

# Charlene R Jackson

## List of Publications by Year in descending order

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107  
papers

2,817  
citations

186265

28  
h-index

206112

48  
g-index

112  
all docs

112  
docs citations

112  
times ranked

3138  
citing authors

#	ARTICLE	IF	CITATIONS
1	Use of a Genus- and Species-Specific Multiplex PCR for Identification of Enterococci. <i>Journal of Clinical Microbiology</i> , 2004, 42, 3558-3565.	3.9	378
2	Genetic mechanisms of antimicrobial resistance identified in <i>Salmonella enterica</i> , <i>Escherichia coli</i> , and <i>Enterococcus</i> spp. isolated from U.S. food animals. <i>Frontiers in Microbiology</i> , 2013, 4, 135.	3.5	147
3	Prevalence and Characterization of Methicillin-Resistant <i>Staphylococcus aureus</i> Isolates from Retail Meat and Humans in Georgia. <i>Journal of Clinical Microbiology</i> , 2013, 51, 1199-1207.	3.9	137
4	Antimicrobial Resistance Genes, Cassettes, and Plasmids Present in <i>Salmonella enterica</i> Associated With United States Food Animals. <i>Frontiers in Microbiology</i> , 2019, 10, 832.	3.5	95
5	DNA microarray detection of antimicrobial resistance genes in diverse bacteria. <i>International Journal of Antimicrobial Agents</i> , 2006, 27, 138-151.	2.5	94
6	Effects of Tylosin Use on Erythromycin Resistance in Enterococci Isolated from Swine. <i>Applied and Environmental Microbiology</i> , 2004, 70, 4205-4210.	3.1	93
7	Prevalence, species distribution and antimicrobial resistance of enterococci isolated from dogs and cats in the United States. <i>Journal of Applied Microbiology</i> , 2009, 107, 1269-1278.	3.1	82
8	Epidemiology and Genotypic Characteristics of Methicillin-Resistant <i>Staphylococcus aureus</i> Strains of Porcine Origin. <i>Journal of Clinical Microbiology</i> , 2012, 50, 3687-3693.	3.9	74
9	Carriage of methicillin-resistant staphylococci by healthy companion animals in the US. <i>Letters in Applied Microbiology</i> , 2014, 59, 1-8.	2.2	65
10	Antimicrobial Resistance in <i>Enterococcus</i> spp. Isolated from Environmental Samples in an Area of Intensive Poultry Production. <i>International Journal of Environmental Research and Public Health</i> , 2013, 10, 1020-1036.	2.6	64
11	Prevalence and Antimicrobial Resistance of Enterococci Isolated from Retail Fruits, Vegetables, and Meats. <i>Journal of Food Protection</i> , 2006, 69, 2976-2982.	1.7	61
12	Comparative Antimicrobial Susceptibility of <i>Listeria monocytogenes</i> , <i>L. innocua</i> , and <i>L. welshimeri</i> . <i>Microbial Drug Resistance</i> , 2009, 15, 27-32.	2.0	55
13	Development of a DNA Microarray to Detect Antimicrobial Resistance Genes Identified in the National Center for Biotechnology Information Database. <i>Microbial Drug Resistance</i> , 2010, 16, 9-19.	2.0	55
14	Detection and Control of a Nosocomial Outbreak Caused by <i>Salmonella</i> Newport at a Large Animal Hospital. <i>Journal of Veterinary Internal Medicine</i> , 2010, 24, 606-616.	1.6	50
15	Transferable Plasmids of <i>Salmonella enterica</i> Associated With Antibiotic Resistance Genes. <i>Frontiers in Microbiology</i> , 2020, 11, 562181.	3.5	49
16	Prevalence, species distribution and antimicrobial resistance of enterococci isolated from US dairy cattle. <i>Letters in Applied Microbiology</i> , 2011, 52, 41-48.	2.2	48
17	Multidrug resistant <i>Mannheimia haemolytica</i> isolated from high-risk beef stocker cattle after antimicrobial metaphylaxis and treatment for bovine respiratory disease. <i>Veterinary Microbiology</i> , 2018, 221, 143-152.	1.9	45
18	Antibiotic Resistance Patterns of Major Zoonotic Pathogens from All-Natural, Antibiotic-Free, Pasture-Raised Broiler Flocks in the Southeastern United States. <i>Journal of Environmental Quality</i> , 2016, 45, 593-603.	2.0	44

