

# Yonghong Xiao

## List of Publications by Year in descending order

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122  
papers

3,479  
citations

159358

30  
h-index

182168

51  
g-index

132  
all docs

132  
docs citations

132  
times ranked

4865  
citing authors

#	ARTICLE	IF	CITATIONS
1	Utility and Applicability of Rapid Diagnostic Testing in Antimicrobial Stewardship in the Asia-Pacific Region: A Delphi Consensus. <i>Clinical Infectious Diseases</i> , 2022, 74, 2067-2076.	2.9	10
2	Genomic epidemiology and characterisation of penicillin-sensitive <i>Staphylococcus aureus</i> isolates from invasive bloodstream infections in China: an increasing prevalence and higher diversity in genetic typing be revealed. <i>Emerging Microbes and Infections</i> , 2022, 11, 326-336.	3.0	8
3	Blood bacterial resistant investigation collaborative system (BRICS) report: a national surveillance in China from 2014 to 2019. <i>Antimicrobial Resistance and Infection Control</i> , 2022, 11, 17.	1.5	20
4	Rapid increase in occurrence of carbapenem-resistant Enterobacteriaceae in healthy rural residents in Shandong Province, China, from 2015 to 2017. <i>Journal of Global Antimicrobial Resistance</i> , 2022, 28, 38-42.	0.9	5
5	Comparison of Genetic Features and Evolution of Global and Chinese Strains of Community-Associated Methicillin-Resistant <i>Staphylococcus aureus</i> ST22. <i>Microbiology Spectrum</i> , 2022, 10, e0203721.	1.2	15
6	It is time to define an organizational model for the prevention and management of infections along the surgical pathway: a worldwide cross-sectional survey. <i>World Journal of Emergency Surgery</i> , 2022, 17, 17.	2.1	11
7	Molecular Characterization of Carbapenem-Resistant <i>Acinetobacter baumannii</i> Isolates Among Intensive Care Unit Patients and Environment. <i>Infection and Drug Resistance</i> , 2022, Volume 15, 1821-1829.	1.1	4
8	The genetic feature and virulence determinant of highly virulent community-associated MRSA ST338-SCCmec Vb in China. <i>Emerging Microbes and Infections</i> , 2021, 10, 1052-1064.	3.0	14
9	Optimal empiric treatment for KPC-2-producing <i>Klebsiella pneumoniae</i> infections in critically ill patients with normal or decreased renal function using Monte Carlo simulation. <i>BMC Infectious Diseases</i> , 2021, 21, 307.	1.3	4
10	The potential impact of the COVID-19 pandemic on global antimicrobial and biocide resistance: an AMR Insights global perspective. <i>JAC-Antimicrobial Resistance</i> , 2021, 3, dlab038.	0.9	48
11	Alteration of the Gut Microbiome in Chronic Kidney Disease Patients and Its Association With Serum Free Immunoglobulin Light Chains. <i>Frontiers in Immunology</i> , 2021, 12, 609700.	2.2	19
12	Optimal Empiric Polymyxin B Treatment of Patients Infected with Gram-Negative Organisms Detected Using a Blood Antimicrobial Surveillance Network in China. <i>Drug Design, Development and Therapy</i> , 2021, Volume 15, 2593-2603.	2.0	6
13	Rapid diagnostic testing for antimicrobial stewardship: Utility in Asia Pacific. <i>Infection Control and Hospital Epidemiology</i> , 2021, 42, 864-868.	1.0	8
14	Performance of different methods for testing polymyxin B: comparison of broth microdilution, agar dilution and MIC test strip in <i>mcrA</i> positive and negative <i>Escherichia coli</i> . <i>Letters in Applied Microbiology</i> , 2021, 73, 197-205.	1.0	4
15	Detection of a new tet(X6)-encoding plasmid in <i>Acinetobacter towneri</i> . <i>Journal of Global Antimicrobial Resistance</i> , 2021, 25, 132-136.	0.9	13
16	Evaluation of Ceftazidime/Avibactam Administration in Enterobacteriaceae and <i>Pseudomonas aeruginosa</i> Bloodstream Infections by Monte Carlo Simulation. <i>Drug Design, Development and Therapy</i> , 2021, Volume 15, 2899-2905.	2.0	3
17	Acquisition of the mcr-1 Gene Lowers the Target Mutation to Impede the Evolution of a High-Level Colistin-Resistant Mutant in <i>Escherichia coli</i> . <i>Infection and Drug Resistance</i> , 2021, Volume 14, 3041-3051.	1.1	1
18	Predicting hosts based on early SARS-CoV-2 samples and analyzing the 2020 pandemic. <i>Scientific Reports</i> , 2021, 11, 17422.	1.6	9

