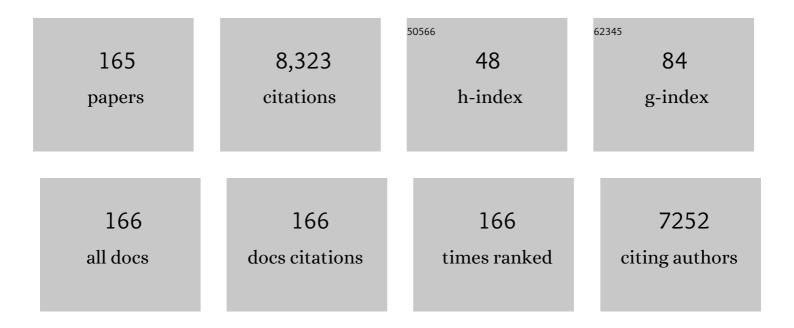
Jianghong Meng

List of Publications by Year in descending order

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IMNCHONG MENC

#	Article	IF	CITATIONS
1	Exploring the predictive capability of advanced machine learning in identifying severe disease phenotype in Salmonella enterica. Food Research International, 2022, 151, 110817.	2.9	10
2	Evaluation of the Persistence and Characterization of Listeria monocytogenes in Foodservice Operations. Foods, 2022, 11, 886.	1.9	2
3	Environmental and anthropogenic factors associated with the likelihood of detecting Salmonella in agricultural watersheds. Environmental Pollution, 2022, 306, 119298.	3.7	11
4	Polishing the Oxford Nanopore long-read assemblies of bacterial pathogens with Illumina short reads to improve genomic analyses. Genomics, 2021, 113, 1366-1377.	1.3	33
5	Diversity of Non-O157 Shiga Toxin-Producing Escherichia coli Isolated from Cattle from Central and Southern Chile. Animals, 2021, 11, 2388.	1.0	4
6	Genetic Determinants of Stress Resistance in Desiccated Salmonella enterica. Applied and Environmental Microbiology, 2021, 87, e0168321.	1.4	8
7	Persistence of Salmonella enterica and Enterococcus faecium NRRL B-2354 on Baby Spinach Subjected to Temperature Abuse after Exposure to Sub-Lethal Stresses. Foods, 2021, 10, 2141.	1.9	2
8	Whole-genome sequencing analysis of uncommon Shiga toxin-producing Escherichia coli from cattle: Virulence gene profiles, antimicrobial resistance predictions, and identification of novel O-serogroups. Food Microbiology, 2021, 99, 103821.	2.1	9
9	The Persistence of Bacterial Pathogens in Surface Water and Its Impact on Global Food Safety. Pathogens, 2021, 10, 1391.	1.2	21
10	Prevention of enteric bacterial infections and modulation of gut microbiota with conjugated linoleic acids producing <i>Lactobacillus</i> in mice. Gut Microbes, 2020, 11, 433-452.	4.3	24
11	Characterization of Mobile Genetic Elements Using Long-Read Sequencing for Tracking Listeria monocytogenes from Food Processing Environments. Pathogens, 2020, 9, 822.	1.2	11
12	Benchmarking hybrid assembly approaches for genomic analyses of bacterial pathogens using Illumina and Oxford Nanopore sequencing. BMC Genomics, 2020, 21, 631.	1.2	40
13	Benchmarking Long-Read Assemblers for Genomic Analyses of Bacterial Pathogens Using Oxford Nanopore Sequencing. International Journal of Molecular Sciences, 2020, 21, 9161.	1.8	26
14	Genomic analyses of multidrug-resistant Salmonella Indiana, Typhimurium, and Enteritidis isolates using MinION and MiSeq sequencing technologies. PLoS ONE, 2020, 15, e0235641.	1.1	16
15	Whole-Genome Phylogenetic Analysis Reveals a Wide Diversity of Non-O157 STEC Isolated From Ground Beef and Cattle Feces. Frontiers in Microbiology, 2020, 11, 622663.	1.5	6
16	<p>Molecular epidemiology and antimicrobial resistance of invasive non-typhoidal Salmonella in China, 2007–2016</p> . Infection and Drug Resistance, 2019, Volume 12, 2885-2897.	1.1	29
17	Characterization of internalin genes in Listeria monocytogenes from food and humans, and their association with the invasion of Caco-2 cells. Gut Pathogens, 2019, 11, 30.	1.6	30
18	Detection of 5â€ <scp>HMF</scp> in apple juice with artificial sensing systems. International Journal of Food Science and Technology, 2019, 54, 2679-2689.	1.3	3

#	Article	IF	CITATIONS
19	Formation of 5-hydroxymethylfurfural in industrial-scale apple juice concentrate processing. Food Control, 2019, 102, 56-68.	2.8	21
