## Poonam Sharma

List of Publications by Year in descending order

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759233 677142 29 537 12 22 citations h-index g-index papers 29 29 29 894 citing authors docs citations times ranked all docs

#	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.	3.3	7
3	Genome analysis of Salmonella strains isolated from imported frozen fish in Burkina Faso. Annals of Microbiology, 2021, 71, .	2.6	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.	3.6	4
5	Genome Analysis of Multidrug-Resistant <i>Escherichia coli</i> Isolated from Poultry in Nigeria. Foodborne Pathogens and Disease, 2020, 17, 1-7.	1.8	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.	2.0	4
7	Circulation of emerging NDMâ€5â€producing <i>Escherichia coli</i> among humans and dogs in Egypt. Zoonoses and Public Health, 2020, 67, 324-329.	2.2	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.	4.3	13
9	Plasmid Replicons and β-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.	2.0	11
10	Genomic comparison of diverse Salmonella serovars isolated from swine. PLoS ONE, 2019, 14, e0224518.	2.5	25
11	Draft genome sequence of a human-associated streptogramin-resistant Staphylococcus aureus. Journal of Global Antimicrobial Resistance, 2019, 16, 72-73.	2.2	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.	2.2	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.	3.4	<b>7</b> 5
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.	1.9	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.	2.4	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.	1.9	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.	3.2	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.	4.4	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.	3.3	22
29	Genome Sequence of Microbacterium yannicii, a Bacterium Isolated from a Cystic Fibrosis Patient. Journal of Bacteriology, 2012, 194, 4785-4785.	2.2	11