

# Poonam Sharma

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4345797/publications.pdf>

Version: 2024-02-01

29  
papers

537  
citations

759233

12  
h-index

677142

22  
g-index

29  
all docs

29  
docs citations

29  
times ranked

894  
citing authors

#	ARTICLE	IF	CITATIONS
1	Use of Sequence-Independent, Single-Primer-Amplification (SISPA) for rapid detection, identification, and characterization of avian RNA viruses. <i>Virology</i> , 2017, 509, 159-166.	2.4	117
2	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
3	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
4	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
5	Whole-Genome Sequence of <i>Chryseobacterium oranimense</i> , a Colistin-Resistant Bacterium Isolated from a Cystic Fibrosis Patient in France. <i>Antimicrobial Agents and Chemotherapy</i> , 2015, 59, 1696-1706.	3.2	29
6	Circulation of emerging NDM-5-producing <i>Escherichia coli</i> among humans and dogs in Egypt. <i>Zoonoses and Public Health</i> , 2020, 67, 324-329.	2.2	26
7	Genomic comparison of diverse <i>Salmonella</i> serovars isolated from swine. <i>PLoS ONE</i> , 2019, 14, e0224518.	2.5	25
8	Phenotypic and genotypic properties of <i>Microbacterium yannicii</i> , a recently described multidrug resistant bacterium isolated from a lung transplanted patient with cystic fibrosis in France. <i>BMC Microbiology</i> , 2013, 13, 97.	3.3	22
9	Complete Genome Sequence of an Avian Paramyxovirus Representative of Putative New Serotype 13. <i>Genome Announcements</i> , 2016, 4, .	0.8	21
10	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
11	Draft genome sequences of two ciprofloxacin-resistant <i>Salmonella enterica</i> subsp. <i>enterica</i> serotype Kentucky ST198 isolated from retail chicken carcasses in Egypt. <i>Journal of Global Antimicrobial Resistance</i> , 2018, 14, 101-103.	2.2	13
12	Comparison of Antimicrobial Resistance and Pan-Genome of Clinical and Non-Clinical <i>Enterococcus cecorum</i> from Poultry Using Whole-Genome Sequencing. <i>Foods</i> , 2020, 9, 686.	4.3	13
13	Genome Analysis of Multidrug-Resistant <i>Escherichia coli</i> Isolated from Poultry in Nigeria. <i>Foodborne Pathogens and Disease</i> , 2020, 17, 1-7.	1.8	12
14	Genome Sequence of <i>Microbacterium yannicii</i> , a Bacterium Isolated from a Cystic Fibrosis Patient. <i>Journal of Bacteriology</i> , 2012, 194, 4785-4785.	2.2	11
15	Whole genome sequencing of bacteria in cystic fibrosis as a model for bacterial genome adaptation and evolution. <i>Expert Review of Anti-Infective Therapy</i> , 2014, 12, 343-355.	4.4	11
16	Plasmid Replicons and $\beta$ -Lactamase-Encoding Genes of Multidrug-Resistant <i>Escherichia coli</i> Isolated from Humans and Food Animals in Lagos, Southwest Nigeria. <i>Microbial Drug Resistance</i> , 2019, 25, 1410-1423.	2.0	11
17	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
18	Serotyping of sub-Saharan Africa <i>Salmonella</i> strains isolated from poultry feces using multiplex PCR and whole genome sequencing. <i>BMC Microbiology</i> , 2021, 21, 29.	3.3	7

#	ARTICLE	IF	CITATIONS
19	Complete Genome Sequence of Genotype VI Newcastle Disease Viruses Isolated from Pigeons in Pakistan. <i>Genome Announcements</i> , 2016, 4, .	0.8	6
20	Identification and Complete Genome Sequence Analysis of a Genotype XIV Newcastle Disease Virus from Nigeria. <i>Genome Announcements</i> , 2016, 4, .	0.8	5
21	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
22	Draft Genome Sequence Analysis of Multidrug-Resistant <i>Escherichia coli</i> Strains Isolated in 2013 from Humans and Chickens in Nigeria. <i>Genome Announcements</i> , 2017, 5, .	0.8	5
23	Genomic Analysis of Multidrug-Resistant <i>Escherichia coli</i> from Surface Water in Northeast Georgia, United States: Presence of an ST131 Epidemic Strain Containing <i>bla</i> <sub>CTX-M-15</sub> on a Phage-Like Plasmid. <i>Microbial Drug Resistance</i> , 2020, 26, 447-455.	2.0	4
24	Genomic Comparison of Conjugative Plasmids from <i>Salmonella enterica</i> and <i>Escherichia coli</i> Encoding Beta-Lactamases and Capable of Mobilizing Kanamycin Resistance Col-like Plasmids. <i>Microorganisms</i> , 2021, 9, 2205.	3.6	4
25	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
26	Draft genome sequence of a human-associated streptogramin-resistant <i>Staphylococcus aureus</i> . <i>Journal of Global Antimicrobial Resistance</i> , 2019, 16, 72-73.	2.2	2
27	Training, Tests, and Tech. Health Information Systems and the Advancement of Medical Practice in Developing Countries, 2022, , 222-261.	0.1	2
28	Genome analysis of <i>Salmonella</i> strains isolated from imported frozen fish in Burkina Faso. <i>Annals of Microbiology</i> , 2021, 71, .	2.6	1
29	Draft Genome Sequences of Eight Streptogramin-Resistant <i>Enterococcus</i> Species Isolated from Animal and Environmental Sources in the United States. <i>Genome Announcements</i> , 2017, 5, .	0.8	0