

# Jonathan Frye

## List of Publications by Citations

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128  
papers

4,981  
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35  
h-index

68  
g-index

141  
ext. papers

6,387  
ext. citations

4.9  
avg, IF

5.25  
L-index

#	Paper	IF	Citations
128	CARD 2017: expansion and model-centric curation of the comprehensive antibiotic resistance database. <i>Nucleic Acids Research</i> , <b>2017</b> , 45, D566-D573	20.1	1209
127	Regulation of Salmonella typhimurium virulence gene expression by cationic antimicrobial peptides. <i>Molecular Microbiology</i> , <b>2003</b> , 50, 219-30	4.1	208
126	Gene expression patterns during swarming in Salmonella typhimurium: genes specific to surface growth and putative new motility and pathogenicity genes. <i>Molecular Microbiology</i> , <b>2004</b> , 52, 169-87	4.1	177
125	Global regulation by CsrA in Salmonella typhimurium. <i>Molecular Microbiology</i> , <b>2003</b> , 48, 1633-45	4.1	174
124	Co-regulation of Salmonella enterica genes required for virulence and resistance to antimicrobial peptides by SlyA and PhoP/PhoQ. <i>Molecular Microbiology</i> , <b>2005</b> , 56, 492-508	4.1	170
123	Identification of new flagellar genes of Salmonella enterica serovar Typhimurium. <i>Journal of Bacteriology</i> , <b>2006</b> , 188, 2233-43	3.5	121
122	Multiplex PCR-based method for identification of common clinical serotypes of Salmonella enterica subsp. enterica. <i>Journal of Clinical Microbiology</i> , <b>2006</b> , 44, 3608-15	9.7	120
121	Genetic mechanisms of antimicrobial resistance identified in Salmonella enterica, Escherichia coli, and Enterococcus spp. isolated from U.S. food animals. <i>Frontiers in Microbiology</i> , <b>2013</b> , 4, 135	5.7	112
120	Enterobacter sakazakii invades brain capillary endothelial cells, persists in human macrophages influencing cytokine secretion and induces severe brain pathology in the neonatal rat. <i>Microbiology (United Kingdom)</i> , <b>2007</b> , 153, 3538-3547	2.9	103
119	The intestinal fatty acid propionate inhibits Salmonella invasion through the post-translational control of HilD. <i>Molecular Microbiology</i> , <b>2013</b> , 87, 1045-60	4.1	102
118	Salmonella enterica serovar Typhimurium requires the Lpf, Pef, and Tafi fimbriae for biofilm formation on HEp-2 tissue culture cells and chicken intestinal epithelium. <i>Infection and Immunity</i> , <b>2006</b> , 74, 3156-69	3.7	102
117	Insights into the complex regulation of rpoS in Borrelia burgdorferi. <i>Molecular Microbiology</i> , <b>2007</b> , 65, 277-93	4.1	101
116	Inc A/C plasmids are prevalent in multidrug-resistant Salmonella enterica isolates. <i>Applied and Environmental Microbiology</i> , <b>2009</b> , 75, 1908-15	4.8	83
115	DNA microarray detection of antimicrobial resistance genes in diverse bacteria. <i>International Journal of Antimicrobial Agents</i> , <b>2006</b> , 27, 138-51	14.3	80
114	Alternative sigma factor interactions in Salmonella: sigma and sigma promote antioxidant defences by enhancing sigma levels. <i>Molecular Microbiology</i> , <b>2005</b> , 56, 811-23	4.1	76
113	Prevalence, distribution and characterisation of ceftiofur resistance in Salmonella enterica isolated from animals in the USA from 1999 to 2003. <i>International Journal of Antimicrobial Agents</i> , <b>2007</b> , 30, 134-42	14.3	75
112	DNA microarray-based typing of an atypical monophasic Salmonella enterica serovar. <i>Journal of Clinical Microbiology</i> , <b>2002</b> , 40, 2074-8	9.7	74

