Chen Moutong

List of Publications by Year in descending order

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77	1,735	24 h-index	37
papers	citations		g-index
78	78	78	1397
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	An ultrasensitive CRISPR/Cas12a based electrochemical biosensor for Listeria monocytogenes detection. Biosensors and Bioelectronics, 2021, 179, 113073.	5.3	151
2	Castor oil-based cationic waterborne polyurethane dispersions: Storage stability, thermo-physical properties and antibacterial properties. Industrial Crops and Products, 2018, 117, 169-178.	2.5	121
3	A Study on Prevalence and Characterization of Bacillus cereus in Ready-to-Eat Foods in China. Frontiers in Microbiology, 2019, 10, 3043.	1.5	84
4	Listeria monocytogenes Prevalence and Characteristics in Retail Raw Foods in China. PLoS ONE, 2015, 10, e0136682.	1.1	81
5	Analysis of Multilocus Sequence Typing and Virulence Characterization of Listeria monocytogenes Isolates from Chinese Retail Ready-to-Eat Food. Frontiers in Microbiology, 2016, 7, 168.	1.5	75
6	Prevalence and characterization of Listeria monocytogenes isolated from retail-level ready-to-eat foods in South China. Food Control, 2014, 38, 1-7.	2.8	69
7	Abundant and Diverse RNA Viruses in Insects Revealed by RNA-Seq Analysis: Ecological and Evolutionary Implications. MSystems, 2020, 5, .	1.7	66
8	Isolation, Potential Virulence, and Population Diversity of Listeria monocytogenes From Meat and Meat Products in China. Frontiers in Microbiology, 2019, 10, 946.	1.5	57
9	Occurrence, Antibiotic Resistance, and Population Diversity of Listeria monocytogenes Isolated From Fresh Aquatic Products in China. Frontiers in Microbiology, 2018, 9, 2215.	1.5	51
10	Prevalence, antibiotic resistance and genetic diversity of Listeria monocytogenes isolated from retail ready-to-eat foods in China. Food Control, 2015, 47, 340-347.	2.8	50
11	Prevalence, Potential Virulence, and Genetic Diversity of Listeria monocytogenes Isolates From Edible Mushrooms in Chinese Markets. Frontiers in Microbiology, 2018, 9, 1711.	1.5	48
12	Cas12aFDet: A CRISPR/Cas12a-based fluorescence platform for sensitive and specific detection of Listeria monocytogenes serotype 4c. Analytica Chimica Acta, 2021, 1151, 338248.	2.6	44
13	Prevalence, enumeration, and pheno- and genotypic characteristics of Listeria monocytogenes isolated from raw foods in South China. Frontiers in Microbiology, 2015, 6, 1026.	1.5	43
14	Polysaccharide from Agrocybe cylindracea prevents diet-induced obesity through inhibiting inflammation mediated by gut microbiota and associated metabolites. International Journal of Biological Macromolecules, 2022, 209, 1430-1438.	3.6	36
15	Prevalence, Antibiotic Susceptibility, and Molecular Characterization of Cronobacter spp. Isolated From Edible Mushrooms in China. Frontiers in Microbiology, 2019, 10, 283.	1.5	35
16	Isolation and Characterization of a Novel Salmonella Phage vB_SalP_TR2. Frontiers in Microbiology, 2021, 12, 664810.	1.5	35
17	Prevalence, virulence, antimicrobial resistance, and molecular characterization of fluoroquinolone resistance of Vibrio parahaemolyticus from different types of food samples in China. International Journal of Food Microbiology, 2020, 317, 108461.	2.1	33
18	Genetic characteristics and virulence of Listeria monocytogenes isolated from fresh vegetables in China. BMC Microbiology, 2019, 19, 119.	1.3	31

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19	Food-Borne Vibrio parahaemolyticus in China: Prevalence, Antibiotic Susceptibility, and Genetic Characterization. Frontiers in Microbiology, 2020, 11, 1670.	1.5	31