20	Analysis of virulence potential of Escherichia coli O145 isolated from cattle feces and hide samples based on whole genome sequencing. PLoS ONE, 2019, 14, e0225057.	1.1	5
21	Diversity of Serotype, Genotype, and Antibiotic Susceptibility of Salmonella Prevalent in Pickled Ready-to-Eat Meat. Frontiers in Microbiology, 2019, 10, 2577.	1.5	13
22	Shiga Toxin-ProducingEscherichia coli. , 2019, , 289-315.		5
23	Title is missing!. , 2019, 14, e0225057.		0
24	Title is missing!. , 2019, 14, e0225057.		0
25	Title is missing!. , 2019, 14, e0225057.		0
26	Title is missing!. , 2019, 14, e0225057.		0
27	Sequence Analysis of IncA/C and Incl1 Plasmids Isolated from Multidrug-Resistant <i>Salmonella</i> Newport Using Single-Molecule Real-Time Sequencing. Foodborne Pathogens and Disease, 2018, 15, 361-371.	0.8	16
28	Characterization of Toxin Genes and Antimicrobial Susceptibility of Staphylococcus aureus from Retail Raw Chicken Meat. Journal of Food Protection, 2018, 81, 528-533.	0.8	13
29	Salmonella enterica Phylogeny Based on Whole-Genome Sequencing Reveals Two New Clades and Novel Patterns of Horizontally Acquired Genetic Elements. MBio, 2018, 9, .	1.8	71
30	Increase in Ceftriaxone Resistance and Widespread Extended-Spectrum β-Lactamases Genes Among <i>Salmonella enterica</i> from Human and Nonhuman Sources. Foodborne Pathogens and Disease, 2018, 15, 770-775.	0.8	13
31	Whole genome shotgun sequencing revealed highly polymorphic genome regions and genes in Escherichia coli O157:H7 isolates collected from a single feedlot. PLoS ONE, 2018, 13, e0202775.	1.1	1
32	Proposed Epidemiological Cutoff Values for Ceftriaxone, Cefepime, and Colistin in <i>Salmonella</i> . Foodborne Pathogens and Disease, 2018, 15, 701-704.	0.8	7
33	Identification and Characterization of Conjugative Plasmids That Encode Ciprofloxacin Resistance in Salmonella. Antimicrobial Agents and Chemotherapy, 2018, 62, .	1.4	18
34	Metagenomic Analysis of Bacteria, Fungi, Bacteriophages, and Helminths in the Gut of Giant Pandas. Frontiers in Microbiology, 2018, 9, 1717.	1.5	55
35	Emerging high-level ciprofloxacin resistance and molecular basis of resistance in Salmonella enterica from humans, food and animals. International Journal of Food Microbiology, 2018, 280, 1-9.	2.1	43
36	Comparative Study on Antibiotic Resistance and DNA Profiles ofSalmonella entericaSerovar Typhimurium Isolated from Humans, Retail Foods, and the Environment in Shanghai, China. Foodborne Pathogens and Disease, 2018, 15, 481-488.	0.8	16

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37	Comparative genomics reveals differences in mobile virulence genes of Escherichia coli O103 pathotypes of bovine fecal origin. PLoS ONE, 2018, 13, e0191362.	1.1	15
38	Antibiotic Susceptibility and Molecular Screening of Class I Integron in <i>Salmonella</i> Isolates Recovered from Retail Raw Chicken Carcasses in China. Microbial Drug Resistance, 2017, 23, 230-235.	0.9	15
39	Draft Genome Sequences of Three Listeria monocytogenes Isolates from Foods in China. Genome Announcements, 2017, 5, .	0.8	2
40	Prevalence and Genomic Characterization of Escherichia coli O157:H7 in Cow-Calf Herds throughout California. Applied and Environmental Microbiology, 2017, 83, .	1.4	14
41	Draft Genome Sequences of Enteropathogenic Escherichia coli O103 Strains Isolated from Feces of Feedlot Cattle. Genome Announcements, 2017, 5, .	0.8	Ο
42	Distribution and Antimicrobial Susceptibility of Foodborne <i>Salmonella</i> Serovars in Eight Provinces in China from 2007 to 2012 (Except 2009). Foodborne Pathogens and Disease, 2017, 14, 393-399.	0.8	25
