Yonghong Xiao

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/8392322/publications.pdf

Version: 2024-02-01

122 papers	3,479 citations	159585 30 h-index	182427 51 g-index
132 all docs	132 docs citations	132 times ranked	4865 citing authors

#	Article	IF	Citations
1	Taking the right measures to control COVID-19. Lancet Infectious Diseases, The, 2020, 20, 523-524.	9.1	251
2	Use and Prescription of Antibiotics in Primary Health Care Settings in China. JAMA Internal Medicine, 2014, 174, 1914.	5.1	210
3	Nationwide high prevalence of CTX-M and an increase of CTX-M-55 in Escherichia coli isolated from patients with community-onset infections in Chinese county hospitals. BMC Infectious Diseases, 2014, 14, 659.	2.9	139
4	Editorial: Horizontal Gene Transfer Mediated Bacterial Antibiotic Resistance. Frontiers in Microbiology, 2019, 10, 1933.	3.5	136
5	Antimicrobials: a global alliance for optimizing their rational use in intra-abdominal infections (AGORA). World Journal of Emergency Surgery, 2016, 11, 33.	5.0	130
6	Changes in Chinese Policies to Promote the Rational Use of Antibiotics. PLoS Medicine, 2013, 10, e1001556.	8.4	126
7	Influence of H7N9 virus infection and associated treatment on human gut microbiota. Scientific Reports, 2015, 5, 14771.	3.3	88
8	Legislation of clinical antibiotic use in China. Lancet Infectious Diseases, The, 2013, 13, 189-191.	9.1	86
9	Plasmon enhanced photocatalytic and antimicrobial activities of Ag-TiO2 nanocomposites under visible light irradiation prepared by DBD cold plasma treatment. Materials Science and Engineering C, 2019, 96, 197-204.	7.3	75
10	Association between antibiotic consumption and the rate of carbapenem-resistant Gram-negative bacteria from China based on 153 tertiary hospitals data in 2014. Antimicrobial Resistance and Infection Control, 2018, 7, 137.	4.1	73
11	Molecular Epidemiology and Colistin Resistant Mechanism of mcr-Positive and mcr-Negative Clinical Isolated Escherichia coli. Frontiers in Microbiology, 2017, 8, 2262.	3.5	65
12	China's national plan to combat antimicrobial resistance. Lancet Infectious Diseases, The, 2016, 16, 1216-1218.	9.1	58
13	Occurrence and Genomic Characterization of ESBL-Producing, MCR-1-Harboring Escherichia coli in Farming Soil. Frontiers in Microbiology, 2017, 8, 2510.	3.5	56
14	High burden of antimicrobial drug resistance in Asia. Journal of Global Antimicrobial Resistance, 2014, 2, 141-147.	2.2	55
15	Molecular Epidemiology and Genetic Diversity of Fluoroquinolone-Resistant Escherichia coli Isolates from Patients with Community-Onset Infections in 30 Chinese County Hospitals. Journal of Clinical Microbiology, 2015, 53, 766-770.	3.9	54
16	Antimicrobial Stewardship in China: Systems, Actions and Future Strategies. Clinical Infectious Diseases, 2018, 67, S135-S141.	5.8	53
17	Characterization of lasR-deficient clinical isolates of Pseudomonas aeruginosa. Scientific Reports, 2018, 8, 13344.	3.3	52
18	Emergence of Raoultella ornithinolytica Coproducing IMP-4 and KPC-2 Carbapenemases in China. Antimicrobial Agents and Chemotherapy, 2015, 59, 7086-7089.	3.2	50

#	Article	IF	Citations
19	High Prevalence of ESBL-Producing Klebsiella pneumoniae Causing Community-Onset Infections in China. Frontiers in Microbiology, 2016, 7, 1830.	3.5	50
20	The potential impact of the COVID-19 pandemic on global antimicrobial and biocide resistance: an AMR Insights global perspective. JAC-Antimicrobial Resistance, 2021, 3, dlab038.	2.1	48