111	A non-redundant microarray of genes for two related bacteria. <i>Nucleic Acids Research</i> , <b>2003</b> , 31, 1869-76	20.1	69
110	Genomic epidemiology of Salmonella enterica serotype Enteritidis based on population structure of prevalent lineages. <i>Emerging Infectious Diseases</i> , <b>2014</b> , 20, 1481-9	10.2	62
109	Identification and transcriptional analysis of a Treponema pallidum operon encoding a putative ABC transport system, an iron-activated repressor protein homolog, and a glycolytic pathway enzyme homolog. <i>Gene</i> , <b>1997</b> , 197, 47-64	3.8	62
108	Host gene expression changes and DNA amplification during temperate phage induction. <i>Journal of Bacteriology</i> , <b>2005</b> , 187, 1485-92	3.5	62
107	High-throughput molecular determination of salmonella enterica serovars by use of multiplex PCR and capillary electrophoresis analysis. <i>Journal of Clinical Microbiology</i> , <b>2009</b> , 47, 1290-9	9.7	61
106	Prevalence, species distribution and antimicrobial resistance of enterococci isolated from dogs and cats in the United States. <i>Journal of Applied Microbiology</i> , <b>2009</b> , 107, 1269-78	4.7	61
105	Detection of KPC-2 in a clinical isolate of Proteus mirabilis and first reported description of carbapenemase resistance caused by a KPC beta-lactamase in P. mirabilis. <i>Journal of Clinical Microbiology</i> , <b>2008</b> , 46, 3080-3	9.7	53
104	Antimicrobial Resistance Genes, Cassettes, and Plasmids Present in Associated With United States Food Animals. <i>Frontiers in Microbiology</i> , <b>2019</b> , 10, 832	5.7	51
103	Analysis of AI-2/LuxS-dependent transcription in Campylobacter jejuni strain 81-176. <i>Foodborne Pathogens and Disease</i> , <b>2008</b> , 5, 399-415	3.8	50
102	Fitness costs and stability of a high-level ciprofloxacin resistance phenotype in Salmonella enterica serotype enteritidis: reduced infectivity associated with decreased expression of Salmonella pathogenicity island 1 genes. <i>Antimicrobial Agents and Chemotherapy</i> , <b>2010</b> , 54, 367-74	5.9	49
101	Analysis of antimicrobial resistance genes detected in multidrug-resistant Salmonella enterica serovar Typhimurium isolated from food animals. <i>Microbial Drug Resistance</i> , <b>2011</b> , 17, 407-18	2.9	49
100	Correlation of phenotype with the genotype of egg-contaminating Salmonella enterica serovar Enteritidis. <i>Applied and Environmental Microbiology</i> , <b>2005</b> , 71, 4388-99	4.8	45
99	Development of a DNA microarray to detect antimicrobial resistance genes identified in the National Center for Biotechnology Information database. <i>Microbial Drug Resistance</i> , <b>2010</b> , 16, 9-19	2.9	44
98	Antimicrobial resistance genes in multidrug-resistant Salmonella enterica isolated from animals, retail meats, and humans in the United States and Canada. <i>Microbial Drug Resistance</i> , <b>2013</b> , 19, 175-84	2.9	41
97	AMRFinderPlus and the Reference Gene Catalog facilitate examination of the genomic links among antimicrobial resistance, stress response, and virulence. <i>Scientific Reports</i> , <b>2021</b> , 11, 12728	4.9	39
96	Antimicrobial resistance and virulence of Enterococcus faecalis isolated from retail food. <i>Journal of Food Protection</i> , <b>2008</b> , 71, 760-9	2.5	37
95	An FDA bioinformatics tool for microbial genomics research on molecular characterization of bacterial foodborne pathogens using microarrays. <i>BMC Bioinformatics</i> , <b>2010</b> , 11 Suppl 6, S4	3.6	35
94	Increased expression of Borrelia burgdorferi vlsE in response to human endothelial cell membranes. <i>Molecular Microbiology</i> , <b>2001</b> , 41, 229-39	4.1	35

