

Minggui Wang

List of Publications by Year in descending order

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113
papers

4,975
citations

117625

34
h-index

106344

65
g-index

115
all docs

115
docs citations

115
times ranked

5238
citing authors

#	ARTICLE	IF	CITATIONS
1	Plasmid-Mediated Quinolone Resistance in Clinical Isolates of <i>Escherichia coli</i> from Shanghai, China. <i>Antimicrobial Agents and Chemotherapy</i> , 2003, 47, 2242-2248.	3.2	423
2	New Plasmid-Mediated Quinolone Resistance Gene, <i>qnrC</i> , Found in a Clinical Isolate of <i>Proteus mirabilis</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2009, 53, 1892-1897.	3.2	279
3	Achieving global targets for antimicrobial resistance. <i>Science</i> , 2016, 353, 874-875.	12.6	233
4	Emerging Plasmid-Mediated Quinolone Resistance Associated with the <i>qnr</i> Gene in <i>Klebsiella pneumoniae</i> Clinical Isolates in the United States. <i>Antimicrobial Agents and Chemotherapy</i> , 2004, 48, 1295-1299.	3.2	219
5	Current Status and Trends of Antibacterial Resistance in China. <i>Clinical Infectious Diseases</i> , 2018, 67, S128-S134.	5.8	205
6	<i>qnr</i> Gene Nomenclature. <i>Antimicrobial Agents and Chemotherapy</i> , 2008, 52, 2297-2299.	3.2	192
7	High Prevalence of Plasmid-Mediated Quinolone Resistance Determinants <i>qnr</i> , <i>aac(6)</i> Tj ETQq1 1 0.784314 rgBT /Ove from Companion and Food-Producing Animals. <i>Antimicrobial Agents and Chemotherapy</i> , 2009, 53, 519-524.	3.2	188
8	Molecular and clinical epidemiology of carbapenem-resistant Enterobacterales in the USA (CRACKLE-2): a prospective cohort study. <i>Lancet Infectious Diseases</i> , The, 2020, 20, 731-741.	9.1	174
9	<i>vanM</i> , a New Glycopeptide Resistance Gene Cluster Found in <i>Enterococcus faecium</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2010, 54, 4643-4647.	3.2	169
10	Emergence of a Plasmid-Encoded Resistance-Nodulation-Division Efflux Pump Conferring Resistance to Multiple Drugs, Including Tigecycline, in <i>Klebsiella pneumoniae</i> . <i>MBio</i> , 2020, 11, .	4.1	153
11	Clinical outcomes and bacterial characteristics of carbapenem-resistant <i>Klebsiella pneumoniae</i> complex among patients from different global regions (CRACKLE-2): a prospective, multicentre, cohort study. <i>Lancet Infectious Diseases</i> , The, 2022, 22, 401-412.	9.1	122
12	Antimicrobial Susceptibility of <i>Mycoplasma pneumoniae</i> Isolates and Molecular Analysis of Macrolide-Resistant Strains from Shanghai, China. <i>Antimicrobial Agents and Chemotherapy</i> , 2009, 53, 2160-2162.	3.2	118
13	Evidence-based Guideline for Therapeutic Drug Monitoring of Vancomycin: 2020 Update by the Division of Therapeutic Drug Monitoring, Chinese Pharmacological Society. <i>Clinical Infectious Diseases</i> , 2020, 71, S363-S371.	5.8	109
14	Prevalence of the <i>oqxAB</i> gene complex in <i>Klebsiella pneumoniae</i> and <i>Escherichia coli</i> clinical isolates. <i>Journal of Antimicrobial Chemotherapy</i> , 2012, 67, 1655-1659.	3.0	86
15	Laboratory diagnosis, clinical management and infection control of the infections caused by extensively drug-resistant Gram-negative bacilli: a Chinese consensus statement. <i>Clinical Microbiology and Infection</i> , 2016, 22, S15-S25.	6.0	85
16	<i>Clostridium difficile</i> infections in a Shanghai hospital: antimicrobial resistance, toxin profiles and ribotypes. <i>International Journal of Antimicrobial Agents</i> , 2009, 33, 339-342.	2.5	84
17	Distribution of 16S rRNA methylases among different species of Gram-negative bacilli with high-level resistance to aminoglycosides. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2010, 29, 1349-1353.	2.9	83
18	The Colonization of Carbapenem-Resistant <i>Klebsiella pneumoniae</i> : Epidemiology, Resistance Mechanisms, and Risk Factors in Patients Admitted to Intensive Care Units in China. <i>Journal of Infectious Diseases</i> , 2020, 221, S206-S214.	4.0	83