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19	Antimicrobial Resistance and Virulence of <i>Enterococcus faecalis</i> Isolated from Retail Food. <i>Journal of Food Protection</i> , 2008, 71, 760-769.	1.7	43
20	<i>Salmonella</i> Enteritidis in Meat, Poultry, and Pasteurized Egg Products Regulated by the U.S. Food Safety and Inspection Service, 1998 through 2003. <i>Journal of Food Protection</i> , 2007, 70, 582-591.	1.7	41
21	Prevalence and Antimicrobial Resistance in <i>Escherichia coli</i> from Food Animals in Lagos, Nigeria. <i>Microbial Drug Resistance</i> , 2015, 21, 358-365.	2.0	41
22	Antimicrobial Resistance, Genetic Diversity and Multilocus Sequence Typing of <i>Escherichia coli</i> from Humans, Retail Chicken and Ground Beef in Egypt. <i>Pathogens</i> , 2020, 9, 357.	2.8	35
23	The prevalence and antimicrobial resistance phenotypes of <i>Salmonella</i> , <i>Escherichia coli</i> and <i>Enterococcus</i> sp. in surface water. <i>Letters in Applied Microbiology</i> , 2020, 71, 3-25.	2.2	35
24	Prevalence and characterization of <i>Escherichia coli</i> isolated from the Upper Oconee Watershed in Northeast Georgia. <i>PLoS ONE</i> , 2018, 13, e0197005.	2.5	34
25	Prevalence of streptogramin resistance in enterococci from animals: identification of <i>vatD</i> from animal sources in the USA. <i>International Journal of Antimicrobial Agents</i> , 2007, 30, 60-66.	2.5	32
26	Related Antimicrobial Resistance Genes Detected in Different Bacterial Species Co-isolated from Swine Fecal Samples. <i>Foodborne Pathogens and Disease</i> , 2011, 8, 663-679.	1.8	32
27	Human-Associated Methicillin-Resistant <i>Staphylococcus aureus</i> from a Subtropical Recreational Marine Beach. <i>Microbial Ecology</i> , 2013, 65, 1039-1051.	2.8	32
28	An assay for determining the susceptibility of <i>Salmonella</i> isolates to commercial and household biocides. <i>PLoS ONE</i> , 2018, 13, e0209072.	2.5	31
29	Comparison of <i>dhgB</i> -linked intergenic sequence ribotyping to DNA microarray hybridization for assigning serotype to <i>Salmonella enterica</i> . <i>FEMS Microbiology Letters</i> , 2012, 337, 61-72.	1.8	30
30	Antimicrobial resistance, virulence determinants and genetic profiles of clinical and nonclinical <i>Enterococcus cecorum</i> from poultry. <i>Letters in Applied Microbiology</i> , 2015, 60, 111-119.	2.2	30
31	Prevalence and multidrug resistance of <i>Escherichia coli</i> from community-acquired infections in Lagos, Nigeria. <i>Journal of Infection in Developing Countries</i> , 2016, 10, 920-931.	1.2	27
32	<i>Salmonella</i> , <i>Campylobacter</i> and <i>Enterococcus</i> spp.: Their Antimicrobial Resistance Profiles and their Spatial Relationships in a Synoptic Study of the Upper Oconee River Basin. <i>Microbial Ecology</i> , 2008, 55, 444-452.	2.8	26
33	Clonally Related Methicillin-Resistant <i>Staphylococcus aureus</i> Isolated from Short-Finned Pilot Whales ( <i>Globicephala macrorhynchus</i> ), Human Volunteers, and a Bayfront Cetacean Rehabilitation Facility. <i>Microbial Ecology</i> , 2013, 65, 1024-1038.	2.8	26
34	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
35	Genetic Relatedness of High-Level Aminoglycoside-Resistant Enterococci Isolated from Poultry Carcasses. <i>Avian Diseases</i> , 2004, 48, 100-107.	1.0	25
36	High-level aminoglycoside resistant enterococci isolated from swine. <i>Epidemiology and Infection</i> , 2005, 133, 367-371.	2.1	25

