List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Staphylococcal chromosomal cassettes mec (SCCmec): A mobile genetic element in methicillin-resistant Staphylococcus aureus. Microbial Pathogenesis, 2016, 101, 56-67.	1.3	197
2	Crystal Violet and XTT Assays on Staphylococcus aureus Biofilm Quantification. Current Microbiology, 2016, 73, 474-482.	1.0	188
3	Vulvovaginal Candidiasis: A Current Understanding and Burning Questions. Journal of Fungi (Basel,) Tj ETQq1 1	0.784314 1.5	rgBT /Overloc 156
4	Development and application of a loop-mediated isothermal amplification method on rapid detection Escherichia coli O157 strains from food samples. Molecular Biology Reports, 2010, 37, 2183-2188.	1.0	149
5	Occurrence and Characteristics of Class 1 and 2 Integrons in <i>Pseudomonas aeruginosa</i> Isolates from Patients in Southern China. Journal of Clinical Microbiology, 2009, 47, 230-234.	1.8	132
6	Development and application of loop-mediated isothermal amplification assays on rapid detection of various types of staphylococci strains. Food Research International, 2012, 47, 166-173.	2.9	129
7	Class 1 integron in staphylococci. Molecular Biology Reports, 2011, 38, 5261-5279.	1.0	111
8	Viable but non-culturable state and toxin gene expression of enterohemorrhagic Escherichia coli O157 under cryopreservation. Research in Microbiology, 2017, 168, 188-193.	1.0	110
9	Integron-bearing methicillin-resistant coagulase-negative staphylococci in South China, 2001–2004. FEMS Microbiology Letters, 2008, 278, 223-230.	0.7	108
10	Development and application of a simple loop-mediated isothermal amplification method on rapid detection of Listeria monocytogenes strains. Molecular Biology Reports, 2012, 39, 445-449.	1.0	104
11	First Confirmation of Integron-Bearing Methicillin-Resistant Staphylococcus aureus. Current Microbiology, 2008, 57, 264-268.	1.0	95
12	First report of class 2 integron in clinical Enterococcus faecalis and class 1 integron in Enterococcus faecium in South China. Diagnostic Microbiology and Infectious Disease, 2010, 68, 315-317.	0.8	95
13	Detection of Foodborne Pathogens by Surface Enhanced Raman Spectroscopy. Frontiers in Microbiology, 2018, 9, 1236.	1.5	94
14	Formation and development of <i>Staphylococcus</i> biofilm: With focus on food safety. Journal of Food Safety, 2017, 37, e12358.	1.1	82
15	Development and application of a loop-mediated isothermal amplification method on rapid detection of Pseudomonas aeruginosa strains. World Journal of Microbiology and Biotechnology, 2011, 27, 181-184.	1.7	76
16	Development and application of a rapid and simple loop-mediated isothermal amplification method for food-borne Salmonella detection. Food Science and Biotechnology, 2010, 19, 1655-1659.	1.2	75
17	Transcriptomic analysis on the formation of the viable putative non-culturable state of beer-spoilage Lactobacillus acetotolerans. Scientific Reports, 2016, 6, 36753.	1.6	74
18	Longitudinal surveillance on antibiogram of important Gram-positive pathogens in Southern China, 2001 to 2015. Microbial Pathogenesis, 2017, 103, 80-86.	1.3	73

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19	Reduction and restoration of culturability of beer-stressed and low-temperature-stressed Lactobacillus acetotolerans strain 2011-8. International Journal of Food Microbiology, 2015, 206, 96-101.	2.1	71
20	Rapid detection of Vibrio parahaemolyticus strains and virulent factors by loop-mediated isothermal amplification assays. Food Science and Biotechnology, 2010, 19, 1191-1197.	1.2	66
21	Antimicrobial Resistance Investigation on <i>Staphylococcus</i> Strains in a Local Hospital in Guangzhou, China, 2001–2010. Microbial Drug Resistance, 2015, 21, 102-104.	0.9	65