43	Draft Genome Sequences of Enterohemorrhagic Escherichia coli O103:H2 Strains Isolated from Feces of Feedlot Cattle. Genome Announcements, 2017, 5, .	0.8	0
44	Antimicrobial Susceptibility and Molecular Typing of Salmonella Senftenberg Isolated from Humans and Other Sources in Shanghai, China, 2005 to 2011. Journal of Food Protection, 2017, 80, 146-150.	0.8	6
45	Evolution and Diversity of Listeria monocytogenes from Clinical and Food Samples in Shanghai, China. Frontiers in Microbiology, 2016, 7, 1138.	1.5	26
46	Molecular Characterization, Antimicrobial Resistance and Caco-2 Cell Invasion Potential of Campylobacter jejuni/coli from Young Children with Diarrhea. Pediatric Infectious Disease Journal, 2016, 35, 330-334.	1.1	23
47	Virulence Gene Profiles and Clonal Relationships of Escherichia coli O26:H11 Isolates from Feedlot Cattle as Determined by Whole-Genome Sequencing. Applied and Environmental Microbiology, 2016, 82, 3900-3912.	1.4	41
48	Turtles as a Possible Reservoir of Nontyphoidal <i>Salmonella</i> in Shanghai, China. Foodborne Pathogens and Disease, 2016, 13, 428-433.	0.8	12
49	Molecular characterization and antimicrobial susceptibility of Listeria monocytogenes isolated from foods and humans. Food Control, 2016, 70, 96-102.	2.8	30
50	Prevalence and Antimicrobial Resistance Patterns of Diarrheagenic Escherichia coli in Shanghai, China. Pediatric Infectious Disease Journal, 2016, 35, 835-839.	1.1	11
51	Tracing Origins of the <i>Salmonella</i> Bareilly Strain Causing a Food-borne Outbreak in the United States. Journal of Infectious Diseases, 2016, 213, 502-508.	1.9	145
52	Escherichia coli O-Antigen Gene Clusters of Serogroups O62, O68, O131, O140, O142, and O163: DNA Sequences and Similarity between O62 and O68, and PCR-Based Serogrouping. Biosensors, 2015, 5, 51-68.	2.3	8
53	Complete Sequences of Six IncA/C Plasmids of Multidrug-Resistant Salmonella enterica subsp. enterica Serotype Newport. Genome Announcements, 2015, 3, .	0.8	18
54	Whole-Genome Sequences of 12 Clinical Strains of Listeria monocytogenes. Genome Announcements, 2015, 3, .	0.8	4

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55	Emergence of β-Lactamases and Extended-Spectrum β-Lactamases (ESBLs) Producing Salmonella in Retail Raw Chicken in China. Foodborne Pathogens and Disease, 2015, 12, 228-234.	0.8	16
56	Molecular analysis and antimicrobial susceptibility of enterotoxigenic Escherichia coli from diarrheal patients. Diagnostic Microbiology and Infectious Disease, 2015, 81, 126-131.	0.8	7
57	Virulence characterization of non-O157 Shiga toxin-producing Escherichia coli isolates from food, humans and animals. Food Microbiology, 2015, 50, 20-27.	2.1	25
58	Prevalence, Toxin Gene Profiles, and Antimicrobial Resistance of Staphylococcus aureus Isolated from Quick-Frozen Dumplings. Journal of Food Protection, 2015, 78, 218-223.	0.8	10
59	Prevalence of antimicrobial resistance of non-typhoidal Salmonella serovars in retail aquaculture products. International Journal of Food Microbiology, 2015, 210, 47-52.	2.1	49
60	Distribution and Molecular Characterization of Hypermutators in Retail Food in China. Journal of Food Protection, 2015, 78, 1481-1487.	0.8	13
61	Genome Sequences of 64 Non-O157:H7 Shiga Toxin-Producing Escherichia coli Strains. Genome Announcements, 2015, 3, .	0.8	4
62	Antimicrobial Resistance and Molecular Typing of <i>Salmonella</i> Stanley Isolated from Humans, Foods, and Environment. Foodborne Pathogens and Disease, 2015, 12, 945-949.	0.8	7