20	Campylobacter jejuni Biofilm Formation Under Aerobic Conditions and Inhibition by ZnO Nanoparticles. Frontiers in Microbiology, 2020, 11, 207.	1.5	31
21	First detection of the plasmid-mediated colistin resistance gene mcr-1 in virulent Vibrio parahaemolyticus. International Journal of Food Microbiology, 2019, 308, 108290.	2.1	28
22	Prevalence and Contamination Patterns of <i>Listeria monocytogenes</i> in <i>Flammulina velutipes</i> Plants. Foodborne Pathogens and Disease, 2014, 11, 620-627.	0.8	27
23	Prevalence and Genetic Diversity of Enterococcus faecalis Isolates from Mineral Water and Spring Water in China. Frontiers in Microbiology, 2017, 8, 1109.	1.5	27
24	Phenotypic and genotypic characterization of PVL-positive Staphylococcus aureus isolated from retail foods in China. International Journal of Food Microbiology, 2019, 304, 119-126.	2.1	26
25	Comparative Genomic Analysis Reveals the Potential Risk of Vibrio parahaemolyticus Isolated From Ready-To-Eat Foods in China. Frontiers in Microbiology, 2019, 10, 186.	1.5	25
26	Quantitative detection of aflatoxin B1 using quantum dots-based immunoassay in a recyclable gravity-driven microfluidic chip. Biosensors and Bioelectronics, 2021, 190, 113394.	5. 3	22
27	Prevalence, genetic analysis and CRISPR typing of Cronobacter spp. isolated from meat and meat products in China. International Journal of Food Microbiology, 2020, 321, 108549.	2.1	21
28	The Genomic Context for the Evolution and Transmission of Community-Associated Staphylococcus aureus ST59 Through the Food Chain. Frontiers in Microbiology, 2020, 11, 422.	1.5	21
29	Cronobacter spp. isolated from aquatic products in China: Incidence, antibiotic resistance, molecular characteristic and CRISPR diversity. International Journal of Food Microbiology, 2020, 335, 108857.	2.1	19
30	Amplified electrochemical antibiotic aptasensing based on electrochemically deposited AuNPs coordinated with PEI-functionalized Fe-based metal-organic framework. Mikrochimica Acta, 2021, 188, 286.	2.5	19
31	Evaluation of the Cholesterol-Lowering Mechanism of Enterococcus faecium Strain 132 and Lactobacillus paracasei Strain 201 in Hypercholesterolemia Rats. Nutrients, 2021, 13, 1982.	1.7	16
32	High prevalence of multidrug-resistant Escherichia coli and first detection of IncHI2/IncX4-plasmid carrying mcr-1 E. coli in retail ready-to-eat foods in China. International Journal of Food Microbiology, 2021, 355, 109349.	2.1	15
33	Novel phage vB_CtuP_B1 for controlling Cronobacter malonaticus and Cronobacter turicensis in ready-to-eat lettuce and powered infant formula. Food Research International, 2021, 143, 110255.	2.9	14
34	Development of a novel RAA-based microfluidic chip for absolute quantitative detection of human norovirus. Microchemical Journal, 2021, 164, 106050.	2.3	12
35	Presence and characterization of methicillin-resistant Staphylococcus aureus co-carrying the multidrug resistance genes cfr and Isa(E) in retail food in China. International Journal of Food Microbiology, 2022, 363, 109512.	2.1	12
36	Characterization of the Novel Phage vB_VpaP_FE11 and Its Potential Role in Controlling Vibrio parahaemolyticus Biofilms. Viruses, 2022, 14, 264.	1.5	12

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37	Characteristics of Antibiotic Resistance Genes and Antibiotic-Resistant Bacteria in Full-Scale Drinking Water Treatment System Using Metagenomics and Culturing. Frontiers in Microbiology, 2021, 12, 798442.	1.5	12
38	Rapid detection of Listeria monocytogenes sequence type 121 strains using a novel multiplex PCR assay. LWT - Food Science and Technology, 2019, 116, 108474.	2.5	11