93	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> , <b>2018</b> , 221, 143-152	3.3	32
92	Rainfall and tillage effects on transport of fecal bacteria and sex hormones 17beta-estradiol and testosterone from broiler litter applications to a Georgia Piedmont Ultisol. <i>Science of the Total Environment</i> , <b>2008</b> , 403, 154-63	10.2	32
91	Microarray analysis of antimicrobial resistance genes in <i>Salmonella enterica</i> from preharvest poultry environment. <i>Journal of Applied Microbiology</i> , <b>2009</b> , 107, 906-14	4.7	29
90	Prevalence and Antimicrobial Resistance in <i>Escherichia coli</i> from Food Animals in Lagos, Nigeria. <i>Microbial Drug Resistance</i> , <b>2015</b> , 21, 358-65	2.9	28
89	Comparison of <i>Salmonella enterica</i> serotype Infantis isolates from a veterinary teaching hospital. <i>Journal of Applied Microbiology</i> , <b>2007</b> , 102, 1527-36	4.7	28
88	In Vivo Transmission of an IncA/C Plasmid in <i>Escherichia coli</i> Depends on Tetracycline Concentration, and Acquisition of the Plasmid Results in a Variable Cost of Fitness. <i>Applied and Environmental Microbiology</i> , <b>2015</b> , 81, 3561-70	4.8	27
87	Comparison of <i>dkgB</i> -linked intergenic sequence ribotyping to DNA microarray hybridization for assigning serotype to <i>Salmonella enterica</i> . <i>FEMS Microbiology Letters</i> , <b>2012</b> , 337, 61-72	2.9	27
86	Related antimicrobial resistance genes detected in different bacterial species co-isolated from swine fecal samples. <i>Foodborne Pathogens and Disease</i> , <b>2011</b> , 8, 663-79	3.8	27
85	Sensitive and rapid molecular detection assays for <i>Salmonella enterica</i> serovars Typhimurium and Heidelberg. <i>Journal of Food Protection</i> , <b>2009</b> , 72, 2350-7	2.5	27
84	Rapid multiplex PCR and real-time TaqMan PCR assays for detection of <i>Salmonella enterica</i> and the highly virulent serovars Choleraesuis and Paratyphi C. <i>Journal of Clinical Microbiology</i> , <b>2008</b> , 46, 4018-22	9.7	27
83	Presence of the KPC carbapenemase gene in Enterobacteriaceae causing bacteremia and its correlation with in vitro carbapenem susceptibility. <i>Journal of Clinical Microbiology</i> , <b>2009</b> , 47, 239-41	9.7	24
82	Analysis of antimicrobial resistance genes detected in multiple-drug-resistant <i>Escherichia coli</i> isolates from broiler chicken carcasses. <i>Microbial Drug Resistance</i> , <b>2012</b> , 18, 453-63	2.9	23
81	Characteristics of plasmids in multi-drug-resistant Enterobacteriaceae isolated during prospective surveillance of a newly opened hospital in Iraq. <i>PLoS ONE</i> , <b>2012</b> , 7, e40360	3.7	23
80	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> , <b>2008</b> , 14, 251-8	2.9	22
79	<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> , <b>2008</b> , 55, 444-52	4.4	22
78	Prevalence and characterization of <i>Escherichia coli</i> isolated from the Upper Oconee Watershed in Northeast Georgia. <i>PLoS ONE</i> , <b>2018</b> , 13, e0197005	3.7	21
77	Microarray-based analysis of IncA/C plasmid-associated genes from multidrug-resistant <i>Salmonella enterica</i> . <i>Applied and Environmental Microbiology</i> , <b>2011</b> , 77, 6991-9	4.8	21
76	Molecular analysis of imipenem-resistant <i>Acinetobacter baumannii</i> isolated from US service members wounded in Iraq, 2003-2008. <i>Epidemiology and Infection</i> , <b>2012</b> , 140, 2302-7	4.3	21