63	Antimicrobial susceptibility, virulence gene profiles and molecular subtypes of Salmonella Newport isolated from humans and other sources. Infection, Genetics and Evolution, 2015, 36, 294-299.	1.0	19
64	Prevalence and characterization of methicillin susceptible Staphylococcus aureus ST398 isolates from retail foods. International Journal of Food Microbiology, 2015, 196, 94-97.	2.1	24
65	Genomic Diversity and Virulence Profiles of Historical Escherichia coli O157 Strains Isolated from Clinical and Environmental Sources. Applied and Environmental Microbiology, 2015, 81, 569-577.	1.4	11
66	Molecular characterization of Salmonella enterica serovar Enteritidis on retail raw poultry in six provinces and two National cities in China. Food Microbiology, 2015, 46, 74-80.	2.1	47
67	Subtyping of Salmonella isolates on retail raw chicken in China by pulsed-field gel electrophoresis and plasmid analysis. Food Control, 2015, 47, 420-426.	2.8	6
68	Evaluation of loop-mediated isothermal amplification for the rapid, reliable, and robust detection of Salmonella in produce. Food Microbiology, 2015, 46, 485-493.	2.1	27
69	Prevalence, Antimicrobial Susceptibility, and Enterotoxin Gene Detection of Staphylococcus aureus Isolates in Ready-to-Eat Foods in Shaanxi, People's Republic of China. Journal of Food Protection, 2014, 77, 331-334.	0.8	21
70	Counts, Serotypes, and Antimicrobial Resistance of Salmonella Isolates on Retail Raw Poultry in the People's Republic of China. Journal of Food Protection, 2014, 77, 894-902.	0.8	51
71	Genetic Diversity of <i>Salmonella</i> Pathogenicity Islands SPI-5 and SPI-6 in <i>Salmonella</i> Newport. Foodborne Pathogens and Disease, 2014, 11, 798-807.	0.8	13
72	Comparative Genomic Analysis and Virulence Differences in Closely Related Salmonella enterica Serotype Heidelberg Isolates from Humans, Retail Meats, and Animals. Genome Biology and Evolution, 2014, 6, 1046-1068.	1.1	123

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73	Evaluation of a Loop-Mediated Isothermal Amplification Suite for the Rapid, Reliable, and Robust Detection of Shiga Toxin-Producing Escherichia coli in Produce. Applied and Environmental Microbiology, 2014, 80, 2516-2525.	1.4	23
74	Characterization of extended-spectrum beta-lactamases-producing Salmonella strains isolated from retail foods in Shaanxi and Henan Province, China. Food Microbiology, 2014, 42, 14-18.	2.1	26
75	Association of Clustered Regularly Interspaced Short Palindromic Repeat (CRISPR) Elements with Specific Serotypes and Virulence Potential of Shiga Toxin-Producing Escherichia coli. Applied and Environmental Microbiology, 2014, 80, 1411-1420.	1.4	41
76	Antimicrobial Susceptibility and Molecular Typing of Methicillin-Resistant <i>Staphylococcus aureus</i> in Retail Foods in Shaanxi, China. Foodborne Pathogens and Disease, 2014, 11, 281-286.	0.8	90
77	Antimicrobial Susceptibility and Molecular Typing of <i>Salmonella</i> Agona Isolated from Humans and Other Sources. Foodborne Pathogens and Disease, 2014, 11, 844-849.	0.8	5
78	Presence of <i>qnr</i> , <i>aac(6′)-lb</i> , <i>qep</i> A, <i>oqx</i> AB, and Mutations in Gyrase and Topoisomerase in Nalidixic Acid–Resistant <i>Salmonella</i> Isolates Recovered from Retail Chicken Carcasses. Foodborne Pathogens and Disease, 2014, 11, 698-705.	0.8	15
79	Presence of disinfectant resistance genes in Escherichia coli isolated from retail meats in the USA. Journal of Antimicrobial Chemotherapy, 2014, 69, 2644-2649.	1.3	102