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19	Socioeconomic burden of bloodstream infections caused by carbapenem-resistant and carbapenem-susceptible <i>Pseudomonas aeruginosa</i> in China. <i>Journal of Global Antimicrobial Resistance</i> , 2021, 26, 101-107.	0.9	5
20	Comparative Genomic Analysis Provides Insights into the Evolution and Genetic Diversity of Community-Genotype Sequence Type 72 <i>Staphylococcus aureus</i> Isolates. <i>MSystems</i> , 2021, 6, e0098621.	1.7	10
21	Clinical Characteristics of Patients and Whole Genome Sequencing-Based Surveillance of <i>Escherichia coli</i> Community-Onset Bloodstream Infections at a Non-tertiary Hospital in CHINA. <i>Frontiers in Microbiology</i> , 2021, 12, 748471.	1.5	3
22	New options for bloodstream infections caused by colistin- or ceftazidime/avibactam-resistant <i>Klebsiella pneumoniae</i> . <i>International Journal of Antimicrobial Agents</i> , 2021, 58, 106458.	1.1	7
23	Genomic Epidemiology and Characterization of Methicillin-Resistant <i>Staphylococcus aureus</i> from Bloodstream Infections in China. <i>MSystems</i> , 2021, 6, e0083721.	1.7	27
24	The Monte Carlo Simulation of Three Antimicrobials for Empiric Treatment of Adult Bloodstream Infections With Carbapenem-Resistant Enterobacterales in China. <i>Frontiers in Microbiology</i> , 2021, 12, 738812.	1.5	2
25	In Vitro Activity Comparison of Ceftazidime-Avibactam and Aztreonam-Avibactam Against Bloodstream Infections With Carbapenem-Resistant Organisms in China. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021, 11, 780365.	1.8	12
26	Comparative genomic and transmission analysis of <i>Clostridioides difficile</i> between environmental, animal, and clinical sources in China. <i>Emerging Microbes and Infections</i> , 2021, 10, 2244-2255.	3.0	9
27	Encephalomyelitis Caused by <i>Balamuthia mandrillaris</i> in a Woman With Breast Cancer: A Case Report and Review of the Literature. <i>Frontiers in Immunology</i> , 2021, 12, 768065.	2.2	5
28	Rifaximin Modulates the Gut Microbiota to Prevent Hepatic Encephalopathy in Liver Cirrhosis Without Impacting the Resistome. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021, 11, 761192.	1.8	19
29	Antibacterial Activity and Optimal Treatment of Ceftazidime-Avibactam and Aztreonam-Avibactam Against Bloodstream Infections Caused by Carbapenem-Resistant <i>Klebsiella pneumoniae</i> . <i>Frontiers in Pharmacology</i> , 2021, 12, 771910.	1.6	1
30	Socioeconomic Burden of Bloodstream Infections Caused by Carbapenem-Resistant Enterobacteriaceae. <i>Infection and Drug Resistance</i> , 2021, Volume 14, 5385-5393.	1.1	11
31	MDR <i>Salmonella enterica</i> serovar Typhimurium ST34 carrying <i>mcr-1</i> isolated from cases of bloodstream and intestinal infection in children in China. <i>Journal of Antimicrobial Chemotherapy</i> , 2020, 75, 92-95.	1.3	33
32	<i>Wza</i> gene knockout decreases <i>Acinetobacter baumannii</i> virulence and affects Wzy-dependent capsular polysaccharide synthesis. <i>Virulence</i> , 2020, 11, 1-13.	1.8	36
33	Emergence of KPC-2-Producing <i>Raoultella ornithinolytica</i> Isolated from a Hospital Wastewater Treatment Plant. <i>Antimicrobial Agents and Chemotherapy</i> , 2020, 64, .	1.4	7
34	Characterization of highly virulent community-associated methicillin-resistant <i>Staphylococcus aureus</i> ST9-SCC <i>mecXII</i> causing bloodstream infection in China. <i>Emerging Microbes and Infections</i> , 2020, 9, 2526-2535.	3.0	17
35	Identification of novel tetracycline resistance gene <i>tet(X14)</i> and its co-occurrence with <i>tet(X2)</i> in a tigecycline-resistant and colistin-resistant <i>Empedobacter stercoris</i> . <i>Emerging Microbes and Infections</i> , 2020, 9, 1843-1852.	3.0	42
36	Economic Burden of Patients with Bloodstream Infections Caused by Extended-Spectrum $\beta$ -Lactamase-Producing <i>Escherichia coli</i> . <i>Infection and Drug Resistance</i> , 2020, Volume 13, 3583-3592.	1.1	8