22	Current methodologies on genotyping for nosocomial pathogen methicillin-resistant Staphylococcus aureus (MRSA). Microbial Pathogenesis, 2017, 107, 17-28.	1.3	64
23	Polymicrobial interaction and biofilms between Staphylococcus aureus and Pseudomonas aeruginosa: an underestimated concern in food safety. Current Opinion in Food Science, 2019, 26, 57-64.	4.1	60
24	Biofilm Formation of Staphylococcus aureus under Food Heat Processing Conditions: First Report on CML Production within Biofilm. Scientific Reports, 2019, 9, 1312.	1.6	57
25	Emerging resistance mechanisms for 4 types of common anti-MRSA antibiotics in Staphylococcus aureus: A comprehensive review. Microbial Pathogenesis, 2021, 156, 104915.	1.3	56
26	Effect of polymyxin resistance (pmr) on biofilm formation of Cronobacter sakazakii. Microbial Pathogenesis, 2017, 106, 16-19.	1.3	55
27	Complete sequence of pBM413, a novel multidrug resistance megaplasmid carrying qnrVC6 and bla IMP-45 from pseudomonas aeruginosa. International Journal of Antimicrobial Agents, 2018, 51, 145-150.	1.1	55
28	First study on the formation and resuscitation of viable but nonculturable state and beer spoilage capability of Lactobacillus lindneri. Microbial Pathogenesis, 2017, 107, 219-224.	1.3	54
29	Chromogenic media for MRSA diagnostics. Molecular Biology Reports, 2016, 43, 1205-1212.	1.0	53
30	Clinical features and antimicrobial resistance profiles of important Enterobacteriaceae pathogens in Guangzhou representative of Southern China, 2001–2015. Microbial Pathogenesis, 2017, 107, 206-211.	1.3	52
31	Transcriptomics Study on Staphylococcus aureus Biofilm Under Low Concentration of Ampicillin. Frontiers in Microbiology, 2018, 9, 2413.	1.5	51
32	Solvent-free enzymatic synthesis of 1, 3-Diacylglycerols by direct esterification of glycerol with saturated fatty acids. Lipids in Health and Disease, 2013, 12, 65.	1.2	50
33	Co-surfactant free microemulsions: Preparation, characterization and stability evaluation for food application. Food Chemistry, 2016, 204, 194-200.	4.2	48
34	Study on spoilage capability and VBNC state formation and recovery of Lactobacillus plantarum. Microbial Pathogenesis, 2017, 110, 257-261.	1.3	48
35	First report of novel genetic array aacA4 - bla IMP-25 - oxa30 - catB3 and identification of novel metallo-β-lactamase gene bla IMP25 : A Retrospective Study of antibiotic resistance surveillance on Psuedomonas aeruginosa in Guangzhou of South China, 2003–2007. Microbial Pathogenesis, 2016, 95, 62-67.	1.3	46
36	Analysis on pathogenic and virulent characteristics of the Cronobacter sakazakii strain BAA-894 by whole genome sequencing and its demonstration in basic biology science. Microbial Pathogenesis, 2017, 109, 280-286.	1.3	46

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37	Draft genome sequence and annotation ofLactobacillus acetotoleransBM-LA14527, a beer-spoilage bacteria. FEMS Microbiology Letters, 2016, 363, fnw201.	0.7	45
38	Molecular epidemiology and evolution of Haemophilus influenzae. Infection, Genetics and Evolution, 2020, 80, 104205.	1.0	43
39	Discovery and control of culturable and viable but non-culturable cells of a distinctive Lactobacillus harbinensis strain from spoiled beer. Scientific Reports, 2018, 8, 11446.	1.6	41
40	A 16-year retrospective surveillance report on the pathogenic features and antimicrobial susceptibility of Pseudomonas aeruginosa isolates from FAHJU in Guangzhou representative of Southern China. Microbial Pathogenesis, 2017, 110, 37-41.	1.3	40
41	Pathogenic features and characteristics of food borne pathogens biofilm: Biomass, viability and matrix. Microbial Pathogenesis, 2017, 111, 285-291.	1.3	38