80	Antimicrobial resistance of Shigella spp. from humans in Shanghai, China, 2004–2011. Diagnostic Microbiology and Infectious Disease, 2014, 78, 282-286.	0.8	27
81	Pathogenicity Islands in Shiga Toxin–ProducingEscherichia coliO26, O103, and O111 Isolates from Humans and Animals. Foodborne Pathogens and Disease, 2014, 11, 342-345.	0.8	9
82	Emergence and Prevalence of Non-H2S-Producing Salmonella enterica Serovar Senftenberg Isolates Belonging to Novel Sequence Type 1751 in China. Journal of Clinical Microbiology, 2014, 52, 2557-2565.	1.8	11
83	First Fully Closed Genome Sequence of Salmonella enterica subsp. enterica Serovar Cubana Associated with a Food-Borne Outbreak. Genome Announcements, 2014, 2, .	0.8	11
84	Isolation and Characterization of <i>Listeria monocytogenes</i> Isolates from Retail Foods in Shaanxi Province, China. Foodborne Pathogens and Disease, 2013, 10, 867-872.	0.8	15
85	Staphylococcus aureus and methicillin-resistant Staphylococcus aureus in retail raw chicken in China. Food Control, 2013, 29, 103-106.	2.8	57
86	Characterization and comparative analysis of a second thermonuclease from Staphylococcus aureus. Microbiological Research, 2013, 168, 174-182.	2.5	34
87	Analysis of pulsed field gel electrophoresis profiles using multiple enzymes for predicting potential source reservoirs for strains of Salmonella Enteritidis and Salmonella Typhimurium isolated from humans. Infection, Genetics and Evolution, 2013, 16, 226-233.	1.0	11
88	Molecular subtyping and virulence gene analysis of Listeria monocytogenes isolates from food. Food Microbiology, 2013, 35, 58-64.	2.1	69
89	Serotyping, antimicrobial susceptibility, pulse field gel electrophoresis analysis of Salmonella isolates from retail foods in Henan Province, China. Food Control, 2013, 32, 228-235.	2.8	64
90	Current Trends in Detecting Non-O157 Shiga Toxin–Producing <i>Escherichia coli</i> in Food. Foodborne Pathogens and Disease, 2013, 10, 665-677.	0.8	96

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91	Prevalence and Quantitative Detection of Salmonella in Retail Raw Chicken in Shaanxi, China. Journal of Food Protection, 2013, 76, 1958-1962.	0.8	16
92	Prevalence of Extended-Spectrum β-Lactamase–Producing Salmonella on Retail Chicken in Six Provinces and Two National Cities in the People's Republic of China. Journal of Food Protection, 2013, 76, 2040-2044.	0.8	32
93	Distribution of Pathogenicity Islands OI-122, OI-43/48, and OI-57 and a High-Pathogenicity Island in Shiga Toxin-Producing Escherichia coli. Applied and Environmental Microbiology, 2013, 79, 3406-3412.	1.4	31
94	Molecular Serogrouping of Shiga Toxin–ProducingEscherichia coliUsing Suspension Array. Foodborne Pathogens and Disease, 2013, 10, 478-480.	0.8	9
95	Rapid Identification and Differentiation of Non-O157 Shiga Toxin–ProducingEscherichia coliUsing Polymerase Chain Reaction Coupled to Electrospray Ionization Mass Spectrometry. Foodborne Pathogens and Disease, 2013, 10, 737-743.	0.8	8
96	Phylogenetics and Differentiation of Salmonella Newport Lineages by Whole Genome Sequencing. PLoS ONE, 2013, 8, e55687.	1.1	63
97	Effects of Tomato Variety, Temperature Differential, and Post–Stem Removal Time on Internalization of Salmonella Enterica Serovar Thompson in Tomatoesâ€. Journal of Food Protection, 2012, 75, 297-303.	0.8	40
98	Draft Genome Sequences of Eight Salmonella enterica Serotype Newport Strains from Diverse Hosts and Locations. Journal of Bacteriology, 2012, 194, 5146-5146.	1.0	10
99	Phylogenetic Analysis of Non-O157 Shiga Toxin-Producing Escherichia coli Strains by Whole-Genome Sequencing. Journal of Clinical Microbiology, 2012, 50, 4123-4127.	1.8	39