75	Antimicrobial resistance, virulence determinants and genetic profiles of clinical and nonclinical <i>Enterococcus cecorum</i> from poultry. <i>Letters in Applied Microbiology</i> , <b>2015</b> , 60, 111-119	2.9	20
74	Use of a promiscuous, constitutively-active bacterial enhancer-binding protein to define the $\square$ (RpoN) regulon of <i>Salmonella</i> Typhimurium LT2. <i>BMC Genomics</i> , <b>2013</b> , 14, 602	4.5	20
73	Transmission of <i>Salmonella enterica</i> serotype Typhimurium in poultry with and without antimicrobial selective pressure. <i>Journal of Applied Microbiology</i> , <b>2006</b> , 101, 1301-8	4.7	19
72	Identification and sequences of the <i>Treponema pallidum</i> flhM, flhY, flhP, flhQ, flhR and flhBS genes. <i>Gene</i> , <b>1995</b> , 166, 57-64	3.8	19
71	Antimicrobial Resistance, Genetic Diversity and Multilocus Sequence Typing of from Humans, Retail Chicken and Ground Beef in Egypt. <i>Pathogens</i> , <b>2020</b> , 9,	4.5	18
70	Development of microarray and multiplex polymerase chain reaction assays for identification of serovars and virulence genes in <i>Salmonella enterica</i> of human or animal origin. <i>Journal of Veterinary Diagnostic Investigation</i> , <b>2010</b> , 22, 559-69	1.5	17
69	Prevalence and multidrug resistance of <i>Escherichia coli</i> from community-acquired infections in Lagos, Nigeria. <i>Journal of Infection in Developing Countries</i> , <b>2016</b> , 10, 920-931	2.3	17
68	An assay for determining the susceptibility of <i>Salmonella</i> isolates to commercial and household biocides. <i>PLoS ONE</i> , <b>2018</b> , 13, e0209072	3.7	17
67	Characterization of multidrug-resistant <i>Escherichia coli</i> by antimicrobial resistance profiles, plasmid replicon typing, and pulsed-field gel electrophoresis. <i>Microbial Drug Resistance</i> , <b>2011</b> , 17, 157-63	2.9	16
66	Prevalence of ColE1-like plasmids and kanamycin resistance genes in <i>Salmonella enterica</i> serovars. <i>Applied and Environmental Microbiology</i> , <b>2010</b> , 76, 6707-14	4.8	16
65	Detection of <i>Salmonella enterica</i> subpopulations by phenotype microarray antibiotic resistance patterns. <i>Applied and Environmental Microbiology</i> , <b>2007</b> , 73, 7753-6	4.8	16
64	Whole genome sequencing of multidrug-resistant serovar Typhimurium isolated from humans and poultry in Burkina Faso. <i>Tropical Medicine and Health</i> , <b>2018</b> , 46, 4	3.4	15
63	Relative survival of four serotypes of <i>Salmonella enterica</i> in low-water activity whey protein powder held at 36 and 70°C at various water activity levels. <i>Journal of Food Protection</i> , <b>2014</b> , 77, 1198-2005	2.5	15
62	Hydrogen-stimulated carbon acquisition and conservation in <i>Salmonella enterica</i> serovar Typhimurium. <i>Journal of Bacteriology</i> , <b>2011</b> , 193, 5824-32	3.5	15
61	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> , <b>2020</b> , 71, 3-25	2.9	14
60	Gene Expression Response of Serotype Enteritidis Phage Type 8 to Subinhibitory Concentrations of the Plant-Derived Compounds -Cinnamaldehyde and Eugenol. <i>Frontiers in Microbiology</i> , <b>2017</b> , 8, 1828	5.7	13
59	Genotypic and phenotypic correlations of multidrug-resistant <i>Acinetobacter baumannii</i> -A. calcoaceticus complex strains isolated from patients at the National Naval Medical Center. <i>Journal of Clinical Microbiology</i> , <b>2010</b> , 48, 4333-6	9.7	13
58	An oligonucleotide microarray to characterize multidrug resistant plasmids. <i>Journal of Microbiological Methods</i> , <b>2010</b> , 81, 96-100	2.8	12