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37	Serotype Is Associated With High Rate of Colistin Resistance Among Clinical Isolates of Salmonella. <i>Frontiers in Microbiology</i> , 2020, 11, 592146.	1.5	6
38	Comparative Analysis of Virulence and Toxin Expression of Vancomycin-Intermediate and Vancomycin-Sensitive Staphylococcus aureus Strains. <i>Frontiers in Microbiology</i> , 2020, 11, 596942.	1.5	6
39	Hypervirulence Markers Among Non-ST11 Strains of Carbapenem- and Multidrug-Resistant Klebsiella pneumoniae Isolated From Patients With Bloodstream Infections. <i>Frontiers in Microbiology</i> , 2020, 11, 1199.	1.5	24
40	Epidemiology and risk factors of infective endocarditis in a tertiary hospital in China from 2007 to 2016. <i>BMC Infectious Diseases</i> , 2020, 20, 428.	1.3	16
41	Change in Antibiotic Use in Secondary and Tertiary Hospitals Nationwide After a National Antimicrobial Stewardship Campaign Was Launched in China, 2011–2016: An Observational Study. <i>Journal of Infectious Diseases</i> , 2020, 221, S148-S155.	1.9	23
42	A Retrospective Analysis of Risk Factors and Outcomes of Carbapenem-Resistant Klebsiella pneumoniae Bacteremia in Nontransplant Patients. <i>Journal of Infectious Diseases</i> , 2020, 221, S174-S183.	1.9	32
43	Taking the right measures to control COVID-19. <i>Lancet Infectious Diseases</i> , The, 2020, 20, 523-524.	4.6	251
44	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. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2020, 39, 1495-1502.	1.3	9
45	Dissemination of a “rare” extended-spectrum $\beta$ -lactamase gene blaSFO-1 mediated by epidemic clones of carbapenemase-producing Enterobacter hormaechei in China. <i>International Journal of Antimicrobial Agents</i> , 2020, 56, 106079.	1.1	13
46	Stool Samples of Acute Diarrhea Inpatients as a Reservoir of ST11 Hypervirulent KPC-2-Producing Klebsiella pneumoniae. <i>MSystems</i> , 2020, 5, .	1.7	42
47	Effect of Short-Term Antimicrobial Therapy on the Tolerance and Antibiotic Resistance of Multidrug-Resistant Staphylococcus capitis. <i>Infection and Drug Resistance</i> , 2020, Volume 13, 2017-2026.	1.1	1
48	Comparative Genomic Analysis of 19 Clinical Isolates of Tigecycline-Resistant Acinetobacter baumannii. <i>Frontiers in Microbiology</i> , 2020, 11, 1321.	1.5	10
49	Complete-Genome Sequencing and Comparative Genomic Characterization of an IMP-4 Producing Citrobacter freundii Isolate from Patient with Diarrhea. <i>Infection and Drug Resistance</i> , 2020, Volume 13, 1057-1065.	1.1	5
50	Association between the rate of fluoroquinolones-resistant gram-negative bacteria and antibiotic consumption from China based on 145 tertiary hospitals data in 2014. <i>BMC Infectious Diseases</i> , 2020, 20, 269.	1.3	35
51	Bloodstream infections caused by Entero-bacteriaceae in China. <i>Lancet Infectious Diseases</i> , The, 2019, 19, 810-811.	4.6	11
52	Comprehensive Genome Analysis of Carbapenem-Resistant Strains of Raoultella Species, an Emerging Multidrug-Resistant Bacterium in Hospitals. <i>Antimicrobial Agents and Chemotherapy</i> , 2019, 63, .	1.4	2
53	The clinical features and prognosis of infective endocarditis in the elderly from 2007 to 2016 in a tertiary hospital in China. <i>BMC Infectious Diseases</i> , 2019, 19, 937.	1.3	9
54	Risk factors and outcomes in non-transplant patients with extended-spectrum beta-lactamase-producing Escherichia coli bacteremia: a retrospective study from 2013 to 2016. <i>Antimicrobial Resistance and Infection Control</i> , 2019, 8, 144.	1.5	15