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19	Clinical and Genomic Analysis of Liver Abscess-Causing <i>Klebsiella pneumoniae</i> Identifies New Liver Abscess-Associated Virulence Genes. <i>Frontiers in Cellular and Infection Microbiology</i> , 2016, 6, 165.	3.9	82
20	Characterization of macrolide resistance in <i>Mycoplasma pneumoniae</i> isolated from children in Shanghai, China. <i>Diagnostic Microbiology and Infectious Disease</i> , 2010, 67, 355-358.	1.8	74
21	First Report of the Multidrug Resistance Gene <i>cfr</i> and the Phenicol Resistance Gene <i>fexA</i> in a <i>Bacillus</i> Strain from Swine Feces. <i>Antimicrobial Agents and Chemotherapy</i> , 2010, 54, 3953-3955.	3.2	71
22	Mechanisms of Tigecycline Resistance among <i>Klebsiella pneumoniae</i> Clinical Isolates. <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 6982-6985.	3.2	71
23	Comparison of Adhesin Genes and Antimicrobial Susceptibilities between Uropathogenic and Intestinal Commensal <i>Escherichia coli</i> Strains. <i>PLoS ONE</i> , 2013, 8, e61169.	2.5	70
24	High ceftazidime hydrolysis activity and porin <i>OmpK35</i> deficiency contribute to the decreased susceptibility to ceftazidime/avibactam in KPC-producing <i>Klebsiella pneumoniae</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2017, 72, 1930-1936.	3.0	64
25	Mapping the resistance-associated mobilome of a carbapenem-resistant <i>Klebsiella pneumoniae</i> strain reveals insights into factors shaping these regions and facilitates generation of a "resistance-disarmed" model organism. <i>Journal of Antimicrobial Chemotherapy</i> , 2015, 70, 2770-2774.	3.0	55
26	Susceptibility of Extended-Spectrum- β -Lactamase-Producing Enterobacteriaceae According to the New CLSI Breakpoints. <i>Journal of Clinical Microbiology</i> , 2011, 49, 3127-3131.	3.9	54
27	Sulbactam-based therapy for <i>Acinetobacter baumannii</i> infection: a systematic review and meta-analysis. <i>Brazilian Journal of Infectious Diseases</i> , 2013, 17, 389-394.	0.6	53
28	Glutathione-S-transferase <i>FosA6</i> of <i>Klebsiella pneumoniae</i> origin conferring fosfomycin resistance in ESBL-producing <i>Escherichia coli</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2016, 71, 2460-2465.	3.0	49
29	Characterization of <i>fosA5</i> , a new plasmid-mediated fosfomycin resistance gene in <i>Escherichia coli</i> . <i>Letters in Applied Microbiology</i> , 2015, 60, 259-264.	2.2	47
30	Development of a Fast Raman-Assisted Antibiotic Susceptibility Test (FRAST) for the Antibiotic Resistance Analysis of Clinical Urine and Blood Samples. <i>Analytical Chemistry</i> , 2021, 93, 5098-5106.	6.5	45
31	Characterization of Fosfomycin Resistance Gene, <i>fosB</i> , in Methicillin-Resistant <i>Staphylococcus aureus</i> Isolates. <i>PLoS ONE</i> , 2016, 11, e0154829.	2.5	42
32	Prevalence of Fosfomycin Resistance and Mutations in <i>murA</i> , <i>glpT</i> , and <i>uhpT</i> in Methicillin-Resistant <i>Staphylococcus aureus</i> Strains Isolated from Blood and Cerebrospinal Fluid Samples. <i>Frontiers in Microbiology</i> , 2015, 6, 1544.	3.5	39
33	Activities of Newer Quinolones against <i>Escherichia coli</i> and <i>Klebsiella pneumoniae</i> Containing the Plasmid-Mediated Quinolone Resistance Determinant <i>qnr</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2004, 48, 1400-1401.	3.2	36
34	Coexistence of <i>qnrB4</i> and <i>qnrS1</i> in a clinical strain of <i>Klebsiella pneumoniae</i> . <i>Acta Pharmacologica Sinica</i> , 2008, 29, 320-324.	6.1	36
35	Shifts in the Antibiotic Susceptibility, Serogroups, and Clonal Complexes of <i>Neisseria meningitidis</i> in Shanghai, China: A Time Trend Analysis of the Pre-Quinolone and Quinolone Eras. <i>PLoS Medicine</i> , 2015, 12, e1001838.	8.4	35
36	Mutations of the Transporter Proteins <i>GlpT</i> and <i>UhpT</i> Confer Fosfomycin Resistance in <i>Staphylococcus aureus</i> . <i>Frontiers in Microbiology</i> , 2017, 8, 914.	3.5	35