57	Evidence of a conjugal erythromycin resistance element in the Lyme disease spirochete <i>Borrelia burgdorferi</i> . <i>International Journal of Antimicrobial Agents</i> , <b>2007</b> , 30, 496-504	14.3	12
56	Gene expression analysis of <i>Salmonella enterica</i> Enteritidis NaI(R) and <i>Salmonella enterica</i> Kentucky 3795 exposed to HCl and acetic acid in rich medium. <i>Foodborne Pathogens and Disease</i> , <b>2012</b> , 9, 331-7	3.8	11
55	Circulation of emerging NDM-5-producing <i>Escherichia coli</i> among humans and dogs in Egypt. <i>Zoonoses and Public Health</i> , <b>2020</b> , 67, 324-329	2.9	11
54	Identification, sequences, and expression of <i>Treponema pallidum</i> chemotaxis genes. <i>DNA Sequence</i> , <b>1997</b> , 7, 267-84		10
53	Novel DNA Binding and Regulatory Activities for $\sigma$ (RpoN) in <i>Salmonella enterica</i> Serovar Typhimurium 14028s. <i>Journal of Bacteriology</i> , <b>2017</b> , 199,	3.5	9
52	Transferable Plasmids of Associated With Antibiotic Resistance Genes. <i>Frontiers in Microbiology</i> , <b>2020</b> , 11, 562181	5.7	9
51	Plasmid Replicons and $\beta$ -Lactamase-Encoding Genes of Multidrug-Resistant Isolated from Humans and Food Animals in Lagos, Southwest Nigeria. <i>Microbial Drug Resistance</i> , <b>2019</b> , 25, 1410-1423	2.9	8
50	Genetic Characterization of Antimicrobial-Resistant Isolated from a Mixed-Use Watershed in Northeast Georgia, USA. <i>International Journal of Environmental Research and Public Health</i> , <b>2019</b> , 16,	4.6	8
49	Pathogenicity of dodecyltrimethylammonium chloride-resistant <i>Salmonella enterica</i> . <i>Applied and Environmental Microbiology</i> , <b>2013</b> , 79, 2371-6	4.8	8
48	Identification and sequences of the <i>Treponema pallidum</i> <i>mglA</i> and <i>mglC</i> genes. <i>DNA Sequence</i> , <b>1996</b> , 6, 293-8		8
47	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> , <b>2015</b> , 62, 479-88	2.9	7
46	Genomic comparison of diverse <i>Salmonella</i> serovars isolated from swine. <i>PLoS ONE</i> , <b>2019</b> , 14, e0224518	3.7	7
45	Characterization of small ColE1-like plasmids conferring kanamycin resistance in <i>Salmonella enterica</i> subsp. <i>enterica</i> serovars Typhimurium and Newport. <i>Plasmid</i> , <b>2010</b> , 63, 150-4	3.3	7
44	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> , <b>2018</b> , 14, 101-103	3.4	7
43	Comparison of Antimicrobial Resistance and Pan-Genome of Clinical and Non-Clinical from Poultry Using Whole-Genome Sequencing. <i>Foods</i> , <b>2020</b> , 9,	4.9	6
42	Sequence analysis of a group of low molecular-weight plasmids carrying multiple IS903 elements flanking a kanamycin resistance <i>aph</i> gene in <i>Salmonella enterica</i> serovars. <i>Plasmid</i> , <b>2011</b> , 65, 246-52	3.3	6
41	Amino acid "little Big Bang": representing amino acid substitution matrices as dot products of Euclidian vectors. <i>BMC Bioinformatics</i> , <b>2010</b> , 11, 4	3.6	6
40	Diversity and antimicrobial resistance of <i>Enterococcus</i> from the Upper Oconee Watershed, Georgia. <i>Journal of Applied Microbiology</i> , <b>2020</b> , 128, 1221-1233	4.7	6

