19  
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330143

37  
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49  
all docs

49  
docs citations

49  
times ranked

1159  
citing authors

#	ARTICLE	IF	CITATIONS
1	Putative Novel Avian Paramyxovirus (AMPV) and Reidentification of APMV-2 and APMV-6 to the Species Level Based on Wild Bird Surveillance (United States, 2016–2018). <i>Applied and Environmental Microbiology</i> , 2022, 88, .	3.1	5
2	Genetic and biological characterization of Newcastle disease viruses circulating in Bangladesh during 2010–2017: further genetic diversification of class II genotype XIII in Southcentral Asia. <i>Journal of General Virology</i> , 2021, 102, .	2.9	11
3	A Novel Recombinant Newcastle Disease Vaccine Improves Post- In Ovo Vaccination Survival with Sustained Protection against Virulent Challenge. <i>Vaccines</i> , 2021, 9, 953.	4.4	4
4	A Pigeon-Derived Sub-Genotype XXI.1.2 Newcastle Disease Virus from Bangladesh Induces High Mortality in Chickens. <i>Viruses</i> , 2021, 13, 1520.	3.3	10
5	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.	4.4	3
6	Complete Genome Sequences of 11 Newcastle Disease Virus Isolates of Subgenotype VII.2 from Indonesia. <i>Microbiology Resource Announcements</i> , 2020, 9, .	0.6	2
7	Complete Genome Sequencing, Molecular Epidemiological, and Pathogenicity Analysis of Pigeon Paramyxoviruses Type 1 Isolated in Guangxi, China during 2012–2018. <i>Viruses</i> , 2020, 12, 366.	3.3	13
8	First Complete Genome Sequence of a Subgenotype Vd Newcastle Disease Virus Isolate. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.6	10
9	Rapid, multiplexed, whole genome and plasmid sequencing of foodborne pathogens using long-read nanopore technology. <i>Scientific Reports</i> , 2019, 9, 16350.	3.3	49
10	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.	3.2	38
11	Updated unified phylogenetic classification system and revised nomenclature for Newcastle disease virus. <i>Infection, Genetics and Evolution</i> , 2019, 74, 103917.	2.3	227
12	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	31
13	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.	2.1	28
14	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.	1.6	5
15	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.	2.4	28
16	Presence of Newcastle disease viruses of sub-genotypes Vc and VI in backyard chickens and in apparently healthy wild birds from Mexico in 2017. <i>Virus Genes</i> , 2019, 55, 479-489.	1.6	14
17	Experimental Infection and Transmission of Newcastle Disease Vaccine Virus in Four Wild Passerines. <i>Avian Diseases</i> , 2019, 63, 389.	1.0	4
18	Newcastle Disease Virus Infection in Quail. <i>Veterinary Pathology</i> , 2018, 55, 682-692.	1.7	16

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19	Evaluation of Protective Efficacy When Combining Turkey Herpesvirusâ€“Vector Vaccines. <i>Avian Diseases</i> , 2018, 63, 75.	1.0	21
20	Rapid virulence prediction and identification of Newcastle disease virus genotypes using third-generation sequencing. <i>Virology Journal</i> , 2018, 15, 179.	3.4	25
21	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.	3.4	31
22	An Epizootiological Report of the Re-emergence and Spread of a Lineage of Virulent Newcastle Disease Virus into Eastern Europe. <i>Transboundary and Emerging Diseases</i> , 2017, 64, 1001-1007.	3.0	31
23	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
24	A robust and cost-effective approach to sequence and analyze complete genomes of small RNA viruses. <i>Virology Journal</i> , 2017, 14, 72.	3.4	75
25	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	42
26	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
27	Newcastle disease vaccinesâ€“A solved problem or a continuous challenge?. <i>Veterinary Microbiology</i> , 2017, 206, 126-136.	1.9	239
28	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.	3.4	17
29	Repeated Challenge with Virulent Newcastle Disease Virus Does Not Decrease the Efficacy of Vaccines. <i>Avian Diseases</i> , 2017, 61, 245-249.	1.0	6
30	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.	1.9	44
31	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
32	Complete Genome Sequence of Genotype VI Newcastle Disease Viruses Isolated from Pigeons in Pakistan. <i>Genome Announcements</i> , 2016, 4, .	0.8	6
33	Complete Genome Sequence of an Avian Paramyxovirus Representative of Putative New Serotype 13. <i>Genome Announcements</i> , 2016, 4, .	0.8	21
34	Phylogenetic analysis of the complete genome of the APMV-13 isolate from Ukraine. <i>International Journal of Infectious Diseases</i> , 2016, 45, 459.	3.3	4
35	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	15
36	Identification and Complete Genome Sequence Analysis of a Genotype XIV Newcastle Disease Virus from Nigeria. <i>Genome Announcements</i> , 2016, 4, .	0.8	5

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37	Repeated isolation of virulent Newcastle disease viruses of sub-genotype VIIId from backyard chickens in Bulgaria and Ukraine between 2002 and 2013. Archives of Virology, 2016, 161, 3345-3353.	2.1	22
38	Reply to "May Newly Defined Subgenotypes Va and Vb of Newcastle Disease Virus in Poultry Be Considered Two Different Genotypes?" Journal of Clinical Microbiology, 2016, 54, 2205-2206.	3.9	1
39	Histopathologic Characterization and Shedding Dynamics of Guineafowl ( <i>Numida meleagris</i> ) Intravenously Infected with a H6N2 Low Pathogenicity Avian Influenza Virus. Avian Diseases, 2016, 60, 279-285.	1.0	1
40	Identification of Avian Paramyxovirus Serotype-1 in Wild Birds in the USA. Journal of Wildlife Diseases, 2016, 52, 657.	0.8	11
41	Temporal, geographic, and host distribution of avian paramyxovirus 1 (Newcastle disease virus). Infection, Genetics and Evolution, 2016, 39, 22-34.	2.3	216
42	Newcastle Disease Viruses Causing Recent Outbreaks Worldwide Show Unexpectedly High Genetic Similarity to Historical Virulent Isolates from the 1940s. Journal of Clinical Microbiology, 2016, 54, 1228-1235.	3.9	39
43	Presence of Vaccine-Derived Newcastle Disease Viruses in Wild Birds. PLoS ONE, 2016, 11, e0162484.	2.5	52
44	International Biological Engagement Programs Facilitate Newcastle Disease Epidemiological Studies. Frontiers in Public Health, 2015, 3, 235.	2.7	29
45	Molecular genetics and SSR markers as a new practice in farm animal genomic analysis for breeding and control of disease disorders. Biotechnology in Animal Husbandry, 2013, 29, 405-429.	0.3	3
46	Status of Wild Birds in Bulgarian Zoos with Regard to Orthomyxovirus and Paramyxovirus Type 1 Infections. Avian Diseases, 2010, 54, 361-364.	1.0	3