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37	High Prevalence of <i>vanM</i> in Vancomycin-Resistant <i>Enterococcus faecium</i> Isolates from Shanghai, China. <i>Antimicrobial Agents and Chemotherapy</i> , 2015, 59, 7795-7798.	3.2	34
38	Prevalence and Expression of the Plasmid-Mediated Quinolone Resistance Determinant <i>qnrA1</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2007, 51, 4105-4110.	3.2	32
39	Molecular and Clinical Characteristics of <i>Clostridium difficile</i> Infection in a University Hospital in Shanghai, China. <i>Clinical Infectious Diseases</i> , 2008, 47, 1606-1608.	5.8	32
40	The Fosfomycin Resistance Gene <i>fosB3</i> Is Located on a Transferable, Extrachromosomal Circular Intermediate in Clinical <i>Enterococcus faecium</i> Isolates. <i>PLoS ONE</i> , 2013, 8, e78106.	2.5	31
41	New Delhi Metallo- β -Lactamase-1 in Carbapenem-Resistant <i>Salmonella</i> Strain, China. <i>Emerging Infectious Diseases</i> , 2013, 19, 2049-2051.	4.3	30
42	Liquid chromatography/tandem mass spectrometry assay for the simultaneous determination of cefoperazone and sulbactam in plasma and its application to a pharmacokinetic study. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2010, 878, 3119-3124.	2.3	27
43	Porin Deficiency in Carbapenem-Resistant <i>Enterobacter aerogenes</i> Strains. <i>Microbial Drug Resistance</i> , 2018, 24, 1277-1283.	2.0	27
44	Efflux pumps AcrAB and OqxAB contribute to nitrofurantoin resistance in an uropathogenic <i>Klebsiella pneumoniae</i> isolate. <i>International Journal of Antimicrobial Agents</i> , 2019, 54, 223-227.	2.5	27
45	Decreased quinolone susceptibility in high percentage of <i>Enterobacter cloacae</i> clinical isolates caused only by Qnr determinants. <i>Diagnostic Microbiology and Infectious Disease</i> , 2010, 67, 110-113.	1.8	26
46	Multiclonal Origin of Macrolide-Resistant <i>Mycoplasma pneumoniae</i> Isolates as Determined by Multilocus Variable-Number Tandem-Repeat Analysis. <i>Journal of Clinical Microbiology</i> , 2012, 50, 2793-2795.	3.9	26
47	Prevalence of the fosfomycin-resistance determinant, <i>fosB3</i> , in <i>Enterococcus faecium</i> clinical isolates from China. <i>Journal of Medical Microbiology</i> , 2014, 63, 1484-1489.	1.8	26
48	Dissemination of blaOXA-23-harboring carbapenem-resistant <i>Acinetobacter baumannii</i> clones in Pakistan. <i>Journal of Global Antimicrobial Resistance</i> , 2020, 21, 357-362.	2.2	25
49	High-Level Fosfomycin Resistance in Vancomycin-Resistant <i>Enterococcus faecium</i> . <i>Emerging Infectious Diseases</i> , 2017, 23, 1902-1904.	4.3	23
50	Clonal dissemination of extensively drug-resistant <i>Acinetobacter baumannii</i> producing an OXA-23 β -lactamase at a teaching hospital in Shanghai, China. <i>Journal of Microbiology, Immunology and Infection</i> , 2015, 48, 101-108.	3.1	22
51	An IncP-2 plasmid sublineage associated with dissemination of <i>bla</i> _{IMP-45} among carbapenem-resistant <i>Pseudomonas aeruginosa</i> . <i>Emerging Microbes and Infections</i> , 2021, 10, 442-449.	6.5	22
52	Prevalence of plasmid-mediated quinolone-resistance determinants in <i>Shigella flexneri</i> isolates from Anhui Province, China. <i>Journal of Antibiotics</i> , 2010, 63, 187-189.	2.0	21
53	High-level tetracycline resistance mediated by efflux pumps Tet(A) and Tet(A)-1 with two start codons. <i>Journal of Medical Microbiology</i> , 2014, 63, 1454-1459.	1.8	21
54	Detection of <i>Mycoplasma pneumoniae</i> P1 subtype variations by denaturing gradient gel electrophoresis. <i>Diagnostic Microbiology and Infectious Disease</i> , 2014, 78, 24-28.	1.8	21