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55	Editorial: Horizontal Gene Transfer Mediated Bacterial Antibiotic Resistance. <i>Frontiers in Microbiology</i> , 2019, 10, 1933.	1.5	136
56	&lt;p&gt;Genomic Analysis Of A KPC-2-Producing &lt;em&gt;Klebsiella Pneumoniae&lt;/em&gt; ST11 Outbreak From A Teaching Hospital In Shandong Province, China&lt;/p&gt;. <i>Infection and Drug Resistance</i> , 2019, Volume 12, 2961-2969.	1.1	16
57	&lt;p&gt;Simulating moxalactam dosage for extended-spectrum Î²-lactamase-producing &lt;em&gt;Enterobacteriaceae&lt;/em&gt; using blood antimicrobial surveillance network data&lt;/p&gt;. <i>Infection and Drug Resistance</i> , 2019, Volume 12, 1199-1208.	1.1	5
58	High prevalence of a globally disseminated hypervirulent clone, <i>Staphylococcus aureus</i> CC121, with reduced vancomycin susceptibility in community settings in China. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 2537-2543.	1.3	12
59	&lt;p&gt;In vitro reduction of colistin susceptibility and comparative genomics reveals multiple differences between MCR-positive and MCR-negative colistin-resistant &lt;em&gt;Escherichia coli&lt;/em&gt;&lt;/p&gt;. <i>Infection and Drug Resistance</i> , 2019, Volume 12, 1665-1674.	1.1	8
60	Detection of an In104-like integron carrying a blaIMP-34 gene in <i>Enterobacter cloacae</i> isolates co-producing IMP-34 and VIM-1. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 2812-2814.	1.3	4
61	A retrospective analysis of risk factors and outcomes in patients with extended-spectrum beta-lactamase-producing <i>Escherichia coli</i> bloodstream infections. <i>Journal of Global Antimicrobial Resistance</i> , 2019, 17, 147-156.	0.9	26
62	A two-step preparation method for nanocrystalline Ag-decorated cotton fabrics and their antibacterial assessment. <i>Journal of Materials Science</i> , 2019, 54, 10447-10456.	1.7	11
63	A retrospective analysis of <i>Pseudomonas aeruginosa</i> bloodstream infections: prevalence, risk factors, and outcome in carbapenem-susceptible and -non-susceptible infections. <i>Antimicrobial Resistance and Infection Control</i> , 2019, 8, 68.	1.5	34
64	Comparison of Tigecycline or Cefoperazone/Sulbactam therapy for bloodstream infection due to Carbapenem-resistant <i>Acinetobacter baumannii</i> . <i>Antimicrobial Resistance and Infection Control</i> , 2019, 8, 52.	1.5	29
65	Occurrence and Genomic Characterization of Two MCR-1-Producing <i>Escherichia coli</i> Isolates from the Same Mink Farmer. <i>MSphere</i> , 2019, 4, .	1.3	13
66	Combined delivery of angiopoietin-1 gene and simvastatin mediated by anti-intercellular adhesion molecule-1 antibody-conjugated ternary nanoparticles for acute lung injury therapy. <i>Nanomedicine: Nanotechnology, Biology, and Medicine</i> , 2019, 15, 25-36.	1.7	34
67	Plasmon enhanced photocatalytic and antimicrobial activities of Ag-TiO <sub>2</sub> nanocomposites under visible light irradiation prepared by DBD cold plasma treatment. <i>Materials Science and Engineering C</i> , 2019, 96, 197-204.	3.8	75
68	Identification and genomic characterization of a KPC-2-, NDM-1- and NDM-5-producing <i>Klebsiella michiganensis</i> isolate. <i>Journal of Antimicrobial Chemotherapy</i> , 2018, 73, 536-538.	1.3	40
69	Silent transmission of an IS 1294b -deactivated mcr-1 gene with inducible colistin resistance. <i>International Journal of Antimicrobial Agents</i> , 2018, 51, 822-828.	1.1	25
70	Complete nucleotide sequences of two KPC-2-encoding plasmids from the same <i>Citrobacter freundii</i> isolate. <i>Journal of Antimicrobial Chemotherapy</i> , 2018, 73, 531-533.	1.3	15
71	Characterization of the population structure, drug resistance mechanisms and plasmids of the community-associated <i>Enterobacter cloacae</i> complex in China. <i>Journal of Antimicrobial Chemotherapy</i> , 2018, 73, 66-76.	1.3	30
72	Study protocol for One Health data collections, analyses and intervention of the Sino-Swedish integrated multisectoral partnership for antibiotic resistance containment (IMPACT). <i>BMJ Open</i> , 2018, 8, e017832.	0.8	26