21	The Global Alliance for Infections in Surgery: defining a model for antimicrobial stewardshipâ€"results from an international cross-sectional survey. World Journal of Emergency Surgery, 2017, 12, 34.	5.0	47
22	Identification of novel tetracycline resistance gene tet (X14) and its co-occurrence with tet (X2) in a tigecycline-resistant and colistin-resistant Empedobacter stercoris. Emerging Microbes and Infections, 2020, 9, 1843-1852.	6.5	42
23	Stool Samples of Acute Diarrhea Inpatients as a Reservoir of ST11 Hypervirulent KPC-2-Producing Klebsiella pneumoniae. MSystems, 2020, 5, .	3.8	42
24	Dysbiosis of urinary microbiota is positively correlated with Type 2 diabetes mellitus. Oncotarget, 2017, 8, 3798-3810.	1.8	41
25	Identification and genomic characterization of a KPC-2-, NDM-1- and NDM-5-producing Klebsiella michiganensis isolate. Journal of Antimicrobial Chemotherapy, 2018, 73, 536-538.	3.0	40
26	Discovery and characterisation of an escherichia coli ST206 strain producing NDM-5 and MCR-1 from a patient with acute diarrhoea in China. International Journal of Antimicrobial Agents, 2018, 51, 273-275.	2.5	38
27	Wza gene knockout decreases Acinetobacter baumannii virulence and affects Wzy-dependent capsular polysaccharide synthesis. Virulence, 2020, 11, 1-13.	4.4	36
28	Complete genome sequencing and genomic characterization of two Escherichia coli strains co-producing MCR-1 and NDM-1 from bloodstream infection. Scientific Reports, 2017, 7, 17885.	3.3	35
29	Association between the rate of fluoroquinolones-resistant gram-negative bacteria and antibiotic consumption from China based on 145 tertiary hospitals data in 2014. BMC Infectious Diseases, 2020, 20, 269.	2.9	35
30	A retrospective analysis of Pseudomonas aeruginosa bloodstream infections: prevalence, risk factors, and outcome in carbapenem-susceptible and -non-susceptible infections. Antimicrobial Resistance and Infection Control, 2019, 8, 68.	4.1	34
31	Combined delivery of angiopoietin-1 gene and simvastatin mediated by anti-intercellular adhesion molecule-1 antibody-conjugated ternary nanoparticles for acute lung injury therapy. Nanomedicine: Nanotechnology, Biology, and Medicine, 2019, 15, 25-36.	3.3	34
32	MDR Salmonella enterica serovar Typhimurium ST34 carrying mcr-1 isolated from cases of bloodstream and intestinal infection in children in China. Journal of Antimicrobial Chemotherapy, 2020, 75, 92-95.	3.0	33
33	Bacterial-resistance among outpatients of county hospitals in China: significant geographic distinctions and minor differences between central cities. Microbes and Infection, 2015, 17, 417-425.	1.9	32
34	A Retrospective Analysis of Risk Factors and Outcomes of Carbapenem-Resistant Klebsiella pneumoniae Bacteremia in Nontransplant Patients. Journal of Infectious Diseases, 2020, 221, S174-S183.	4.0	32
35	Analysis of tigecycline resistance development in clinical Acinetobacter baumannii isolates through a combined genomic and transcriptomic approach. Scientific Reports, 2016, 6, 26930.	3.3	31
36	In vitro antibacterial activity of fosfomycin combined with other antimicrobials against KPC-producing Klebsiella pneumoniae. International Journal of Antimicrobial Agents, 2017, 50, 237-241.	2.5	31

#	ARTICLE	IF	CITATIONS
37	Alterations of Urinary Microbiota in Type 2 Diabetes Mellitus with Hypertension and/or Hyperlipidemia. Frontiers in Physiology, 2017, 8, 126.	2.8	31