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55	CTX-M-190, a Novel β -Lactamase Resistant to Tazobactam and Sulbactam, Identified in an <i>Escherichia coli</i> Clinical Isolate. <i>Antimicrobial Agents and Chemotherapy</i> , 2017, 61, .	3.2	21
56	Macrolide-resistant <i>Mycoplasma pneumoniae</i> prevalence and clinical aspects in adult patients with community-acquired pneumonia in China: a prospective multicenter surveillance study. <i>Journal of Thoracic Disease</i> , 2017, 9, 3774-3781.	1.4	21
57	The prevalence of plasmid-mediated quinolone resistance determinants among clinical isolates of ESBL or AmpC-producing <i>Escherichia coli</i> from Chinese pediatric patients. <i>Microbiology and Immunology</i> , 2010, 54, 123-128.	1.4	21
58	In vivo development of tigecycline resistance in <i>Klebsiella pneumoniae</i> owing to deletion of the ramR ribosomal binding site. <i>International Journal of Antimicrobial Agents</i> , 2017, 50, 523-528.	2.5	20
59	Selection and characterisation of <i>Staphylococcus aureus</i> mutants with reduced susceptibility to the investigational oxazolidinone MRX-I. <i>International Journal of Antimicrobial Agents</i> , 2014, 43, 418-422.	2.5	19
60	CHINET efforts to control antimicrobial resistance in China. <i>Journal of Global Antimicrobial Resistance</i> , 2020, 21, 76-77.	2.2	19
61	RamA upregulates multidrug resistance efflux pumps AcrAB and OqxAB in <i>Klebsiella pneumoniae</i> . <i>International Journal of Antimicrobial Agents</i> , 2021, 57, 106251.	2.5	19
62	Comparison of empirical therapy with cefoperazone/sulbactam or a carbapenem for bloodstream infections due to ESBL-producing Enterobacteriaceae. <i>Journal of Antimicrobial Chemotherapy</i> , 2018, 73, 3176-3180.	3.0	18
63	Molecular Characteristics and Antimicrobial Susceptibility Profiles of <i>Elizabethkingia</i> Clinical Isolates in Shanghai, China. <i>Infection and Drug Resistance</i> , 2020, Volume 13, 247-256.	2.7	17
64	Co-production of SFO-1 and DHA-1 β -lactamases and 16S rRNA methylase ArmA in clinical isolates of <i>Klebsiella pneumoniae</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2012, 67, 2361-2366.	3.0	16
65	Sequence Types Harboring Genes Encoding Aminoglycoside Modifying Enzymes and 16SrRNA Methylase; a Multicenter Study from Pakistan. <i>Infection and Drug Resistance</i> , 2020, Volume 13, 2855-2862.	2.7	15
66	A mutational analysis and molecular dynamics simulation of quinolone resistance proteins QnrA1 and QnrC from <i>Proteus mirabilis</i> . <i>BMC Structural Biology</i> , 2010, 10, 33.	2.3	14
67	Establishment of norvancomycin fluorescence polarization immunoassay for therapeutic drug monitoring. <i>Journal of Antibiotics</i> , 2012, 65, 35-39.	2.0	14
68	Association Between Single-Nucleotide Polymorphism in CISH Gene and Susceptibility to Tuberculosis in Chinese Han Population. <i>Cell Biochemistry and Biophysics</i> , 2014, 68, 529-534.	1.8	14
69	IncX2 and IncX1-X2 Hybrid Plasmids Coexisting in a FosA6-Producing <i>Escherichia coli</i> Strain. <i>Antimicrobial Agents and Chemotherapy</i> , 2017, 61, .	3.2	14
70	Characterization of a bla _{NDM-1} -harboring plasmid from a <i>Salmonella enterica</i> clinical isolate in China. <i>Molecular Medicine Reports</i> , 2017, 16, 1087-1092.	2.4	13
71	Structures of Class I and Class II Transcription Complexes Reveal the Molecular Basis of RamA-Dependent Transcription Activation. <i>Advanced Science</i> , 2022, 9, e2103669.	11.2	13
72	Type II and type IV topoisomerase mutations in clinical isolates of <i>Morganella morganii</i> harbouring the qnrD gene. <i>Annals of Clinical Microbiology and Antimicrobials</i> , 2014, 13, 34.	3.8	12