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73	Discovery and characterisation of an escherichia coli ST206 strain producing NDM-5 and MCR-1 from a patient with acute diarrhoea in China. <i>International Journal of Antimicrobial Agents</i> , 2018, 51, 273-275.	1.1	38
74	Low prevalence of MCR-1-producing <i>Klebsiella pneumoniae</i> in bloodstream infections in China. <i>Clinical Microbiology and Infection</i> , 2018, 24, 205-206.	2.8	19
75	Association between antibiotic consumption and the rate of carbapenem-resistant Gram-negative bacteria from China based on 153 tertiary hospitals data in 2014. <i>Antimicrobial Resistance and Infection Control</i> , 2018, 7, 137.	1.5	73
76	Retrospective comparative analysis of risk factors and outcomes in patients with carbapenem resistant <i>Acinetobacter baumannii</i> bloodstream infections: cefoperazone&ndash;sulbactam associated with resistance and tigecycline increased the mortality. <i>Infection and Drug Resistance</i> , 2018, Volume 11, 2021-2030.	1.1	26
77	Antimicrobial Stewardship in China: Systems, Actions and Future Strategies. <i>Clinical Infectious Diseases</i> , 2018, 67, S135-S141.	2.9	53
78	Characterization of lasR-deficient clinical isolates of <i>Pseudomonas aeruginosa</i> . <i>Scientific Reports</i> , 2018, 8, 13344.	1.6	52
79	Antibacterial effect evaluation of moxalactam against extended-spectrum $\beta$ -lactamase-producing <i>Escherichia coli</i> and <i>Klebsiella pneumoniae</i> with in vitro pharmacokinetics/pharmacodynamics simulation. <i>Infection and Drug Resistance</i> , 2018, Volume 11, 103-112.	1.1	9
80	In vitro antibacterial effect of fosfomycin combination therapy against colistin-resistant <i>Klebsiella pneumoniae</i> . <i>Infection and Drug Resistance</i> , 2018, Volume 11, 577-585.	1.1	7
81	A retrospective, comparative analysis of risk factors and outcomes in carbapenem-susceptible and carbapenem-nonsusceptible <i>Klebsiella pneumoniae</i> bloodstream infections: tigecycline significantly increases the mortality. <i>Infection and Drug Resistance</i> , 2018, Volume 11, 595-606.	1.1	23
82	Emergence of a novel <i>Enterobacter kobei</i> clone carrying chromosomal-encoded CTX-M-12 with diversified pathogenicity in northeast China. <i>New Microbes and New Infections</i> , 2017, 17, 7-10.	0.8	9
83	Genome sequence of <i>Shigella flexneri</i> strain SP1, a diarrheal isolate that encodes an extended-spectrum $\beta$ -lactamase (ESBL). <i>Annals of Clinical Microbiology and Antimicrobials</i> , 2017, 16, 37.	1.7	7
84	Evolution of Drug-resistant <i>Acinetobacter baumannii</i> After DCD Renal Transplantation. <i>Scientific Reports</i> , 2017, 7, 1968.	1.6	1
85	Clinical features and treatment of patients with <i>Vibrio vulnificus</i> infection. <i>International Journal of Infectious Diseases</i> , 2017, 59, 1-6.	1.5	21
86	Community-associated methicillin-resistant <i>Staphylococcus aureus</i> pneumonia in China. <i>Lancet Infectious Diseases</i> , The, 2017, 17, 26.	4.6	6
87	Complete nucleotide sequence of pSKLX3330, an IncI1 plasmid carrying bla CTX-M-55 isolated from community-onset <i>Escherichia coli</i> infection. <i>Journal of Global Antimicrobial Resistance</i> , 2017, 11, 120-122.	0.9	1
88	Genome characterization of two bile-isolated <i>Vibrio fluvialis</i> strains: an insight into pathogenicity and bile salt adaption. <i>Scientific Reports</i> , 2017, 7, 11827.	1.6	14
89	A novel Tn1696-like composite transposon (Tn6404) harboring bla IMP-4 in a <i>Klebsiella pneumoniae</i> isolate carrying a rare ESBL gene bla SFO-1. <i>Scientific Reports</i> , 2017, 7, 17321.	1.6	20
90	In vitro antibacterial activity of fosfomycin combined with other antimicrobials against KPC-producing <i>Klebsiella pneumoniae</i> . <i>International Journal of Antimicrobial Agents</i> , 2017, 50, 237-241.	1.1	31