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37	Mechanisms of antimicrobial resistance and genetic relatedness among enterococci isolated from dogs and cats in the United States. <i>Journal of Applied Microbiology</i> , 2009, 108, 2171-9.	3.1	25
38	Genomic comparison of diverse <i>Salmonella</i> serovars isolated from swine. <i>PLoS ONE</i> , 2019, 14, e0224518.	2.5	25
39	Carriage and Gene Content Variability of the pESI-Like Plasmid Associated with <i>Salmonella</i> Infantis Recently Established in United States Poultry Production. <i>Genes</i> , 2020, 11, 1516.	2.4	25
40	Analysis of <i>Salmonella enterica</i> with Reduced Susceptibility to the Third-Generation Cephalosporin Ceftriaxone Isolated from U.S. Cattle During 2000–2004. <i>Microbial Drug Resistance</i> , 2008, 14, 251-258.	2.0	23
41	Coproduction of Tet(X7) Conferring High-Level Tigecycline Resistance, Fosfomycin FosA4, and Colistin Mcr-1.1 in <i>Escherichia coli</i> Strains from Chickens in Egypt. <i>Antimicrobial Agents and Chemotherapy</i> , 2021, 65, .	3.2	22
42	Introduction to United States Department of Agriculture VetNet: Status of <i>Salmonella</i> and <i>Campylobacter</i> Databases from 2004 Through 2005. <i>Foodborne Pathogens and Disease</i> , 2007, 4, 241-248.	1.8	21
43	Genetic relatedness of a rarely isolated <i>Salmonella</i> : <i>Salmonella enterica</i> serotype Niakhar from NARMS animal isolates. <i>Journal of Antimicrobial Chemotherapy</i> , 2006, 57, 190-198.	3.0	20
44	Genetic Characterization of Antimicrobial-Resistant <i>Escherichia coli</i> Isolated from a Mixed-Use Watershed in Northeast Georgia, USA. <i>International Journal of Environmental Research and Public Health</i> , 2019, 16, 3761.	2.6	19
45	Emergence of Multidrug-Resistant <i>Escherichia coli</i> Producing CTX-M, MCR-1, and FosA in Retail Food From Egypt. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021, 11, 681588.	3.9	19
46	Effect of Growth Promotant Usage on Enterococci Species on a Poultry Farm. <i>Avian Diseases</i> , 2005, 49, 361-365.	1.0	18
47	Antibiotic-resistant bacteria and gut microbiome communities associated with wild-caught shrimp from the United States versus imported farm-raised retail shrimp. <i>Scientific Reports</i> , 2021, 11, 3356.	3.3	18
48	Detection and Molecular Characterization of Methicillin-Resistant <i>Staphylococcus aureus</i> from Table Eggs in Haripur, Pakistan. <i>Foodborne Pathogens and Disease</i> , 2018, 15, 86-93.	1.8	17
49	Non-point source fecal contamination from aging wastewater infrastructure is a primary driver of antibiotic resistance in surface waters. <i>Water Research</i> , 2022, 222, 118853.	11.3	17
50	<i>Staphylococci</i> in poultry intestines: a comparison between farmed and household chickens. <i>Poultry Science</i> , 2020, 99, 4549-4557.	3.4	16
51	Diversity and antimicrobial resistance of <i>Enterococcus</i> from the Upper Oconee Watershed, Georgia. <i>Journal of Applied Microbiology</i> , 2020, 128, 1221-1233.	3.1	15
52	Antimicrobial Resistance Gene Detection and Plasmid Typing Among Multidrug Resistant Enterococci Isolated from Freshwater Environment. <i>Microorganisms</i> , 2020, 8, 1338.	3.6	15
53	Effect of media, temperature and culture conditions on the species population and antibiotic resistance of enterococci from broiler chickens*. <i>Letters in Applied Microbiology</i> , 2005, 41, 262-268.	2.2	13
54	Evidence of a conjugal erythromycin resistance element in the Lyme disease spirochete <i>Borrelia burgdorferi</i> . <i>International Journal of Antimicrobial Agents</i> , 2007, 30, 496-504.	2.5	13