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73	Characterization of a Novel IncHI2 Plasmid Carrying Tandem Copies of <i>bla</i> _{CTX-M-2} in a <i>fosA6</i> -Harboring <i>Escherichia coli</i> Sequence Type 410 Strain. <i>Antimicrobial Agents and Chemotherapy</i> , 2016, 60, 6742-6747.	3.2	12
74	Evaluation of anidulafungin in the treatment of intra-abdominal candidiasis: a pooled analysis of patient-level data from 5 prospective studies. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2019, 38, 1849-1856.	2.9	12
75	Clinical and molecular characteristics of <i>Chryseobacterium indologenes</i> isolates at a teaching hospital in Shanghai, China. <i>Annals of Translational Medicine</i> , 2021, 9, 668-668.	1.7	12
76	Four Carbapenem-Resistant Gram-Negative Species Carrying Distinct Carbapenemases in a Single Patient. <i>Journal of Clinical Microbiology</i> , 2015, 53, 1031-1033.	3.9	11
77	In vivo Acquisition of Carbapenemase Gene <i>bla</i> KPC-2 in Multiple Species of Enterobacteriaceae through Horizontal Transfer of Insertion Sequence or Plasmid. <i>Frontiers in Microbiology</i> , 2016, 7, 1651.	3.5	11
78	Evolution of Sequence Type 4821 Clonal Complex Meningococcal Strains in China from Prequinolone to Quinolone Era, 1972–2013. <i>Emerging Infectious Diseases</i> , 2018, 24, 683-690.	4.3	11
79	Outbreak Of <i>Klebsiella pneumoniae</i> Carbapenemase-Producing <i>Klebsiella aerogenes</i> Strains In A Tertiary Hospital In China. <i>Infection and Drug Resistance</i> , 2019, Volume 12, 3283-3290.	2.7	11
80	Prevalence of <i>fusB</i> in <i>Staphylococcus aureus</i> clinical isolates. <i>Journal of Medical Microbiology</i> , 2013, 62, 1199-1203.	1.8	10
81	Efflux Pump <i>AcrAB</i> Confers Decreased Susceptibility to Piperacillin–Tazobactam and Ceftolozane–Tazobactam in Tigecycline-Non-Susceptible <i>Klebsiella pneumoniae</i> . <i>Infection and Drug Resistance</i> , 2020, Volume 13, 4309-4319.	2.7	10
82	Plasmids and genes contributing to high-level quinolone resistance in <i>Escherichia coli</i> . <i>International Journal of Antimicrobial Agents</i> , 2020, 56, 105987.	2.5	10
83	Characterization of the novel plasmid-encoded MBL gene <i>bla</i> AFM-1, integrated into a <i>bla</i> IMP-45-bearing transposon Tn6485e in a carbapenem-resistant <i>Pseudomonas aeruginosa</i> clinical isolate. <i>Journal of Antimicrobial Chemotherapy</i> , 2021, 77, 83-88.	3.0	10
84	Prevalence and phenotypes of erythromycin-resistant <i>Streptococcus pneumoniae</i> in Shanghai, China. <i>Diagnostic Microbiology and Infectious Disease</i> , 2001, 39, 187-189.	1.8	9
85	Antimicrobial Resistance in China: Challenges and Actions. <i>Clinical Infectious Diseases</i> , 2018, 67, S127-S127.	5.8	9
86	Diagnosis and Management of Intraabdominal Infection: Guidelines by the Chinese Society of Surgical Infection and Intensive Care and the Chinese College of Gastrointestinal Fistula Surgeons. <i>Clinical Infectious Diseases</i> , 2020, 71, S337-S362.	5.8	9
87	Vancomycin Heteroresistance in <i>vanM</i> -type <i>Enterococcus faecium</i> . <i>Microbial Drug Resistance</i> , 2020, 26, 776-782.	2.0	8
88	Determination of norvancomycin epidemiological cut-off values (ECOFFs) for <i>Staphylococcus aureus</i> , <i>Staphylococcus epidermidis</i> , <i>Staphylococcus haemolyticus</i> and <i>Staphylococcus hominis</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2021, 76, 152-159.	3.0	8
89	Transmission barrier of the <i>bla</i> KPC plasmid mediated by type I restriction-modification systems in <i>Escherichia coli</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2022, 77, 952-956.	3.0	7
90	A Clinical Practice Guideline for the Emergency Management of Anaphylaxis (2020). <i>Frontiers in Pharmacology</i> , 2022, 13, 845689.	3.5	7