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91	Complete genome sequencing and genomic characterization of two <i>Escherichia coli</i> strains co-producing MCR-1 and NDM-1 from bloodstream infection. <i>Scientific Reports</i> , 2017, 7, 17885.	1.6	35
92	Alterations of Urinary Microbiota in Type 2 Diabetes Mellitus with Hypertension and/or Hyperlipidemia. <i>Frontiers in Physiology</i> , 2017, 8, 126.	1.3	31
93	Determining optimal dosing regimen of oral administration of dicloxacillin using Monte Carlo simulation. <i>Drug Design, Development and Therapy</i> , 2017, Volume 11, 1951-1956.	2.0	4
94	In vitro Pharmacokinetics/Pharmacodynamics Evaluation of Fosfomycin Combined with Amikacin or Colistin against KPC2-Producing <i>Klebsiella pneumoniae</i> . <i>Frontiers in Cellular and Infection Microbiology</i> , 2017, 7, 246.	1.8	15
95	Molecular Epidemiology and Colistin Resistant Mechanism of mcr-Positive and mcr-Negative Clinical Isolated <i>Escherichia coli</i> . <i>Frontiers in Microbiology</i> , 2017, 8, 2262.	1.5	65
96	Occurrence and Genomic Characterization of ESBL-Producing, MCR-1-Harboring <i>Escherichia coli</i> in Farming Soil. <i>Frontiers in Microbiology</i> , 2017, 8, 2510.	1.5	56
97	The Global Alliance for Infections in Surgery: defining a model for antimicrobial stewardship—results from an international cross-sectional survey. <i>World Journal of Emergency Surgery</i> , 2017, 12, 34.	2.1	47
98	Characterization of the urinary microbiota of elderly women and the effects of type 2 diabetes and urinary tract infections on the microbiota. <i>Oncotarget</i> , 2017, 8, 100678-100690.	0.8	31
99	Dysbiosis of urinary microbiota is positively correlated with Type 2 diabetes mellitus. <i>Oncotarget</i> , 2017, 8, 3798-3810.	0.8	41
100	High Prevalence of ESBL-Producing <i>Klebsiella pneumoniae</i> Causing Community-Onset Infections in China. <i>Frontiers in Microbiology</i> , 2016, 7, 1830.	1.5	50
101	Identification of <i>Raoultella terrigena</i> as a Rare Causative Agent of Subungual Abscess Based on 16S rRNA and Housekeeping Gene Sequencing. <i>Canadian Journal of Infectious Diseases and Medical Microbiology</i> , 2016, 2016, 1-4.	0.7	12
102	Building bridges to operationalise one health – A Sino-Swedish collaboration to tackle antibiotic resistance. <i>One Health</i> , 2016, 2, 139-143.	1.5	18
103	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. <i>International Journal of Antimicrobial Agents</i> , 2016, 48, 409-414.	1.1	24
104	Antimicrobials: a global alliance for optimizing their rational use in intra-abdominal infections (AGORA). <i>World Journal of Emergency Surgery</i> , 2016, 11, 33.	2.1	130
105	Analysis of tigecycline resistance development in clinical <i>Acinetobacter baumannii</i> isolates through a combined genomic and transcriptomic approach. <i>Scientific Reports</i> , 2016, 6, 26930.	1.6	31
106	China's national plan to combat antimicrobial resistance. <i>Lancet Infectious Diseases</i> , The, 2016, 16, 1216-1218.	4.6	58
107	Intrinsic colistin resistance. <i>Lancet Infectious Diseases</i> , The, 2016, 16, 1227-1228.	4.6	5
108	Using Monte Carlo simulation to determine optimal dosing regimen for cefetamet sodium for injection. <i>Journal of Chemotherapy</i> , 2016, 28, 172-179.	0.7	3