100	Non-O157 Shiga toxin-producing Escherichia coli in retail ground beef and pork in the Washington D.C. area. Food Microbiology, 2012, 32, 371-377.	2.1	52
101	Antimicrobial Susceptibility Testing and Genotypic Characterization of <i>Staphylococcus aureus</i> from Food and Food Animals. Foodborne Pathogens and Disease, 2012, 9, 95-101.	0.8	24
102	Inactivation of foodborne pathogens in raw milk using high hydrostatic pressure. Food Control, 2012, 28, 273-278.	2.8	70
103	Mutations in gyrase and topoisomerase genes associated with fluoroquinolone resistance in Salmonella serovars from retail meats. Food Research International, 2012, 45, 935-939.	2.9	26
104	Characterization of Staphylococcus aureus isolated from powdered infant formula milk and infant rice cereal in China. International Journal of Food Microbiology, 2012, 153, 142-147.	2.1	70
105	Differential Gene Expression by RamA in Ciprofloxacin-Resistant Salmonella Typhimurium. PLoS ONE, 2011, 6, e22161.	1.1	27
106	Identification of a Salmonellosis Outbreak by Means of Molecular Sequencing. New England Journal of Medicine, 2011, 364, 981-982.	13.9	155
107	Simultaneous Analysis of Multiple Enzymes Increases Accuracy of Pulsed-Field Gel Electrophoresis in Assigning Genetic Relationships among Homogeneous Salmonella Strains. Journal of Clinical Microbiology, 2011, 49, 85-94.	1.8	35
108	Identification and Antimicrobial Resistance of Extraintestinal Pathogenic Escherichia coli from Retail Meats. Journal of Food Protection, 2011, 74, 38-44.	0.8	36

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109	Prevalence of Salmonella on Raw Poultry at Retail Markets in China. Journal of Food Protection, 2011, 74, 1724-1728.	0.8	82
110	Antimicrobial Susceptibility of Staphylococcus aureus from Retail Ground Meats. Journal of Food Protection, 2011, 74, 1625-1629.	0.8	38
111	Osmoregulated periplasmic glucans synthesis gene family of Shigella flexneri. Archives of Microbiology, 2010, 192, 167-174.	1.0	8
112	Prevalence and characterization of Salmonella serovars in retail meats of marketplace in Shaanxi, China. International Journal of Food Microbiology, 2010, 141, 63-72.	2.1	238
113	Presence and Characterization of Shiga Toxin-Producing <i>Escherichia coli</i> and Other Potentially Diarrheagenic <i>E. coli</i> Strains in Retail Meats. Applied and Environmental Microbiology, 2010, 76, 1709-1717.	1.4	64
114	Osmoregulated periplasmic glucans of Salmonella enterica serovar Typhimurium are required for optimal virulence in mice. Microbiology (United Kingdom), 2009, 155, 229-237.	0.7	48
115	Effect of transcriptional activators RamA and SoxS on expression of multidrug efflux pumps AcrAB and AcrEF in fluoroquinolone-resistant Salmonella Typhimurium. Journal of Antimicrobial Chemotherapy, 2009, 63, 95-102.	1.3	63
116	Characterization of Salmonella isolates from retail foods based on serotyping, pulse field gel electrophoresis, antibiotic resistance and other phenotypic properties. International Journal of Food Microbiology, 2009, 129, 93-98.	2.1	24
117	Osmoregulated periplasmic glucans are needed for competitive growth and biofilm formation by <i>Salmonella enterica</i> serovar Typhimurium in leafy-green vegetable wash waters and colonization in mice. FEMS Microbiology Letters, 2009, 292, 13-20.	0.7	21
118	Characterisation of antimicrobial resistance-associated integrons and mismatch repair gene mutations in Salmonella serotypes. International Journal of Antimicrobial Agents, 2009, 33, 120-124.	1.1	20