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91	Identification of <i>qnrE3</i> and <i>qnrE4</i> , New Transferable Quinolone Resistance <i>qnrE</i> Family Genes Originating from <i>Enterobacter mori</i> and <i>Enterobacter asburiae</i> , Respectively. <i>Antimicrobial Agents and Chemotherapy</i> , 2021, 65, e0045621.	3.2	6
92	Treatment of extensively drug-resistant Gram-negative infections in critically ill patients: Outcome of a consensus meeting at the 13th Asia-Pacific Congress of Clinical Microbiology and Infection, October 2012. <i>Journal of Global Antimicrobial Resistance</i> , 2013, 1, 117-122.	2.2	5
93	Emergence of tigecycline- and carbapenem-nonsusceptible <i>Klebsiella pneumoniae</i> ST11 clone in patients without exposure to tigecycline. <i>Journal of Microbiology, Immunology and Infection</i> , 2016, 49, 962-968.	3.1	5
94	Treatment and economic burden of mucormycosis in China: Case report review and burden estimation. <i>Journal of Clinical Pharmacy and Therapeutics</i> , 2022, 47, 905-914.	1.5	5
95	A Case-Control Study: Clinical Characteristics of Nosocomial Bloodstream Infections Versus Non-bloodstream Infections of <i>Acinetobacter</i> spp.. <i>Clinical Infectious Diseases</i> , 2018, 67, S189-S195.	5.8	4
96	In Vivo Evolution of CTX-M-215, a Novel Narrow-Spectrum β -Lactamase in an <i>Escherichia coli</i> Clinical Isolate Conferring Resistance to Mecillinam. <i>Antimicrobial Agents and Chemotherapy</i> , 2020, 64, .	3.2	4
97	Establishment of epidemiological cut-off values for cefoselis, a new fourth-generation cephalosporin, against <i>Escherichia coli</i> , <i>Klebsiella pneumoniae</i> , <i>Enterobacter cloacae</i> , <i>Proteus mirabilis</i> and <i>Pseudomonas aeruginosa</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2021, 76, 2593-2599.	3.0	4
98	Combined PK/PD Index May Be a More Appropriate PK/PD Index for Cefoperazone/Sulbactam against <i>Acinetobacter baumannii</i> in Patients with Hospital-Acquired Pneumonia. <i>Antibiotics</i> , 2022, 11, 703.	3.7	4
99	Carbapenemase-Encoding Gene Copy Number Estimator (CCNE): a Tool for Carbapenemase Gene Copy Number Estimation. <i>Microbiology Spectrum</i> , 2022, 10, .	3.0	4
100	Genetic diversity of fluoroquinolone-nonsusceptible <i>Streptococcus pneumoniae</i> clinical isolates and the first identification of serotype 20B in China. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2014, 33, 465-470.	2.9	3
101	Molecular characteristics of <i>Clostridium difficile</i> strains from patients with a first recurrence more than 8 weeks after the primary infection. <i>Journal of Microbiology, Immunology and Infection</i> , 2017, 50, 532-536.	3.1	3
