

Yunsong Yu

List of Publications by Year in descending order

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191
papers

5,322
citations

117625

34
h-index

138484

58
g-index

196
all docs

196
docs citations

196
times ranked

4551
citing authors

#	ARTICLE	IF	CITATIONS
1	European Society of Clinical Microbiology and Infectious Diseases (ESCMID) guidelines for the treatment of infections caused by multidrug-resistant Gram-negative bacilli (endorsed by European) TJ ETQq1 1 0.784314 rg324 Overl	10.7	140
2	Prevalence of <i>mcr-1</i> in <i>Escherichia coli</i> and <i>Klebsiella pneumoniae</i> recovered from bloodstream infections in China: a multicentre longitudinal study. <i>Lancet Infectious Diseases</i> , The, 2017, 17, 400-410.	9.1	177
3	Novel Genetic Environment of the Carbapenem-Hydrolyzing $\hat{1}^2$ -Lactamase KPC-2 among <i>Enterobacteriaceae</i> in China. <i>Antimicrobial Agents and Chemotherapy</i> , 2009, 53, 4333-4338.	3.2	173
4	Resistance of strains producing extended-spectrum $\hat{1}^2$ -lactamases and genotype distribution in China. <i>Journal of Infection</i> , 2007, 54, 53-57.	3.3	164
5	Contamination-free visual detection of SARS-CoV-2 with CRISPR/Cas12a: A promising method in the point-of-care detection. <i>Biosensors and Bioelectronics</i> , 2020, 169, 112642.	10.1	136
6	BacWGSTdb 2.0: a one-stop repository for bacterial whole-genome sequence typing and source tracking. <i>Nucleic Acids Research</i> , 2021, 49, D644-D650.	14.5	129
7	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
8	Complete Nucleotide Sequence of <i>Klebsiella pneumoniae</i> Multidrug Resistance Plasmid pKP048, Carrying <i>bla</i> _{KPC-2} , <i>bla</i> _{DHA-1} , <i>qnrB4</i> , and <i>armA</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2010, 54, 3967-3969.	3.2	121
9	Genomic Analysis of the Multidrug-Resistant <i>Acinetobacter baumannii</i> Strain MDR-ZJ06 Widely Spread in China. <i>Antimicrobial Agents and Chemotherapy</i> , 2011, 55, 4506-4512.	3.2	116
10	A global perspective on the convergence of hypervirulence and carbapenem resistance in <i>Klebsiella pneumoniae</i> . <i>Journal of Global Antimicrobial Resistance</i> , 2021, 25, 26-34.	2.2	110
11	High prevalence of ESBL-producing <i>Escherichia coli</i> and <i>Klebsiella pneumoniae</i> in community-onset bloodstream infections in China. <i>Journal of Antimicrobial Chemotherapy</i> , 2017, 72, 273-280.	3.0	93
12	High Incidence and Endemic Spread of NDM-1-Positive <i>Enterobacteriaceae</i> in Henan Province, China. <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 4275-4282.	3.2	90
13	Wide distribution of CC92 carbapenem-resistant and OXA-23-producing <i>Acinetobacter baumannii</i> in multiple provinces of China. <i>International Journal of Antimicrobial Agents</i> , 2013, 42, 322-328.	2.5	88
14	Dissemination of <i>bla</i> _{NDM-5} gene via an <i>IncX3</i> -type plasmid among non-clonal <i>Escherichia coli</i> in China. <i>Antimicrobial Resistance and Infection Control</i> , 2018, 7, 59.	4.1	84
15	Dissemination of a clone carrying a <i>fosA3</i> -harbouring plasmid mediates high fosfomycin resistance rate of KPC-producing <i>Klebsiella pneumoniae</i> in China. <i>International Journal of Antimicrobial Agents</i> , 2015, 45, 66-70.	2.5	77
16	Triclosan resistance in clinical isolates of <i>Acinetobacter baumannii</i> . <i>Journal of Medical Microbiology</i> , 2009, 58, 1086-1091.	1.8	74
17	The Rapid Emergence of Tigecycline Resistance in <i>bla</i> _{KPC-2} Harboring <i>Klebsiella pneumoniae</i> , as Mediated in Vivo by Mutation in <i>tetA</i> During Tigecycline Treatment. <i>Frontiers in Microbiology</i> , 2018, 9, 648.	3.5	73
18	Decreased susceptibility to tigecycline in <i>Acinetobacter baumannii</i> mediated by a mutation in <i>trm</i> encoding SAM-dependent methyltransferase. <i>Journal of Antimicrobial Chemotherapy</i> , 2014, 69, 72-76.	3.0	72