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55	Diversity of Plasmids and Antimicrobial Resistance Genes in Multidrug-Resistant <i>Escherichia coli</i> Isolated from Healthy Companion Animals. <i>Zoonoses and Public Health</i> , 2015, 62, 479-488.	2.2	13
56	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
57	Comparison of Antimicrobial Resistance and Pan-Genome of Clinical and Non-Clinical <i>Enterococcus faecium</i> from Poultry Using Whole-Genome Sequencing. <i>Foods</i> , 2020, 9, 686.	4.3	13
58	Characterization of <i>Salmonella enterica</i> Serovar Agona Slaughter Isolates from the Animal Arm of the National Antimicrobial Resistance Monitoring System <sup>®</sup> Enteric Bacteria (NARMS): 1997 through 2003. <i>Microbial Drug Resistance</i> , 2008, 14, 55-63.	2.0	12
59	Contribution of Healthy Chickens to Antimicrobial-Resistant <i>Escherichia coli</i> Associated with Human Extraintestinal Infections in Egypt. <i>Vector-Borne and Zoonotic Diseases</i> , 2018, 18, 408-416.	1.5	12
60	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
61	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
62	First report of <i>vatB</i> and <i>vgaB</i> from <i>Enterococcus gallinarum</i> in the USA. <i>International Journal of Antimicrobial Agents</i> , 2008, 31, 175-176.	2.5	9
63	Anatomical distribution and genetic relatedness of antimicrobial-resistant <i>Escherichia coli</i> from healthy companion animals. <i>Journal of Applied Microbiology</i> , 2011, 110, 597-604.	3.1	9
64	Emergence of <i>Salmonella</i> Genomic Island 1 Variant SGI1-C in a Multidrug-Resistant Clinical Isolate of <i>Klebsiella pneumoniae</i> ST485 from Egypt. <i>Antimicrobial Agents and Chemotherapy</i> , 2020, 64, .	3.2	8
65	Draft Genome Sequence of <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Bardo Strain CRJJGF_00099 (Phylum Gammaproteobacteria). <i>Genome Announcements</i> , 2016, 4, .	0.8	7
66	Incidence, species and antimicrobial resistance of naturally occurring <i>Campylobacter</i> isolates from quail carcasses sampled in a commercial processing facility. <i>Journal of Food Safety</i> , 2018, 38, e12438.	2.3	7
67	Detection and Molecular Characterization of Staphylococci from Eggs of Household Chickens. <i>Foodborne Pathogens and Disease</i> , 2019, 16, 550-557.	1.8	7
68	Draft genome sequence of a <i>bla</i> NDM-1- and <i>bla</i> OXA-244-carrying multidrug-resistant <i>Escherichia coli</i> D-ST69 clinical isolate from Egypt. <i>Journal of Global Antimicrobial Resistance</i> , 2020, 22, 832-834.	2.2	7
69	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
70	Effect of Subtherapeutic Antimicrobials on Genetic Diversity of <i>Enterococcus faecium</i> from Chickens. <i>Avian Diseases</i> , 2006, 50, 115-119.	1.0	6
71	A Comparison of BOX-PCR and Pulsed-Field Gel Electrophoresis to Determine Genetic Relatedness of <i>Enterococci</i> from Different Environments. <i>Microbial Ecology</i> , 2012, 64, 378-387.	2.8	6
72	Draft Genome Sequence of <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Orion Strain CRJJGF_00093 (Phylum Gammaproteobacteria). <i>Genome Announcements</i> , 2016, 4, .	0.8	6

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73	A newly developed <i>Escherichia coli</i> isolate panel from a cross section of U.S. animal production systems reveals geographic and commodity-based differences in antibiotic resistance gene carriage. <i>Journal of Hazardous Materials</i> , 2020, 382, 120991.	12.4	6
74	Isolation and characterization of two novel groups of kanamycin-resistance ColE1-like plasmids in <i>Salmonella enterica</i> serotypes from food animals. <i>PLoS ONE</i> , 2018, 13, e0193435.	2.5	6
75	Analysis of <i>Salmonella enterica</i> Isolated from a Mixed-Use Watershed in Georgia, USA: Antimicrobial Resistance, Serotype Diversity, and Genetic Relatedness to Human Isolates. <i>Applied and Environmental Microbiology</i> , 2022, 88, e0039322.	3.1	6
76	Detection of <i>Salmonella</i> Serotypes by Overnight Incubation of Entire Broiler Carcass. <i>Journal of Food Safety</i> , 2017, 37, e12298.	2.3	5
77	Carcass orientation and drip time affect potential surface water carryover for broiler carcasses subjected to a post-chill water dip or spray. <i>Poultry Science</i> , 2017, 96, 241-245.	3.4	5
78	Diversity of Plasmids and Genes Encoding Resistance to Extended-Spectrum $\beta$ -Lactamase in <i>Escherichia coli</i> from Different Animal Sources. <i>Microorganisms</i> , 2021, 9, 1057.	3.6	5
79	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
80	Resistance Genes, Plasmids, Multilocus Sequence Typing (MLST), and Phenotypic Resistance of Non-Typhoidal <i>Salmonella</i> (NTS) Isolated from Slaughtered Chickens in Burkina Faso. <i>Antibiotics</i> , 2022, 11, 782.	3.7	5
81	Draft Genome Sequence of <i>Salmonella enterica</i> subsp. <i>diarizonae</i> Serovar 61:k:1,5,(7) Strain CRJJGF_00165 (Phylum <i>Gammaproteobacteria</i> ). <i>Genome Announcements</i> , 2016, 4, .	0.8	4
82	Draft Genome Sequence of <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Putten Strain CRJJGF_00159 (Phylum <i>Gammaproteobacteria</i> ). <i>Genome Announcements</i> , 2016, 4, .	0.8	4
83	Draft Genome Sequence of <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Blockley Strain CRJJGF_00147 (Phylum <i>Gammaproteobacteria</i> ). <i>Genome Announcements</i> , 2016, 4, .	0.8	4
84	Draft Genome Sequence of <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Kiambu Strain CRJJGF_00061 (Phylum <i>Gammaproteobacteria</i> ). <i>Genome Announcements</i> , 2016, 4, .	0.8	4
85	Draft Genome Sequence of <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Lille Strain CRJJGF_000101 (Phylum <i>Gammaproteobacteria</i> ). <i>Genome Announcements</i> , 2016, 4, .	0.8	4
86	Draft Genome Sequence of <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Widemarsh Strain CRJJGF_00058 (Phylum <i>Gammaproteobacteria</i> ). <i>Genome Announcements</i> , 2016, 4, .	0.8	4
87	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
88	Antimicrobial interventions to reduce <i>Salmonella</i> and <i>Campylobacter</i> populations and improve shelf life of quail carcasses. <i>Poultry Science</i> , 2020, 99, 5977-5982.	3.4	4
89	Whole-Genome Sequence Analysis of Multidrug-Resistant <i>Enterobacter hormaechei</i> Isolated from Imported Retail Shrimp. <i>Microbiology Resource Announcements</i> , 2020, 9, .	0.6	4
90	Screening and Rapid Identification of <i>Campylobacter</i> Spp. DNA by FlaA PCR Based Method on Chicken and Human Fecal Samples in Egypt. <i>International Journal of Poultry Science</i> , 2015, 14, 252-256.	0.1	4