42	The viable but nonculturable state induction and genomic analyses of <i>Lactobacillus casei</i> BMâ€LC14617, a beerâ€spoilage bacterium. MicrobiologyOpen, 2017, 6, e00506.	1.2	37
43	Induction and Recovery of the Viable but Nonculturable State of Hop-Resistance Lactobacillus brevis. Frontiers in Microbiology, 2018, 9, 2076.	1.5	37
44	Study the Features of 57 Confirmed CRISPR Loci in 38 Strains of Staphylococcus aureus. Frontiers in Microbiology, 2018, 9, 1591.	1,5	36
45	A variant ECE1 allele contributes to reduced pathogenicity of Candida albicans during vulvovaginal candidiasis. PLoS Pathogens, 2021, 17, e1009884.	2.1	35
46	Virulent and pathogenic features on the Cronobacter sakazakii polymyxin resistant pmr mutant strain s-3. Microbial Pathogenesis, 2017, 110, 359-364.	1.3	31
47	Effect of ultrasound treatment conditions on Saccharomyces cerevisiae by response surface methodology. Microbial Pathogenesis, 2017, 111, 497-502.	1.3	30
48	Whole-genome resequencing of Bacillus cereus and expression of genes functioning in sodium chloride stress. Microbial Pathogenesis, 2017, 104, 248-253.	1.3	29
49	Evaluation and application of molecular genotyping on nosocomial pathogen-methicillin-resistant Staphylococcus aureus isolates in Guangzhou representative of Southern China. Microbial Pathogenesis, 2017, 107, 397-403.	1.3	28
50	Inhibitory effects of two types of food additives on biofilm formation by foodborne pathogens. MicrobiologyOpen, 2019, 8, e00853.	1.2	25
51	Polymicrobial interaction between <i>Lactobacillus</i> and <i>Saccharomyces cerevisiae</i> : coexistence-relevant mechanisms. Critical Reviews in Microbiology, 2021, 47, 386-396.	2.7	24
52	Complete genome sequence and bioinformatics analyses of Bacillus thuringiensis strain BM-BT15426. Microbial Pathogenesis, 2017, 108, 55-60.	1.3	23
53	Formation and elimination of pyrraline in the Maillard reaction in a saccharide–lysine model system. Journal of the Science of Food and Agriculture, 2016, 96, 2555-2564.	1.7	22
54	Expression and purification of gp41-gp36 fusion protein and application in serological screening assay of HIV-1 and HIV-2. African Journal of Microbiology Research, 2012, 6, .	0.4	21

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55	Effect of aminoglycosides on the pathogenic characteristics of microbiology. Microbial Pathogenesis, 2017, 113, 357-364.	1.3	21
56	Investigation of Formation of Bacterial Biofilm upon Dead Siblings. Langmuir, 2019, 35, 7405-7413.	1.6	19
57	Spoilage Lactic Acid Bacteria in the Brewing Industry. Journal of Microbiology and Biotechnology, 2020, 30, 955-961.	0.9	18
58	Physical relation and mechanism of ultrasonic bactericidal activity on pathogenic E. coli with WPI. Microbial Pathogenesis, 2018, 117, 73-79.	1.3	17
59	Correlation and in vitro mechanism of bactericidal activity on E. coli with whey protein isolate during ultrasonic treatment. Microbial Pathogenesis, 2018, 115, 154-158.	1.3	16
60	Complete Sequence of pCY-CTX, a Plasmid Carrying a Phage-Like Region and an ISEcp1-Mediated Tn2Element fromEnterobacter cloacae. Microbial Drug Resistance, 2018, 24, 307-313.	0.9	16
61	Microbial virulence, molecular epidemiology and pathogenic factors of fluoroquinolone-resistant Haemophilus influenzae infections in Guangzhou, China. Annals of Clinical Microbiology and Antimicrobials, 2018, 17, 41.	1.7	16
62	Rapid Detection of Food-Borne Escherichia coli O157:H7 with Visual Inspection by Crossing Priming Amplification (CPA). Food Analytical Methods, 2020, 13, 474-481.	1.3	16
63	Phenotypic characterization of pathogenic Cronobacter spp. strains. Microbial Pathogenesis, 2018, 121, 232-237.	1.3	15
64	Formation of Peptide Bound Pyrraline in the Maillard Model Systems with Different Lys-Containing Dipeptides and Tripeptides. Molecules, 2016, 21, 463.	1.7	14
