

Chunjiang Zhao

List of Publications by Year in descending order

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Version: 2024-02-01

38
papers

1,592
citations

361413

20
h-index

315739

38
g-index

44
all docs

44
docs citations

44
times ranked

2077
citing authors

#	ARTICLE	IF	CITATIONS
1	A Practical Approach for Predicting Antimicrobial Phenotype Resistance in <i>Staphylococcus aureus</i> Through Machine Learning Analysis of Genome Data. <i>Frontiers in Microbiology</i> , 2022, 13, 841289.	3.5	15
2	Clinical and microbiological characteristics of adults with hospital-acquired pneumonia: a 10-year prospective observational study in China. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2021, 40, 683-690.	2.9	26
3	Bloodstream Infections Caused by Carbapenem-Resistant Enterobacterales: Risk Factors for Mortality, Antimicrobial Therapy and Treatment Outcomes from a Prospective Multicenter Study. <i>Infection and Drug Resistance</i> , 2021, Volume 14, 731-742.	2.7	23
4	Clinical Profile, Prognostic Factors, and Outcome Prediction in Hospitalized Patients With Bloodstream Infection: Results From a 10-Year Prospective Multicenter Study. <i>Frontiers in Medicine</i> , 2021, 8, 629671.	2.6	16
5	The role of interspecies recombination in the evolution of antibiotic-resistant pneumococci. <i>ELife</i> , 2021, 10, .	6.0	21
6	Evolution of <i>Acinetobacter baumannii</i> in Clinical Bacteremia Patients. <i>Infection and Drug Resistance</i> , 2021, Volume 14, 3553-3562.	2.7	3
7	Investigation of Antibiotic Resistance, Serotype Distribution, and Genetic Characteristics of 164 Invasive <i>Streptococcus pneumoniae</i> from North China Between April 2016 and October 2017. <i>Infection and Drug Resistance</i> , 2020, Volume 13, 2117-2128.	2.7	11
8	Long-Term Continuous Antimicrobial Resistance Surveillance Among Nosocomial Gram-Negative Bacilli in China from 2010 to 2018 (CMSS). <i>Infection and Drug Resistance</i> , 2020, Volume 13, 2617-2629.	2.7	12
9	In vitro activity of the novel β -lactamase inhibitor taniborbactam (VNRX-5133), in combination with cefepime or meropenem, against MDR Gram-negative bacterial isolates from China. <i>Journal of Antimicrobial Chemotherapy</i> , 2020, 75, 1850-1858.	3.0	32
10	Evaluation of the in vitro activity of new polymyxin B analogue SPR206 against clinical MDR, colistin-resistant and tigecycline-resistant Gram-negative bacilli. <i>Journal of Antimicrobial Chemotherapy</i> , 2020, 75, 2609-2615.	3.0	30
11	Prospective multi-center evaluation on risk factors, clinical characteristics and outcomes due to carbapenem resistance in <i>Acinetobacter baumannii</i> complex bacteraemia: experience from the Chinese Antimicrobial Resistance Surveillance of Nosocomial Infections (CARES) Network. <i>Journal of Medical Microbiology</i> , 2020, 69, 949-959.	1.8	22
12	A retrospective study on Xpert MTB/RIF for detection of tuberculosis in a teaching hospital in China. <i>BMC Infectious Diseases</i> , 2020, 20, 362.	2.9	10
13	In vitro activities of Eravacycline against 336 isolates collected from 2012 to 2016 from 11 teaching hospitals in China. <i>BMC Infectious Diseases</i> , 2019, 19, 508.	2.9	16
14	A multicenter epidemiology study on the risk factors and clinical outcomes of nosocomial intra-abdominal infections in China: results from the Chinese antimicrobial resistance surveillance of nosocomial infections (CARES) 2007–2016. <i>Infection and Drug Resistance</i> , 2018, Volume 11, 2311-2319.	2.7	16
15	The Changing Pattern of Population Structure of <i>Staphylococcus aureus</i> from Bacteremia in China from 2013 to 2016: ST239-030-MRSA Replaced by ST59-t437. <i>Frontiers in Microbiology</i> , 2018, 9, 332.	3.5	95
16	Serotype distribution and antibiotic resistance of <i>Streptococcus pneumoniae</i> isolates from 17 Chinese cities from 2011 to 2016. <i>BMC Infectious Diseases</i> , 2017, 17, 804.	2.9	45
17	Antimicrobial susceptibility of <i>Streptococcus pneumoniae</i> , <i>Haemophilus influenzae</i> and <i>Moraxella catarrhalis</i> isolated from community-acquired respiratory tract infections in China: Results from the CARTIPS Antimicrobial Surveillance Program. <i>Journal of Global Antimicrobial Resistance</i> , 2016, 5, 36-41.	2.2	27
18	High Prevalence of Hypervirulent <i>Klebsiella pneumoniae</i> Infection in China: Geographic Distribution, Clinical Characteristics, and Antimicrobial Resistance. <i>Antimicrobial Agents and Chemotherapy</i> , 2016, 60, 6115-6120.	3.2	222