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91	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
92	Distribution and Transfer of Plasmid Replicon Families among Multidrug-Resistant <i>Enterococcus faecalis</i> and <i>Enterococcus faecium</i> from Poultry. <i>Microorganisms</i> , 2022, 10, 1244.	3.6	4
93	Detection of Plasmids and Class 1 Integrons in <i>Salmonella enterica</i> Serovar Agona Isolated from NARMS Slaughter Samples Collected in the Years 1997–2003. <i>Microbial Drug Resistance</i> , 2007, 13, 212-219.	2.0	3
94	Evaluation of a new chromogenic agar for the detection of environmental <i>Enterococcus</i> . <i>Journal of Microbiological Methods</i> , 2020, 178, 106082.	1.6	3
95	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
96	Genetic Diversity of <i>Staphylococcus aureus</i> Strains from a Tertiary Care Hospital in Rawalpindi, Pakistan. <i>Microorganisms</i> , 2021, 9, 2301.	3.6	2
97	Routes of transmission of <i>Salmonella</i> and <i>Campylobacter</i> in breeder turkeys. <i>Journal of Applied Poultry Research</i> , 2016, 25, 591-609.	1.2	1
98	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
99	Application of Metagenomic Technologies for Antimicrobial Resistance and Food Safety Research and Beyond—CYC and XY contributed equally to this review. The opinions expressed in this review are entirely those of the authors and do not represent those of the USDA.. , 2015, , 401-422.		0
100	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
101	Comparison of two commercially available rapid detection methods and a conventional culture method to detect naturally occurring salmonellae on broiler carcasses. <i>Journal of Food Safety</i> , 2019, 39, e12702.	2.3	0
102	Antimicrobial-resistant pathogens in water. <i>Letters in Applied Microbiology</i> , 2020, 71, 2-2.	2.2	0
103	Draft Genomic Sequences of Three <i>Escherichia coli</i> Sequence Type 131 Isolates (H45, H43ii, and H43iii) from Patients in Lagos, Nigeria. <i>Microbiology Resource Announcements</i> , 2020, 9, .	0.6	0
104	Genomic comparison of diverse <i>Salmonella</i> serovars isolated from swine. , 2019, 14, e0224518.		0
105	Genomic comparison of diverse <i>Salmonella</i> serovars isolated from swine. , 2019, 14, e0224518.		0
106	Genomic comparison of diverse <i>Salmonella</i> serovars isolated from swine. , 2019, 14, e0224518.		0
107	Genomic comparison of diverse <i>Salmonella</i> serovars isolated from swine. , 2019, 14, e0224518.		0