38	Characterization of the urinary microbiota of elderly women and the effects of type 2 diabetes and urinary tract infections on the microbiota. Oncotarget, 2017, 8, 100678-100690.	1.8	31
39	Characterization of the population structure, drug resistance mechanisms and plasmids of the community-associated Enterobacter cloacae complex in China. Journal of Antimicrobial Chemotherapy, 2018, 73, 66-76.	3.0	30
40	Comparison of Tigecycline or Cefoperazone/Sulbactam therapy for bloodstream infection due to Carbapenem-resistant Acinetobacter baumannii. Antimicrobial Resistance and Infection Control, 2019, 8, 52.	4.1	29
41	The Major Aminoglycoside-Modifying Enzyme AAC(3)-II Found inEscherichia coliDetermines a Significant Disparity in Its Resistance to Gentamicin and Amikacin in China. Microbial Drug Resistance, 2012, 18, 42-46.	2.0	27
42	Genomic Epidemiology and Characterization of Methicillin-Resistant Staphylococcus aureus from Bloodstream Infections in China. MSystems, 2021, 6, e0083721.	3.8	27
43	Study protocol for One Health data collections, analyses and intervention of the Sino-Swedish integrated multisectoral partnership for antibiotic resistance containment (IMPACT). BMJ Open, 2018, 8, e017832.	1.9	26
44	Retrospective comparative analysis of risk factors and outcomes in patients with carbapenem resistant Acinetobacter baumannii bloodstream infections: cefoperazone–sulbactam associated with resistance and tigecycline increased the mortality. Infection and Drug Resistance, 2018, Volume 11, 2021-2030.	2.7	26
45	A retrospective analysis of risk factors and outcomes in patients with extended-spectrum beta-lactamase-producing Escherichia coli bloodstream infections. Journal of Global Antimicrobial Resistance, 2019, 17, 147-156.	2.2	26
46	Silent transmission of an IS 1294b -deactivated mcr-1 gene with inducible colistin resistance. International Journal of Antimicrobial Agents, 2018, 51, 822-828.	2.5	25
47	Retrospective survey of the efficacy of mandatory implementation of the Essential Medicine Policy in the primary healthcare setting in China: failure to promote the rational use of antibiotics in clinics. International Journal of Antimicrobial Agents, 2016, 48, 409-414.	2.5	24
48	Hypervirulence Markers Among Non-ST11 Strains of Carbapenem- and Multidrug-Resistant Klebsiella pneumoniae Isolated From Patients With Bloodstream Infections. Frontiers in Microbiology, 2020, 11, 1199.	3.5	24
49	A retrospective, comparative analysis of risk factors and outcomes in carbapenem-susceptible and carbapenem-nonsusceptible Klebsiella pneumoniae bloodstream infections: tigecycline significantly increases the mortality. Infection and Drug Resistance, 2018, Volume 11, 595-606.	2.7	23
50	Change in Antibiotic Use in Secondary and Tertiary Hospitals Nationwide After a National Antimicrobial Stewardship Campaign Was Launched in China, 2011â€“2016: An Observational Study. Journal of Infectious Diseases, 2020, 221, S148-S155.	4.0	23
51	Clinical features and treatment of patients with Vibrio vulnificus infection. International Journal of Infectious Diseases, 2017, 59, 1-6.	3.3	21
52	A novel Tn1696-like composite transposon (Tn6404) harboring bla IMP-4 in a Klebsiella pneumoniae isolate carrying a rare ESBL gene bla SFO-1. Scientific Reports, 2017, 7, 17321.	3.3	20
53	Blood bacterial resistant investigation collaborative system (BRICS) report: a national surveillance in China from 2014 to 2019. Antimicrobial Resistance and Infection Control, 2022, 11, 17.	4.1	20