39	Genome- and Proteome-Wide Analysis of Lysine Acetylation in Vibrio vulnificus Vv180806 Reveals Its Regulatory Roles in Virulence and Antibiotic Resistance. Frontiers in Microbiology, 2020, 11, 591287.	1.5	11
40	Cronobacter sakazakii, Cronobacter malonaticus, and Cronobacter dublinensis Genotyping Based on CRISPR Locus Diversity. Frontiers in Microbiology, 2019, 10, 1989.	1.5	10
41	Real-time PCR identification of Listeria monocytogenes serotype 4c using primers for novel target genes obtained by comparative genomic analysis. LWT - Food Science and Technology, 2021, 138, 110774.	2.5	10
42	Distribution, contamination routes, and seasonal influence of persistent Listeria monocytogenes in a commercial fresh Hypsizigus marmoreus production facility. Food Control, 2021, 127, 108118.	2.8	10
43	Proteomic analysis of trichloroethylene-induced alterations in expression, distribution, and interactions of SET/TAF-ll± and two SET/TAF-ll±-binding proteins, eEF1A1 and eEF1A2, in hepatic L-02 cells. Toxicology and Applied Pharmacology, 2012, 263, 259-272.	1.3	9
44	Multiplex PCR for the Identification of Pathogenic Listeria in Flammulina velutipes Plant Based on Novel Specific Targets Revealed by Pan-Genome Analysis. Frontiers in Microbiology, 2020, 11, 634255.	1.5	9
45	Pediococcus pentosaceus IM96 Exerts Protective Effects against Enterohemorrhagic Escherichia coli O157:H7 Infection In Vivo. Foods, 2021, 10, 2945.	1.9	9
46	Exploration of the Molecular Mechanisms Underlying the Anti-Photoaging Effect of Limosilactobacillus fermentum XJC60. Frontiers in Cellular and Infection Microbiology, 2022, 12, 838060.	1.8	9
47	A database for risk assessment and comparative genomic analysis of foodborne Vibrio parahaemolyticus in China. Scientific Data, 2020, 7, 321.	2.4	8
48	Mining of novel target genes through pan-genome analysis for multiplex PCR differentiation of the major Listeria monocytogenes serotypes. International Journal of Food Microbiology, 2021, 339, 109026.	2.1	8
49	Identification of Novel Sensitive and Reliable Serovar-Specific Targets for PCR Detection of Salmonella Serovars Hadar and Albany by Pan-Genome Analysis. Frontiers in Microbiology, 2021, 12, 605984.	1.5	8
50	Molecular characterisation of antimicrobial resistance determinants and class 1 integrons of Salmonella enterica subsp. enterica serotype Enteritidis strains from retail food in China. Food Control, 2021, 128, 108191.	2.8	8
51	A microfluidic genoserotyping strategy for fast and objective identification of common Salmonella serotypes isolated from retail food samples in China. Analytica Chimica Acta, 2022, 1201, 339657.	2.6	8
52	Microbial Communities and Physiochemical Properties of Four Distinctive Traditionally Fermented Vegetables from North China and Their Influence on Quality and Safety. Foods, 2022, 11, 21.	1.9	8
53	Whole <i>Agrocybe cylindracea</i> Prevented Obesity Linking with Modification of Gut Microbiota and Associated Fecal Metabolites in Highâ€Fat Dietâ€Fed Mice. Molecular Nutrition and Food Research, 2022, 66, e2100897.	1.5	7
54	Exploration of the molecular mechanisms underlying the antibiotic resistance of <i>Helicobacter pylori</i> : A wholeâ€genome sequencingâ€based study in Southern China. Helicobacter, 2022, 27, e12879.	1.6	7

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55	Characterization of class 1 integrons harboring blaVEB-1 in Vibrio parahaemolyticus isolated from ready-to-eat foods in China. International Journal of Food Microbiology, 2020, 318, 108473.	2.1	6
56	Identification of the Potential Biological Preservative Tetramycin A-Producing Strain and Enhancing Its Production. Frontiers in Microbiology, 2019, 10, 2925.	1.5	6