39	Carriage and Gene Content Variability of the pESI-Like Plasmid Associated with Infantis Recently Established in United States Poultry Production. <i>Genes</i> , <b>2020</b> , 11,	4.2	6
38	Coproduction of Tet(X7) Conferring High-Level Tigecycline Resistance, Fosfomycin FosA4, and Colistin Mcr-1.1 in Escherichia coli Strains from Chickens in Egypt. <i>Antimicrobial Agents and Chemotherapy</i> , <b>2021</b> , 65,	5.9	6
37	Genotyping Campylobacter jejuni by comparative genome indexing: an evaluation with pulsed-field gel electrophoresis and flaA SVR sequencing. <i>Foodborne Pathogens and Disease</i> , <b>2009</b> , 6, 337-49	3.8	5
36	Identification and sequences of the Treponema pallidum flhA, flhF, and orf304 genes. <i>DNA Sequence</i> , <b>1997</b> , 7, 107-16		5
35	Genome Analysis of Multidrug-Resistant Isolated from Poultry in Nigeria. <i>Foodborne Pathogens and Disease</i> , <b>2020</b> , 17, 1-7	3.8	5
34	Detection of Salmonella Serotypes by Overnight Incubation of Entire Broiler Carcass. <i>Journal of Food Safety</i> , <b>2017</b> , 37, e12298	2	4
33	Carcass orientation and drip time affect potential surface water carryover for broiler carcasses subjected to a post-chill water dip or spray1. <i>Poultry Science</i> , <b>2017</b> , 96, 241-245	3.9	4
32	Incidence, species and antimicrobial resistance of naturally occurring Campylobacter isolates from quail carcasses sampled in a commercial processing facility. <i>Journal of Food Safety</i> , <b>2018</b> , 38, e12438	2	4
31	Expression and sequence analysis of a Treponema pallidum gene, tpn38(b), encoding an exported protein with homology to T. pallidum and Borrelia burgdorferi proteins. <i>FEMS Microbiology Letters</i> , <b>1996</b> , 135, 57-63	2.9	4
30	Draft Genome Sequence Analysis of Multidrug-Resistant Strains Isolated in 2013 from Humans and Chickens in Nigeria. <i>Genome Announcements</i> , <b>2017</b> , 5,		4
29	A newly developed Escherichia coli 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> , <b>2020</b> , 382, 120991	12.8	4
28	Sequences of the Salmonella typhimurium mglA and mglC genes. <i>Gene</i> , <b>1996</b> , 171, 131-2	3.8	3
27	Draft genome sequence of a bla- and bla-carrying multidrug-resistant Escherichia coli D-ST69 clinical isolate from Egypt. <i>Journal of Global Antimicrobial Resistance</i> , <b>2020</b> , 22, 832-834	3.4	3
26	Antimicrobial Resistance Gene Detection and Plasmid Typing Among Multidrug Resistant Enterococci Isolated from Freshwater Environment. <i>Microorganisms</i> , <b>2020</b> , 8,	4.9	3
25	Diversity of Plasmids and Genes Encoding Resistance to Extended-Spectrum Lactamase in from Different Animal Sources. <i>Microorganisms</i> , <b>2021</b> , 9,	4.9	3
24	Serotyping of sub-Saharan Africa Salmonella strains isolated from poultry feces using multiplex PCR and whole genome sequencing. <i>BMC Microbiology</i> , <b>2021</b> , 21, 29	4.5	3
23	Analysis of Campylobacter jejuni whole-genome DNA microarrays: significance of prophage and hypervariable regions for discriminating isolates. <i>Foodborne Pathogens and Disease</i> , <b>2012</b> , 9, 473-9	3.8	2
22	Genomic Analysis of Multidrug-Resistant from Surface Water in Northeast Georgia, United States: Presence of an ST131 Epidemic Strain Containing on a Phage-Like Plasmid. <i>Microbial Drug Resistance</i> , <b>2020</b> , 26, 447-455	2.9	2