65	The fingerprint mapping and genotyping systems application on methicillin-resistant Staphylococcus aureus. Microbial Pathogenesis, 2018, 125, 246-251.	1.3	14
66	Identification of the KPC plasmid pCT-KPC334: New insights on the evolutionary pathway of epidemic plasmids harboring fosA3-blaKPC-2 genes. International Journal of Antimicrobial Agents, 2018, 52, 510-511.	1.1	12
67	Intense pulsed light for inactivation of foodborne gram-positive bacteria in planktonic cultures and bacterial biofilms. LWT - Food Science and Technology, 2021, 152, 112374.	2.5	12
68	Determination of Free-Form and Peptide Bound Pyrraline in the Commercial Drinks Enriched with Different Protein Hydrolysates. International Journal of Molecular Sciences, 2016, 17, 1053.	1.8	10
69	Lithium Hydroxide Hydrolysis Combined with MALDI TOF Mass Spectrometry for Rapid Sphingolipid Detection. Journal of the American Society for Mass Spectrometry, 2021, 32, 289-300.	1.2	10
70	Emergence of Clinical <i>Pseudomonas aeruginosa</i> Isolate Guangzhou-PaeC79 Carrying <i>crpP</i> , <i>bla</i> _{GES-5} , and <i>bla</i> _{KPC-2} in Guangzhou of China. Microbial Drug Resistance, 2021, 27, 965-970.	0.9	10
71	Impact of pmrA on Cronobacter sakazakii planktonic and biofilm cells: A comprehensive transcriptomic study. Food Microbiology, 2021, 98, 103785.	2.1	10
72	Detection of culturable and viable but non-culturable cells of beer spoilage lactic acid bacteria by combined use of propidium monoazide and <i>horA</i> specific polymerase chain reaction. Journal of the Institute of Brewing, 2016, 122, 29-33.	0.8	9

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73	Complete genomic analysis of multidrug-resistance Pseudomonas aeruginosa Guangzhou-Pae617, the host of megaplasmid pBM413. Microbial Pathogenesis, 2018, 117, 265-269.	1.3	9
74	Complete Sequence of a Novel Multidrug-Resistant Pseudomonas putida Strain Carrying Two Copies of qnrVC6. Microbial Drug Resistance, 2019, 25, 1-7.	0.9	9
75	Antibiotic Susceptibility, Biofilm-Forming Ability, and Incidence of Class 1 Integron of <i>Salmonella</i> spp., <i>Escherichia coli</i> , and <i>Staphylococcus aureus</i> Isolated from Various Foods in a School Canteen in China. Foodborne Pathogens and Disease, 2020, 17, 269-275.	0.8	9
76	Regulatory network controls microbial biofilm development, with <i>Candida albicans</i> as a representative: from adhesion to dispersal. Bioengineered, 2022, 13, 253-267.	1.4	9
77	A novel procedure in combination of genomic sequencing, flow cytometry and routine culturing for confirmation of beer spoilage caused by Pediococcus damnosus in viable but nonculturable state. LWT - Food Science and Technology, 2022, 154, 112623.	2.5	8
78	Development of a propidium monoazide-polymerase chain reaction assay for detection of viable Lactobacillus brevis in beer. Brazilian Journal of Microbiology, 2017, 48, 740-746.	0.8	7
79	Effects of magnetic fields on the enzymatic synthesis of naringin palmitate. RSC Advances, 2018, 8, 13364-13369.	1.7	7
80	Effect of ultrasonic field on the enzyme activities and ion balance of potential pathogen Saccharomyces cerevisiae. Microbial Pathogenesis, 2018, 119, 216-220.	1.3	7
81	Comparative genomic analyses of two novel qnrVC6 carrying multidrug-resistant Pseudomonas. spp strains. Microbial Pathogenesis, 2018, 123, 269-274.	1.3	7
82	Genomic analysis of a hop-resistance Lactobacillus brevis strain responsible for food spoilage and capable of entering into the VBNC state. Microbial Pathogenesis, 2020, 145, 104186.	1.3	7
83	Adaptive behaviors of planktonic Pseudomonas aeruginosa in response to the surface-deposited dead siblings. Colloids and Surfaces B: Biointerfaces, 2021, 197, 111408.	2.5	7