54	Low prevalence of MCR-1-producing Klebsiella pneumoniae in bloodstream infections in China. Clinical Microbiology and Infection, 2018, 24, 205-206.	6.0	19

#	Article	IF	Citations
55	Alteration of the Gut Microbiome in Chronic Kidney Disease Patients and Its Association With Serum Free Immunoglobulin Light Chains. Frontiers in Immunology, 2021, 12, 609700.	4.8	19
56	Rifaximin Modulates the Gut Microbiota to Prevent Hepatic Encephalopathy in Liver Cirrhosis Without Impacting the Resistome. Frontiers in Cellular and Infection Microbiology, 2021, 11, 761192.	3.9	19
57	Building bridges to operationalise one health â€“ A Sino-Swedish collaboration to tackle antibiotic resistance. One Health, 2016, 2, 139-143.	3.4	18
58	Characterization of highly virulent community-associated methicillin-resistant Staphylococcus aureus ST9-SCC mec XII causing bloodstream infection in China. Emerging Microbes and Infections, 2020, 9, 2526-2535.	6.5	17
59	Genomic Analysis Of A KPC-2-Producing Klebsiella Pneumoniae ST11 Outbreak From A Teaching Hospital In Shandong Province, China . Infection and Drug Resistance, 2019, Volume 12, 2961-2969.	2.7	16
60	Epidemiology and risk factors of infective endocarditis in a tertiary hospital in China from 2007 to 2016. BMC Infectious Diseases, 2020, 20, 428.	2.9	16
61	In vitro Pharmacokinetics/Pharmacodynamics Evaluation of Fosfomycin Combined with Amikacin or Colistin against KPC2-Producing Klebsiella pneumoniae. Frontiers in Cellular and Infection Microbiology, 2017, 7, 246.	3.9	15
62	Complete nucleotide sequences of two KPC-2-encoding plasmids from the same Citrobacter freundii isolate. Journal of Antimicrobial Chemotherapy, 2018, 73, 531-533.	3.0	15
63	Risk factors and outcomes in non-transplant patients with extended-spectrum beta-lactamase-producing Escherichia coli bacteremia: a retrospective study from 2013 to 2016. Antimicrobial Resistance and Infection Control, 2019, 8, 144.	4.1	15
64	Comparison of Genetic Features and Evolution of Global and Chinese Strains of Community-Associated Methicillin-Resistant Staphylococcus aureus ST22. Microbiology Spectrum, 2022, 10, e0203721.	3.0	15
65	Genome sequencing and genomic characterization of a tigecycline-resistant Klebsiella pneumoniae strain isolated from the bile samples of a cholangiocarcinoma patient. Gut Pathogens, 2014, 6, 40.	3.4	14
66	Genome characterization of two bile-isolated Vibrio fluvialis strains: an insight into pathogenicity and bile salt adaption. Scientific Reports, 2017, 7, 11827.	3.3	14
67	The genetic feature and virulence determinant of highly virulent community-associated MRSA ST338-SCCmec Vb in China. Emerging Microbes and Infections, 2021, 10, 1052-1064.	6.5	14
68	Severe infective endocarditis with systemic embolism due to community associated methicillin-resistant Staphylococcus aureus ST630. Brazilian Journal of Infectious Diseases, 2015, 19, 85-89.	0.6	13
69	Occurrence and Genomic Characterization of Two MCR-1-Producing Escherichia coli Isolates from the Same Mink Farmer. MSphere, 2019, 4, .	2.9	13
70	Dissemination of a â€˜rareâ€™ extended-spectrum Î²-lactamase gene blaSFO-1 mediated by epidemic clones of carbapenemase-producing Enterobacter hormaechei in China. International Journal of Antimicrobial Agents, 2020, 56, 106079.	2.5	13