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19	In vitro activities of tedizolid compared with other antibiotics against Gram-positive pathogens associated with hospital-acquired pneumonia, skin and soft tissue infection and bloodstream infection collected from 26 hospitals in China. <i>Journal of Medical Microbiology</i> , 2016, 65, 1215-1224.	1.8	20
20	Comparative evaluation of tigecycline susceptibility testing methods for <i>Acinetobacter baumannii</i> and <i>Enterobacteriaceae</i> . <i>Journal of Global Antimicrobial Resistance</i> , 2015, 3, 75-79.	2.2	4
21	Genetic characterisation of clinical <i>Klebsiella pneumoniae</i> isolates with reduced susceptibility to tigecycline: Role of the global regulator RamA and its local repressor RamR. <i>International Journal of Antimicrobial Agents</i> , 2015, 45, 635-640.	2.5	52
22	Emergence of a hypervirulent carbapenem-resistant <i>Klebsiella pneumoniae</i> isolate from clinical infections in China. <i>Journal of Infection</i> , 2015, 71, 553-560.	3.3	176
23	Evolution of Carbapenem-Resistant <i>Acinetobacter baumannii</i> Revealed through Whole-Genome Sequencing and Comparative Genomic Analysis. <i>Antimicrobial Agents and Chemotherapy</i> , 2015, 59, 1168-1176.	3.2	56
24	<i>In Vitro</i> Activities of Ceftazidime-Avibactam and Aztreonam-Avibactam against 372 Gram-Negative Bacilli Collected in 2011 and 2012 from 11 Teaching Hospitals in China. <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 1774-1778.	3.2	81
25	An Outbreak of a Nosocomial NDM-1-Producing <i>Klebsiella pneumoniae</i> ST147 at a Teaching Hospital in Mainland China. <i>Microbial Drug Resistance</i> , 2014, 20, 144-149.	2.0	57
26	In vitro antibacterial activities of two novel oral antibiotics, tebipenem and cefditoren, and other comparators against community-acquired respiratory tract infection-associated bacterial pathogens: A multicentre study in China. <i>International Journal of Antimicrobial Agents</i> , 2014, 43, 92-93.	2.5	0
27	Efficacy and safety of daptomycin for the treatment of infectious disease: a meta-analysis based on randomized controlled trials. <i>Journal of Antimicrobial Chemotherapy</i> , 2014, 69, 3181-3189.	3.0	26
28	In vitro antimicrobial activity of the novel oxazolidinone tedizolid and comparator agents against <i>Staphylococcus aureus</i> and linezolid-resistant Gram-positive pathogens: a multicentre study in China. <i>International Journal of Antimicrobial Agents</i> , 2014, 44, 276-277.	2.5	18
29	Insights on evolution of virulence and resistance from the whole-genome analysis of a predominant methicillin-resistant <i>Staphylococcus aureus</i> clone sequence type 239 in China. <i>Science Bulletin</i> , 2014, 59, 1104-1112.	1.7	7
30	Novel NDM-9 metallo- β -lactamase identified from a ST107 <i>Klebsiella pneumoniae</i> strain isolated in China. <i>International Journal of Antimicrobial Agents</i> , 2014, 44, 90-91.	2.5	48
31	Population structure and characterisation of <i>Staphylococcus aureus</i> from bacteraemia at multiple hospitals in China: association between antimicrobial resistance, toxin genes and genotypes. <i>International Journal of Antimicrobial Agents</i> , 2013, 42, 211-219.	2.5	84
32	Prevalence and molecular typing of oxacillin-susceptible <i>mecA</i> -positive <i>Staphylococcus aureus</i> from multiple hospitals in China. <i>Diagnostic Microbiology and Infectious Disease</i> , 2013, 77, 267-269.	1.8	22
33	Phenotypic and Genotypic Characteristic of Invasive Pneumococcal Isolates from Both Children and Adult Patients from a Multicenter Surveillance in China 2005–2011. <i>PLoS ONE</i> , 2013, 8, e82361.	2.5	38
34	Identification of Gene Clusters Associated with Host Adaptation and Antibiotic Resistance in Chinese <i>Staphylococcus aureus</i> Isolates by Microarray-Based Comparative Genomics. <i>PLoS ONE</i> , 2013, 8, e53341.	2.5	12
35	Comparative Proteomics-Based Identification of Genes Associated with Glycopeptide Resistance in Clinically Derived Heterogeneous Vancomycin-Intermediate <i>Staphylococcus aureus</i> Strains. <i>PLoS ONE</i> , 2013, 8, e66880.	2.5	23
36	Antimicrobial resistance trends among 5608 clinical Gram-positive isolates in China: results from the Gram-Positive Cocci Resistance Surveillance program (2005–2010). <i>Diagnostic Microbiology and Infectious Disease</i> , 2012, 73, 174-181.	1.8	65

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37	Characterization of Community Acquired <i>Staphylococcus aureus</i> Associated with Skin and Soft Tissue Infection in Beijing: High Prevalence of PVL+ ST398. PLoS ONE, 2012, 7, e38577.	2.5	98
38	Prevalence and Characterization of Heterogeneous Vancomycin-Intermediate <i>Staphylococcus aureus</i> Isolates from 14 Cities in China. Antimicrobial Agents and Chemotherapy, 2009, 53, 3642-3649.	3.2	51