84	Understanding of food biofilms by the application of omics techniques. Future Microbiology, 2021, 16, 257-269.	1.0	7
85	Molecular Epidemiology, Microbial Virulence, and Resistance of Carbapenem-Resistant <i>Enterobacterales</i> Isolates in a Teaching Hospital in Guangzhou, China. Microbial Drug Resistance, 2022, 28, 698-709.	0.9	7
86	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
87	Sheaolein-based cold-soluble powder fats with medium- and long-chain triacylglycerol: production via chemical interesterification using sheaolein and palm kernel stearin. RSC Advances, 2016, 6, 18632-18640.	1.7	5
88	Microbial infection pattern, pathogenic features and resistance mechanism of carbapenem-resistant Gram negative bacilli during long-term hospitalization. Microbial Pathogenesis, 2018, 117, 356-360.	1.3	5
89	Chromosome and Plasmid Features of Two ST37 <i>Clostridioides difficile</i> Strains Isolated in China Reveal Distinct Multidrug Resistance and Virulence Determinants. Microbial Drug Resistance, 2020, 26, 1503-1508.	0.9	5
90	Development and evaluation of reverse-transcription loop-mediated isothermal amplification for rapid detection of human immunodeficiency virus type 1. Indian Journal of Medical Microbiology, 2012, 30, 391-396.	0.3	4

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91	Occurrence of Serratia marcescens Carrying blaIMP-26 and mcr-9 in Southern China: New Insights in the Evolution of Megaplasmid IMP-26. Antibiotics, 2022, 11, 869.	1.5	4
92	Lowâ€Temperature Chemical Glycerolysis: An Evaluation of Substrates Miscibility on Reaction Rate. JAOCS, Journal of the American Oil Chemists' Society, 2011, 88, 1077-1079.	0.8	3
93	Staphylococcal Food Poisoning and Novel Perspectives in Food Safety. , 2016, , .		3
94	Antioxidant Profile of 1â€Monocaffeoyl Glycerol in Lipophobic/Lipophilic Media. Journal of Food Science, 2019, 84, 2091-2100.	1.5	3
95	Significant downtrend of antimicrobial resistance rate and rare β-lactamase genes and plasmid replicons carriage in clinical Pseudomonas aeruginosa in Southern China. Microbial Pathogenesis, 2021, 159, 105124.	1.3	3
96	Resistome and virulome study on pathogenic Streptococcus agalactiae Guangzhou-SAG036. Microbial Pathogenesis, 2020, 147, 104258.	1.3	1
97	Editorial: The Prevalence of MDR Non-Fermenting Gram Negative Bacteria and Their Chemotherapy. Frontiers in Microbiology, 2021, 12, 664336.	1.5	1
98	Letter to the Editor: Four Novel Types of Gene Cassettes from Carbapenem-Resistant Pseudomonas aeruginosa in Southern China—First Report of qnrVC7. Microbial Drug Resistance, 2021, 27, 1011-1012.	0.9	1
99	Food pathogens. , 2021, , 295-321.		1
100	Studies on Salt Tolerance of Bacillus Isolated from the Industrial Soy Sauce Residue. , 2016, , .		1
101	Verification and application of a modified carbapenem inactivation method (mCIM) on <i>Pseudomonas aeruginosa</i> : a potential screening methodology on carbapenemases phenotype in <i>Bacillus cereus</i> . Bioengineered, 2022, 13, 12088-12098.	1.4	1
102	Antimicrobial susceptibility and genetic features of a heterogeneous vancomycin intermediate-resistant Staphylococcus aureus strain. Infection, Genetics and Evolution, 2020, 85, 104565.	1.0	0
103	Editorial. Bioprocess and Biosystems Engineering, 2021, 44, 927-928.	1.7	0
104	Editorial: Emerging Frontiers in the Formation of Viable but Non-culturable Microorganisms and Biofilms During Food Processing. Frontiers in Microbiology, 2021, 12, 726348.	1.5	0
105	A strategy design based on antibiotic‑resistance and plasmid replicons genes of clinical Escherichia coli strains. Bioengineered, 2022, 13, 7499-7513.	1.4	0