57	Genetic Diversity and Population Structure of Vibrio parahaemolyticus Isolated From Clinical and Food Sources. Frontiers in Microbiology, 2021, 12, 708795.	1.5	6
58	Genomic Analysis and Stability Evaluation of the Phenol-Degrading Bacterium Acinetobacter sp. DW-1 During Water Treatment. Frontiers in Microbiology, 2021, 12, 687511.	1.5	6
59	Microbial Communities and Physicochemical Characteristics of Traditional Dajiang and Sufu in North China Revealed by High-Throughput Sequencing of 16S rRNA. Frontiers in Microbiology, 2021, 12, 665243.	1.5	6
60	A novel multiplex PCR method for simultaneous identification of hypervirulent Listeria monocytogenes clonal complex 87 and CC88 strains in China. International Journal of Food Microbiology, 2022, 366, 109558.	2.1	6
61	Establishment and application of a rapid visual detection method for <i>Listeria monocytogenes</i> based on polymerase spiral reaction (PSR). Bioengineered, 2022, 13, 7860-7867.	1.4	6
62	Identification of new serovar-specific detection targets against salmonella B serogroup using large-scale comparative genomics. Food Control, 2021, 124, 107862.	2.8	5
63	Trichloroethylene induces biphasic concentration-dependent changes in cell proliferation and the expression of SET-associated proteins in human hepatic L-02 cells. Biomedical and Environmental Sciences, 2013, 26, 618-21.	0.2	5
64	A Novel Gene vp0610 Negatively Regulates Biofilm Formation in Vibrio parahaemolyticus. Frontiers in Microbiology, 2021, 12, 656380.	1.5	4
65	Development of a High-Efficiency Immunomagnetic Enrichment Method for Detection of Human Norovirus via PAMAM Dendrimer/SA-Biotin Mediated Cascade-Amplification. Frontiers in Microbiology, 2021, 12, 673872.	1.5	4
66	Evolutionary Divergence of the Novel Staphylococcal Species Staphylococcus argenteus. Frontiers in Microbiology, 2021, 12, 769642.	1.5	4
67	Determination of Antiviral Mechanism of Centenarian Gut-Derived Limosilactobacillus fermentum Against Norovirus. Frontiers in Nutrition, 2022, 9, 812623.	1.6	4
68	Presence and Characterization of a Novel cfr-Carrying Tn558 Transposon Derivative in Staphylococcus delphini Isolated From Retail Food. Frontiers in Microbiology, 2020, 11, 598990.	1.5	3
69	Development and Application of a Novel Rapid and Throughput Method for Broad-Spectrum Anti-Foodborne Norovirus Antibody Testing. Frontiers in Microbiology, 2021, 12, 670488.	1.5	3
70	Recent Advances in Glycosidase Probes Used in Escherichia Coli Detection. Current Medicinal Chemistry, 2021, 28, 5386-5410.	1.2	2
71	Pseudomonas protegens FJKB0103 Isolated from Rhizosphere Exhibits Anti-Methicillin-Resistant Staphylococcus aureus Activity. Microorganisms, 2022, 10, 315.	1.6	2
72	Pseudotargeted Metabolomic Fingerprinting and Deep Learning for Identification and Visualization of Common Pathogens. Frontiers in Microbiology, 2022, 13, 830832.	1.5	2

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73	Evolutionary Mechanism of Immunological Cross-Reactivity Between Different GII.17 Variants. Frontiers in Microbiology, 2021, 12, 653719.	1.5	1
74	Editorial: The Prevalence of MDR Non-Fermenting Gram Negative Bacteria and Their Chemotherapy. Frontiers in Microbiology, 2021, 12, 664336.	1.5	1
75	Isolation and Characterization of Non-O157 Shiga Toxin–Producing Escherichia coli in Foods Sold at Retail Markets in China. Journal of Food Protection, 2020, 83, 460-466.	0.8	1
76	Imbalanced Dermic Microbiome Aggravates Inflammation in Toenail Paronychia. Frontiers in Cellular and Infection Microbiology, 2021, 11, 781927.	1.8	1
77	Molecular Characterization of Rifampicin-Resistant Staphylococcus aureus Isolates from Retail Foods in China. Antibiotics, 2021, 10, 1487.	1.5	1