71	Detection of a new tet(X6)-encoding plasmid in Acinetobacter towneri. Journal of Global Antimicrobial Resistance, 2021, 25, 132-136.	2.2	13
72	Identification ofRaoultella terrigenaas a Rare Causative Agent of Subungual Abscess Based on 16S rRNA and Housekeeping Gene Sequencing. Canadian Journal of Infectious Diseases and Medical Microbiology, 2016, 2016, 1-4.	1.9	12

#	Article	IF	Citations
73	High prevalence of a globally disseminated hypervirulent clone, Staphylococcus aureus CC121, with reduced vancomycin susceptibility in community settings in China. Journal of Antimicrobial Chemotherapy, 2019, 74, 2537-2543.	3.0	12
74	In Vitro Activity Comparison of Ceftazidimeâ€“Avibactam and Aztreonamâ€“Avibactam Against Bloodstream Infections With Carbapenem-Resistant Organisms in China. Frontiers in Cellular and Infection Microbiology, 2021, 11, 780365.	3.9	12
75	Bloodstream infections caused by Entero-bacteriaceae in China. Lancet Infectious Diseases, The, 2019, 19, 810-811.	9.1	11
76	A two-step preparation method for nanocrystalline Ag-decorated cotton fabrics and their antibacterial assessment. Journal of Materials Science, 2019, 54, 10447-10456.	3.7	11
77	It is time to define an organizational model for the prevention and management of infections along the surgical pathway: a worldwide cross-sectional survey. World Journal of Emergency Surgery, 2022, 17, 17.	5.0	11
78	Socioeconomic Burden of Bloodstream Infections Caused by Carbapenem-Resistant Enterobacteriaceae. Infection and Drug Resistance, 2021, Volume 14, 5385-5393.	2.7	11
79	Comparative Genomic Analysis of 19 Clinical Isolates of Tigecycline-Resistant Acinetobacter baumannii. Frontiers in Microbiology, 2020, 11, 1321.	3.5	10
80	Comparative Genomic Analysis Provides Insights into the Evolution and Genetic Diversity of Community-Genotype Sequence Type 72 Staphylococcus aureus Isolates. MSystems, 2021, 6, e0098621.	3.8	10
81	Utility and Applicability of Rapid Diagnostic Testing in Antimicrobial Stewardship in the Asia-Pacific Region: A Delphi Consensus. Clinical Infectious Diseases, 2022, 74, 2067-2076.	5.8	10
82	Emergence of a novel Enterobacter kobei clone carrying chromosomal-encoded CTX-M-12 with diversified pathogenicity in northeast China. New Microbes and New Infections, 2017, 17, 7-10.	1.6	9
83	Antibacterial effect evaluation of moxalactam against extended-spectrum β-lactamase-producing Escherichia coli and Klebsiella pneumoniae with in vitro pharmacokinetics/pharmacodynamics simulation. Infection and Drug Resistance, 2018, Volume 11, 103-112.	2.7	9
84	The clinical features and prognosis of infective endocarditis in the elderly from 2007 to 2016 in a tertiary hospital in China. BMC Infectious Diseases, 2019, 19, 937.	2.9	9
85	Association between the rate of third generation cephalosporin-resistant Escherichia coli and Klebsiella pneumoniae and antibiotic consumption based on 143 Chinese tertiary hospitals data in 2014. European Journal of Clinical Microbiology and Infectious Diseases, 2020, 39, 1495-1502.	2.9	9
86	Predicting hosts based on early SARS-CoV-2 samples and analyzing the 2020 pandemic. Scientific Reports, 2021, 11, 17422.	3.3	9
87	Comparative genomic and transmission analysis of Clostridioides difficile between environmental, animal, and clinical sources in China. Emerging Microbes and Infections, 2021, 10, 2244-2255.	6.5	9
88	In vitro reduction of colistin susceptibility and comparative genomics reveals multiple differences between MCR-positive and MCR-negative colistin-resistant Escherichia coli . Infection and Drug Resistance, 2019, Volume 12, 1665-1674.	2.7	8
