

# 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

759055

12  
h-index

677027

22  
g-index

29  
all docs

29  
docs citations

29  
times ranked

894  
citing authors

#	ARTICLE	IF	CITATIONS
1	Training, Tests, and Tech. Health Information Systems and the Advancement of Medical Practice in Developing Countries, 2022, , 222-261.	0.1	2
2	Serotyping of sub-Saharan Africa Salmonella strains isolated from poultry feces using multiplex PCR and whole genome sequencing. BMC Microbiology, 2021, 21, 29.	1.3	7
3	Genome analysis of Salmonella strains isolated from imported frozen fish in Burkina Faso. Annals of Microbiology, 2021, 71, .	1.1	1
4	Genomic Comparison of Conjugative Plasmids from Salmonella enterica and Escherichia coli Encoding Beta-Lactamases and Capable of Mobilizing Kanamycin Resistance Col-like Plasmids. Microorganisms, 2021, 9, 2205.	1.6	4
5	Genome Analysis of Multidrug-Resistant <i>Escherichia coli</i> Isolated from Poultry in Nigeria. Foodborne Pathogens and Disease, 2020, 17, 1-7.	0.8	12
6	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. Microbial Drug Resistance, 2020, 26, 447-455.	0.9	4
7	Circulation of emerging NDM-5-producing <i>Escherichia coli</i> among humans and dogs in Egypt. Zoonoses and Public Health, 2020, 67, 324-329.	0.9	26
8	Comparison of Antimicrobial Resistance and Pan-Genome of Clinical and Non-Clinical Enterococcus cecorum from Poultry Using Whole-Genome Sequencing. Foods, 2020, 9, 686.	1.9	13
9	Plasmid Replicons and $\beta$ -Lactamase-Encoding Genes of Multidrug-Resistant <i>Escherichia coli</i> Isolated from Humans and Food Animals in Lagos, Southwest Nigeria. Microbial Drug Resistance, 2019, 25, 1410-1423.	0.9	11
10	Genomic comparison of diverse Salmonella serovars isolated from swine. PLoS ONE, 2019, 14, e0224518.	1.1	25
11	Draft genome sequence of a human-associated streptogramin-resistant Staphylococcus aureus. Journal of Global Antimicrobial Resistance, 2019, 16, 72-73.	0.9	2
12	Draft Genome Sequences of Five Novel Ochrobactrum spp. Isolated from Different Avian Hosts in Nigeria. Genome Announcements, 2018, 6, .	0.8	5
13	Draft genome sequences of two ciprofloxacin-resistant Salmonella enterica subsp. enterica serotype Kentucky ST198 isolated from retail chicken carcasses in Egypt. Journal of Global Antimicrobial Resistance, 2018, 14, 101-103.	0.9	13
14	Draft Genome Sequences of Three Ochrobactrum spp. Isolated from Different Avian Hosts in Pakistan. Genome Announcements, 2018, 6, .	0.8	2
15	A robust and cost-effective approach to sequence and analyze complete genomes of small RNA viruses. Virology Journal, 2017, 14, 72.	1.4	75
16	Repeated isolation of virulent Newcastle disease viruses in poultry and captive non-poultry avian species in Pakistan from 2011 to 2016. Preventive Veterinary Medicine, 2017, 142, 1-6.	0.7	42
17	Complete Genome Sequences of Four Avian Paramyxoviruses of Serotype 10 Isolated from Rockhopper Penguins on the Falkland Islands. Genome Announcements, 2017, 5, .	0.8	7
18	Use of Sequence-Independent, Single-Primer-Amplification (SISPA) for rapid detection, identification, and characterization of avian RNA viruses. Virology, 2017, 509, 159-166.	1.1	117

#	ARTICLE	IF	CITATIONS
19	Phylogenetic assessment reveals continuous evolution and circulation of pigeon-derived virulent avian avulaviruses 1 in Eastern Europe, Asia, and Africa. BMC Veterinary Research, 2017, 13, 291.	0.7	44
20	Draft Genome Sequences of Eight Streptogramin-Resistant Enterococcus Species Isolated from Animal and Environmental Sources in the United States. Genome Announcements, 2017, 5, .	0.8	0
21	Draft Genome Sequence Analysis of Multidrug-Resistant Escherichia coli Strains Isolated in 2013 from Humans and Chickens in Nigeria. Genome Announcements, 2017, 5, .	0.8	5
22	Complete Genome Sequence of Genotype VI Newcastle Disease Viruses Isolated from Pigeons in Pakistan. Genome Announcements, 2016, 4, .	0.8	6
23	Complete Genome Sequence of an Avian Paramyxovirus Representative of Putative New Serotype 13. Genome Announcements, 2016, 4, .	0.8	21
24	Complete Genome Sequence of a Genotype XVII Newcastle Disease Virus, Isolated from an Apparently Healthy Domestic Duck in Nigeria. Genome Announcements, 2016, 4, .	0.8	15
25	Identification and Complete Genome Sequence Analysis of a Genotype XIV Newcastle Disease Virus from Nigeria. Genome Announcements, 2016, 4, .	0.8	5
26	Whole-Genome Sequence of Chryseobacterium oranimense, a Colistin-Resistant Bacterium Isolated from a Cystic Fibrosis Patient in France. Antimicrobial Agents and Chemotherapy, 2015, 59, 1696-1706.	1.4	29
27	Whole genome sequencing of bacteria in cystic fibrosis as a model for bacterial genome adaptation and evolution. Expert Review of Anti-Infective Therapy, 2014, 12, 343-355.	2.0	11
28	Phenotypic and genotypic properties of Microbacterium yannicii, a recently described multidrug resistant bacterium isolated from a lung transplanted patient with cystic fibrosis in France. BMC Microbiology, 2013, 13, 97.	1.3	22
29	Genome Sequence of Microbacterium yannicii, a Bacterium Isolated from a Cystic Fibrosis Patient. Journal of Bacteriology, 2012, 194, 4785-4785.	1.0	11