119	Fitness cost of macrolide resistance in Campylobacter jejuni. International Journal of Antimicrobial Agents, 2009, 34, 462-466.	1.1	43
120	Identification and Characterization of Shiga Toxin Type 2 Variants in <i>Escherichia coli</i> Isolates from Animals, Food, and Humans. Applied and Environmental Microbiology, 2008, 74, 5645-5652.	1.4	39
121	<i>Campylobacter</i> -Induced Interleukin-8 Secretion in Polarized Human Intestinal Epithelial Cells Requires <i>Campylobacter</i> -Secreted Cytolethal Distending Toxin- and Toll-Like Receptor-Mediated Activation of NF-I [®] B. Infection and Immunity, 2008, 76, 4498-4508.	1.0	109
122	An Enhanced Discriminatory Pulsed-Field Gel Electrophoresis Scheme for Subtyping Salmonella Serotypes Heidelberg, Kentucky, SaintPaul, and Hadar. Journal of Food Protection, 2008, 71, 2067-2072.	0.8	20
123	Contribution of Target Gene Mutations and Efflux to Decreased Susceptibility of Salmonella enterica Serovar Typhimurium to Fluoroquinolones and Other Antimicrobials. Antimicrobial Agents and Chemotherapy, 2007, 51, 535-542.	1.4	137
124	Identification and Antimicrobial Susceptibility of Lactic Acid Bacteria from Retail Fermented Foods. Journal of Food Protection, 2007, 70, 2606-2612.	0.8	26
125	Enhanced Subtyping Scheme for <i>Salmonella</i> Enteritidis. Emerging Infectious Diseases, 2007, 13, 1932-1935.	2.0	55
126	Characterization of Listeria monocytogenes isolated from retail foods. International Journal of Food Microbiology, 2007, 113, 47-53.	2.1	112

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127	Heterogeneity of vat(E)-carrying plasmids in Enterococcus faecium recovered from human and animal sources. International Journal of Antimicrobial Agents, 2006, 28, 200-205.	1.1	5
128	Adherence to and Invasion of Human Intestinal Epithelial Cells by Campylobacter jejuni and Campylobacter coli Isolates from Retail Meat Products. Journal of Food Protection, 2006, 69, 768-774.	0.8	72
129	AN IMPROVED METHOD FOR RAPID ISOLATION OF SALMONELLA AGAINST PROTEUS IN CHICKEN CARCASSES. Journal of Food Safety, 2006, 26, 49-61.	1.1	14
130	Isolation and Characterization of Listeria monocytogenes Isolates from Ready-To-Eat Foods in Florida. Applied and Environmental Microbiology, 2006, 72, 5073-5076.	1.4	56
131	Antimicrobial susceptibility and molecular characterization of avian pathogenic Escherichia coli isolates. Veterinary Microbiology, 2005, 107, 215-224.	0.8	124
132	Use of Ramification Amplification Assay for Detection of Escherichia coli O157:H7 and Other E. coli Shiga Toxin-Producing Strains. Journal of Clinical Microbiology, 2005, 43, 6086-6090.	1.8	34
133	Role of Efflux Pumps and Topoisomerase Mutations in Fluoroquinolone Resistance in Campylobacter jejuni and Campylobacter coli. Antimicrobial Agents and Chemotherapy, 2005, 49, 3347-3354.	1.4	131
134	Prevalence and Antimicrobial Resistance of Campylobacter spp. and Salmonella Serovars in Organic Chickens from Maryland Retail Stores. Applied and Environmental Microbiology, 2005, 71, 4108-4111.	1.4	186
135	Identification of antimicrobial resistance and class 1 integrons in Shiga toxin-producing Escherichia coli recovered from humans and food animals. Journal of Antimicrobial Chemotherapy, 2005, 56, 216-219.	1.3	79
136	A DNA microarray for identification of virulence and antimicrobial resistance genes in Salmonella serovars and Escherichia coli. Molecular and Cellular Probes, 2005, 19, 195-201.	0.9	72