89	Economic Burden of Patients with Bloodstream Infections Caused by Extended-Spectrum Î²-Lactamase-Producing Escherichia coli . Infection and Drug Resistance, 2020, Volume 13, 3583-3592.	2.7	8
90	Rapid diagnostic testing for antimicrobial stewardship: Utility in Asia Pacific. Infection Control and Hospital Epidemiology, 2021, 42, 864-868.	1.8	8

#	Article	IF	Citations
91	Genomic epidemiology and characterisation of penicillin-sensitive Staphylococcus aureus isolates from invasive bloodstream infections in China: an increasing prevalence and higher diversity in genetic typing be revealed. Emerging Microbes and Infections, 2022, 11, 326-336.	6.5	8
92	Genome sequence of Shigella flexneri strain SP1, a diarrheal isolate that encodes an extended-spectrum Î²-lactamase (ESBL). Annals of Clinical Microbiology and Antimicrobials, 2017, 16, 37.	3.8	7
93	In vitro antibacterial effect of fosfomycin combination therapy against colistin-resistant Klebsiella pneumoniae. Infection and Drug Resistance, 2018, Volume 11, 577-585.	2.7	7
94	Emergence of KPC-2-Producing Raoultella ornithinolytica Isolated from a Hospital Wastewater Treatment Plant. Antimicrobial Agents and Chemotherapy, 2020, 64, .	3.2	7
95	New options for bloodstream infections caused by colistinâ€•or ceftazidime/avibactamâ€•resistant Klebsiella pneumoniae. International Journal of Antimicrobial Agents, 2021, 58, 106458.	2.5	7
96	Community-associated meticillin-resistant Staphylococcus aureus pneumonia in China. Lancet Infectious Diseases, The, 2017, 17, 26.	9.1	6
97	Serotype Is Associated With High Rate of Colistin Resistance Among Clinical Isolates of Salmonella. Frontiers in Microbiology, 2020, 11, 592146.	3.5	6
98	Comparative Analysis of Virulence and Toxin Expression of Vancomycin-Intermediate and Vancomycin-Sensitive Staphylococcus aureus Strains. Frontiers in Microbiology, 2020, 11, 596942.	3.5	6
99	Optimal Empiric Polymyxin B Treatment of Patients Infected with Gram-Negative Organisms Detected Using a Blood Antimicrobial Surveillance Network in China. Drug Design, Development and Therapy, 2021, Volume 15, 2593-2603.	4.3	6
100	Identification and characterization of cfr-positive Staphylococcus aureus isolates from community-onset infectious patients in a county hospital in China. Journal of Medical Microbiology, 2015, 64, 910-915.	1.8	6
101	Intrinsic colistin resistance. Lancet Infectious Diseases, The, 2016, 16, 1227-1228.	9.1	5
102	Simulating moxalactam dosage for extended-spectrum Î²-lactamase-producing Enterobacteriaceae using blood antimicrobial surveillance network data . Infection and Drug Resistance, 2019, Volume 12, 1199-1208.	2.7	5
103	Complete-Genome Sequencing and Comparative Genomic Characterization of an IMP-4 Producing Citrobacter freundii Isolate from Patient with Diarrhea . Infection and Drug Resistance, 2020, Volume 13, 1057-1065.	2.7	5
104	Socioeconomic burden of bloodstream infections caused by carbapenem-resistant and carbapenem-susceptible Pseudomonas aeruginosa in China. Journal of Global Antimicrobial Resistance, 2021, 26, 101-107.	2.2	5
105	Encephalomyelitis Caused by Balamuthia mandrillaris in a Woman With Breast Cancer: A Case Report and Review of the Literature. Frontiers in Immunology, 2021, 12, 768065.	4.8	5