102	New Subclass B1 Metallo- β -Lactamase Gene from a Clinical Pathogenic <i>Myroides odoratus</i> Strain. <i>Microbial Drug Resistance</i> , 2018, 24, 909-914.	2.0	3
103	China's United States Research Collaborations in Antimicrobial Resistance. <i>Clinical Infectious Diseases</i> , 2018, 67, S142-S145.	5.8	3
104	The Predominance of Strain Replacement Among Enterobacteriaceae Pairs With Emerging Carbapenem Resistance During Hospitalization. <i>Journal of Infectious Diseases</i> , 2020, 221, S215-S219.	4.0	3
105	Antibiotic Exposure during the Preceding Six Months Is Related to Intestinal ESBL-Producing Enterobacteriaceae Carriage in the Elderly. <i>Antibiotics</i> , 2022, 11, 953.	3.7	2
106	Comparison of the genetic structures surrounding <i>qnrA1</i> in Korean <i>Enterobacter cloacae</i> and Chinese <i>Escherichia coli</i> Strains isolated in the early 2000s: Evidence for <i>qnrA</i> mobilization via Inc HI2 type plasmid. <i>Journal of Microbiology</i> , 2012, 50, 166-169.	2.8	1
107	Molecular Characteristics of <i>Escherichia coli</i> Causing Bloodstream Infections During 2010-2015 in Tertiary Hospital, Shanghai, China. <i>Infection and Drug Resistance</i> , 2021, Volume 14, 2079-2086.	2.7	1
108	Coexistence of <i>armA</i> and genes encoding aminoglycoside-modifying enzymes in <i>Acinetobacter baumannii</i> . <i>African Journal of Microbiology Research</i> , 2012, 6, .	0.4	1

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109	Serogroup Y Clonal Complex 23 Meningococcus in China Acquiring Penicillin Resistance from Commensal <i>Neisseria lactamica</i> Species. <i>Antimicrobial Agents and Chemotherapy</i> , 2022, 66, .	3.2	1
110	The prevalence of plasmid-mediated quinolone resistance determinants among clinical isolates of ESBL or AmpC-producing <i>Escherichia coli</i> from Chinese pediatric patients. <i>Microbiology and Immunology</i> , 2009, 54, 123-128.	1.4	0
111	Draft Genome Sequence of Linezolid-Resistant <i>Enterococcus faecalis</i> Clinical Isolate HS0914. <i>Genome Announcements</i> , 2014, 2, .	0.8	0
112	2037. A Novel Strategy of Antimicrobial Stewardship in Shanghai: Preliminary Practice with Integration of Three Surveillance Networks. <i>Open Forum Infectious Diseases</i> , 2019, 6, S685-S685.	0.9	0
113	toxin B-induced colonic inflammation is mediated by the FOXO3/PPM1B pathway in fetal human colon epithelial cells. <i>American Journal of Translational Research (discontinued)</i> , 2020, 12, 6204-6219.	0.0	0