21	Emergence of Multidrug-Resistant Producing CTX-M, MCR-1, and FosA in Retail Food From Egypt. <i>Frontiers in Cellular and Infection Microbiology</i> , <b>2021</b> , 11, 681588	5.9	2
20	Draft Genome Sequence of Salmonella enterica subsp. enterica Serovar Bardo Strain CRJJGF_00099 (Phylum Gammaproteobacteria). <i>Genome Announcements</i> , <b>2016</b> , 4,		2
19	Draft genome sequence of a human-associated streptogramin-resistant Staphylococcus aureus. <i>Journal of Global Antimicrobial Resistance</i> , <b>2019</b> , 16, 72-73	3.4	1
18	Draft Genome Sequence of Salmonella enterica subsp. enterica Serovar Orion Strain CRJJGF_00093 (Phylum Gammaproteobacteria). <i>Genome Announcements</i> , <b>2016</b> , 4,		1
17	Routes of transmission of Salmonella and Campylobacter in breeder turkeys. <i>Journal of Applied Poultry Research</i> , <b>2016</b> , 25, 591-609	2	1
16	Draft Genome Sequence of Salmonella enterica subsp. enterica Serovar Putten Strain CRJJGF_00159 (Phylum Gammaproteobacteria). <i>Genome Announcements</i> , <b>2016</b> , 4,		1
15	Draft Genome Sequence of Salmonella enterica subsp. enterica Serovar Blockley Strain CRJJGF_00147 (Phylum Gammaproteobacteria). <i>Genome Announcements</i> , <b>2016</b> , 4,		1
14	Draft Genome Sequence of Salmonella enterica subsp. enterica Serovar Kiambu Strain CRJJGF_00061 (Phylum Gammaproteobacteria). <i>Genome Announcements</i> , <b>2016</b> , 4,		1
13	Draft Genome Sequence of Salmonella enterica subsp. enterica Serovar Lille Strain CRJJGF_000101 (Phylum Gammaproteobacteria). <i>Genome Announcements</i> , <b>2016</b> , 4,		1
12	Draft Genome Sequence of Salmonella enterica subsp. enterica Serovar Widemarsh Strain CRJJGF_00058 (Phylum Gammaproteobacteria). <i>Genome Announcements</i> , <b>2016</b> , 4,		1
11	Whole-Genome Sequencing of Salmonella enterica subsp. enterica Serovar Cubana Strains Isolated from Agricultural Sources. <i>Genome Announcements</i> , <b>2014</b> , 2,		1
10	Antimicrobial interventions to reduce Salmonella and Campylobacter populations and improve shelf life of quail carcasses. <i>Poultry Science</i> , <b>2020</b> , 99, 5977-5982	3.9	1
9	Evaluation of a new chromogenic agar for the detection of environmental Enterococcus. <i>Journal of Microbiological Methods</i> , <b>2020</b> , 178, 106082	2.8	1
8	Draft Genome Sequence of Salmonella enterica subsp. diarizonae Serovar 61:k:1,5,(7) Strain CRJJGF_00165 (Phylum Gammaproteobacteria). <i>Genome Announcements</i> , <b>2016</b> , 4,		1
7	Analysis of Salmonella enterica 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> , <b>2022</b> , e0039322	4.8	0
6	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> , <b>2019</b> , 39, e12702	2	
5	Sequence of the Leptospira biflexa serovar patoc recA gene. <i>Gene</i> , <b>1995</b> , 167, 339-40	3.8	
4	Genomic comparison of diverse Salmonella serovars isolated from swine <b>2019</b> , 14, e0224518		



- 3 Genomic comparison of diverse Salmonella serovars isolated from swine **2019**, 14, e0224518
- 2 Genomic comparison of diverse Salmonella serovars isolated from swine **2019**, 14, e0224518
- 1 Genomic comparison of diverse Salmonella serovars isolated from swine **2019**, 14, e0224518