137	Evaluation of Molecular Typing Methods for Escherichia coli O157:H7 Isolates from Cattle, Food, and Humans. Journal of Food Protection, 2004, 67, 651-657.	0.8	43
138	Characterization of Multiple-Antimicrobial-Resistant Escherichia coli Isolates from Diseased Chickens and Swine in China. Journal of Clinical Microbiology, 2004, 42, 3483-3489.	1.8	210
139	Retail meat and poultry as a reservoir of antimicrobial-resistant Escherichia coli. Food Microbiology, 2004, 21, 249-255.	2.1	68
140	Characterization of Multiple-Antimicrobial-Resistant Salmonella Serovars Isolated from Retail Meats. Applied and Environmental Microbiology, 2004, 70, 1-7.	1.4	387
141	Isolation of antimicrobial-resistant Escherichia coli from retail meats purchased in Greater Washington, DC, USA. International Journal of Food Microbiology, 2003, 85, 197-202.	2.1	97
142	Antimicrobial-Resistant Campylobacter Species from Retail Raw Meats. Applied and Environmental Microbiology, 2003, 69, 3005-3007.	1.4	132
143	Isolation and Characterization of Escherichia coli Recovered from Maryland Apple Cider and the Cider Production Environment. Journal of Food Protection, 2003, 66, 2237-2244.	0.8	7
144	Comparison of the Etest and agar dilution for in vitro antimicrobial susceptibility testing of Campylobacter. Journal of Antimicrobial Chemotherapy, 2002, 50, 487-494.	1.3	71

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145	Antimicrobial Resistance of Escherichia coli O157 Isolated from Humans, Cattle, Swine, and Food. Applied and Environmental Microbiology, 2002, 68, 576-581.	1.4	245
146	A PCR–ELISA for detecting Shiga toxin-producing Escherichia coli. Microbes and Infection, 2002, 4, 285-290.	1.0	39
147	Introduction. Microbiological food safety. Microbes and Infection, 2002, 4, 395-397.	1.0	79
148	Antimicrobial Resistance of <i>Escherichia coli</i> O26, O103, O111, O128, and O145 from Animals and Humans. Emerging Infectious Diseases, 2002, 8, 1409-1414.	2.0	106
149	The Isolation of Antibiotic-Resistant Salmonella from Retail Ground Meats. New England Journal of Medicine, 2001, 345, 1147-1154.	13.9	442
150	Prevalence of Campylobacter spp. , Escherichia coli , and Salmonella Serovars in Retail Chicken, Turkey, Pork, and Beef from the Greater Washington, D.C., Area. Applied and Environmental Microbiology, 2001, 67, 5431-5436.	1.4	500
151	Identification and Characterization of Integron-Mediated Antibiotic Resistance among Shiga Toxin-Producing Escherichia coli Isolates. Applied and Environmental Microbiology, 2001, 67, 1558-1564.	1.4	201
152	Availability of Glutamate and Arginine during Acid Challenge Determines Cell Density-Dependent Survival Phenotype of Escherichia coli Strains. Applied and Environmental Microbiology, 2001, 67, 4914-4918.	1.4	74
153	Identification and Expression of Cephamycinase bla CMY Genes in Escherichia coli and Salmonella Isolates from Food Animals and Ground Meat. Antimicrobial Agents and Chemotherapy, 2001, 45, 3647-3650.	1.4	190
154	Genomic typing of Escherichia coli O157:H7 by semi-automated fluorescent AFLP analysis. Microbes and Infection, 2000, 2, 107-113.	1.0	50
155	Use of Hazard Analysis Critical Control Point and Alternative Treatments in the Production of Apple Cider. Journal of Food Protection, 1999, 62, 778-785.	0.8	24
156	Virulence genes of Shiga toxin-producing Escherichia coli isolated from food, animals and humans. International Journal of Food Microbiology, 1998, 45, 229-235.	2.1	85
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#	Article	IF	CITATIONS
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