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109	Influence of H7N9 virus infection and associated treatment on human gut microbiota. <i>Scientific Reports</i> , 2015, 5, 14771.	1.6	88
110	Severe infective endocarditis with systemic embolism due to community associated methicillin-resistant <i>Staphylococcus aureus</i> ST630. <i>Brazilian Journal of Infectious Diseases</i> , 2015, 19, 85-89.	0.3	13
111	Molecular Epidemiology and Genetic Diversity of Fluoroquinolone-Resistant <i>Escherichia coli</i> Isolates from Patients with Community-Onset Infections in 30 Chinese County Hospitals. <i>Journal of Clinical Microbiology</i> , 2015, 53, 766-770.	1.8	54
112	Bacterial-resistance among outpatients of county hospitals in China: significant geographic distinctions and minor differences between central cities. <i>Microbes and Infection</i> , 2015, 17, 417-425.	1.0	32
113	Emergence of <i>Raoultella ornithinolytica</i> Coproducing IMP-4 and KPC-2 Carbapenemases in China. <i>Antimicrobial Agents and Chemotherapy</i> , 2015, 59, 7086-7089.	1.4	50
114	Complete genome sequence of <i>Lactobacillus heilongjiangensis</i> DSM 28069T: Insight into its probiotic potential. <i>Journal of Biotechnology</i> , 2015, 216, 65-66.	1.9	3
115	Identification and characterization of <i>cfr</i> -positive <i>Staphylococcus aureus</i> isolates from community-onset infectious patients in a county hospital in China. <i>Journal of Medical Microbiology</i> , 2015, 64, 910-915.	0.7	6
116	Nationwide high prevalence of CTX-M and an increase of CTX-M-55 in <i>Escherichia coli</i> isolated from patients with community-onset infections in Chinese county hospitals. <i>BMC Infectious Diseases</i> , 2014, 14, 659.	1.3	139
117	Genome sequencing and genomic characterization of a tigecycline-resistant <i>Klebsiella pneumoniae</i> strain isolated from the bile samples of a cholangiocarcinoma patient. <i>Gut Pathogens</i> , 2014, 6, 40.	1.6	14
118	Use and Prescription of Antibiotics in Primary Health Care Settings in China. <i>JAMA Internal Medicine</i> , 2014, 174, 1914.	2.6	210
119	High burden of antimicrobial drug resistance in Asia. <i>Journal of Global Antimicrobial Resistance</i> , 2014, 2, 141-147.	0.9	55
120	Legislation of clinical antibiotic use in China. <i>Lancet Infectious Diseases</i> , The, 2013, 13, 189-191.	4.6	86
121	Changes in Chinese Policies to Promote the Rational Use of Antibiotics. <i>PLoS Medicine</i> , 2013, 10, e1001556.	3.9	126
122	The Major Aminoglycoside-Modifying Enzyme AAC(3)-II Found in <i>Escherichia coli</i> Determines a Significant Disparity in Its Resistance to Gentamicin and Amikacin in China. <i>Microbial Drug Resistance</i> , 2012, 18, 42-46.	0.9	27