106	Rapid increase in occurrence of carbapenem-resistant Enterobacteriaceae in healthy rural residents in Shandong Province, China, from 2015 to 2017. Journal of Global Antimicrobial Resistance, 2022, 28, 38-42.	2.2	5
107	Determining optimal dosing regimen of oral administration of dicloxacillin using Monte Carlo simulation. Drug Design, Development and Therapy, 2017, Volume 11, 1951-1956.	4.3	4
108	Detection of an In104-like integron carrying a blaIMP-34 gene in Enterobacter cloacae isolates co-producing IMP-34 and VIM-1. Journal of Antimicrobial Chemotherapy, 2019, 74, 2812-2814.	3.0	4

#	ARTICLE	IF	CITATIONS
109	Optimal empiric treatment for KPC-2-producing Klebsiella pneumoniae infections in critically ill patients with normal or decreased renal function using Monte Carlo simulation. BMC Infectious Diseases, 2021, 21, 307.	2.9	4
110	Performance of different methods for testing polymyxin B: comparison of broth microdilution, agar dilution and MIC test strip in mcrâ€1 positive and negative Escherichia coli. Letters in Applied Microbiology, 2021, 73, 197-205.	2.2	4
111	Molecular Characterization of Carbapenem-Resistant Acinetobacter baumannii Isolates Among Intensive Care Unit Patients and Environment. Infection and Drug Resistance, 2022, Volume 15, 1821-1829.	2.7	4
112	Complete genome sequence of Lactobacillus heilongjiangensis DSM 28069T: Insight into its probiotic potential. Journal of Biotechnology, 2015, 216, 65-66.	3.8	3
113	Using Monte Carlo simulation to determine optimal dosing regimen for cefetamet sodium for injection. Journal of Chemotherapy, 2016, 28, 172-179.	1.5	3
114	Evaluation of Ceftazidime/Avibactam Administration in Enterobacteriaceae and Pseudomonas aeruginosa Bloodstream Infections by Monte Carlo Simulation. Drug Design, Development and Therapy, 2021, Volume 15, 2899-2905.	4.3	3
115	Clinical Characteristics of Patients and Whole Genome Sequencing-Based Surveillance of Escherichia coli Community-Onset Bloodstream Infections at a Non-tertiary Hospital in CHINA. Frontiers in Microbiology, 2021, 12, 748471.	3.5	3
116	Comprehensive Genome Analysis of Carbapenem-Resistant Strains of Raoultella Species, an Emerging Multidrug-Resistant Bacterium in Hospitals. Antimicrobial Agents and Chemotherapy, 2019, 63, .	3.2	2
117	The Monte Carlo Simulation of Three Antimicrobials for Empiric Treatment of Adult Bloodstream Infections With Carbapenem-Resistant Enterobacterales in China. Frontiers in Microbiology, 2021, 12, 738812.	3.5	2
118	Evolution of Drug-resistant Acinetobacter baumannii After DCD Renal Transplantation. Scientific Reports, 2017, 7, 1968.	3.3	1
119	Complete nucleotide sequence of pSKLX3330, an IncI1 plasmid carrying bla CTX-M-55 isolated from community-onset Escherichia coli infection. Journal of Global Antimicrobial Resistance, 2017, 11, 120-122.	2.2	1
120	Effect of Short-Term Antimicrobial Therapy on the Tolerance and Antibiotic Resistance of Multidrug-Resistant Staphylococcus capitis . Infection and Drug Resistance, 2020, Volume 13, 2017-2026.	2.7	1
121	Acquisition of the mcr-1 Gene Lowers the Target Mutation to Impede the Evolution of a High-Level Colistin-Resistant Mutant in Escherichia coli. Infection and Drug Resistance, 2021, Volume 14, 3041-3051.	2.7	1
122	Antibacterial Activity and Optimal Treatment of Ceftazidime-Avibactam and Aztreonam-Avibactam Against Bloodstream Infections Caused by Carbapenem-Resistant Klebsiella pneumoniae. Frontiers in Pharmacology, 2021, 12, 771910.	3.5	1