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19	Tigecycline Susceptibility and the Role of Efflux Pumps in Tigecycline Resistance in KPC-Producing <i>Klebsiella pneumoniae</i> . PLoS ONE, 2015, 10, e0119064.	2.5	72
20	The global dissemination of bacterial infections necessitates the study of reverse genomic epidemiology. Briefings in Bioinformatics, 2020, 21, 741-750.	6.5	56
21	A Highly Efficient CRISPR-Cas9-Based Genome Engineering Platform in <i>Acinetobacter baumannii</i> to Understand the H ₂ O ₂ -Sensing Mechanism of OxyR. Cell Chemical Biology, 2019, 26, 1732-1742.e5.	5.2	55
22	Characterization of a Novel Plasmid Type and Various Genetic Contexts of bla _{OXA-58} in <i>Acinetobacter</i> spp. from Multiple Cities in China. PLoS ONE, 2014, 9, e84680.	2.5	52
23	Colistin Resistance in <i>Acinetobacter baumannii</i> MDR-ZJ06 Revealed by a Multiomics Approach. Frontiers in Cellular and Infection Microbiology, 2017, 7, 45.	3.9	50
24	Risk factors for acquisition and mortality of multidrug-resistant <i>Acinetobacter baumannii</i> bacteremia. Medicine (United States), 2019, 98, e14937.	1.0	50
25	Colistin-phage combinations decrease antibiotic resistance in <i>Acinetobacter baumannii</i> via changes in envelope architecture. Emerging Microbes and Infections, 2021, 10, 2205-2219.	6.5	50
26	Resistance evolution of hypervirulent carbapenem-resistant <i>Klebsiella pneumoniae</i> ST11 during treatment with tigecycline and polymyxin. Emerging Microbes and Infections, 2021, 10, 1129-1136.	6.5	49
27	Distribution of the ACME-arcA gene among methicillin-resistant <i>Staphylococcus haemolyticus</i> and identification of a novel ccr allotype in ACME-arcA-positive isolates. Journal of Medical Microbiology, 2009, 58, 731-736.	1.8	48
28	The role of the type VI secretion system vgrG gene in the virulence and antimicrobial resistance of <i>Acinetobacter baumannii</i> ATCC 19606. PLoS ONE, 2018, 13, e0192288.	2.5	48
29	Species Distribution of Clinical <i>Acinetobacter</i> Isolates Revealed by Different Identification Techniques. PLoS ONE, 2014, 9, e104882.	2.5	48
30	The Effect of Colistin Resistance-Associated Mutations on the Fitness of <i>Acinetobacter baumannii</i> . Frontiers in Microbiology, 2016, 7, 1715.	3.5	47
31	OXA-23 Is a Prevalent Mechanism Contributing to Sulbactam Resistance in Diverse <i>Acinetobacter baumannii</i> Clinical Strains. Antimicrobial Agents and Chemotherapy, 2019, 63, .	3.2	47
32	Diversity of virulence level phenotype of hypervirulent <i>Klebsiella pneumoniae</i> from different sequence type lineage. BMC Microbiology, 2018, 18, 94.	3.3	46
33	Clinical Impact of Metagenomic Next-Generation Sequencing of Bronchoalveolar Lavage in the Diagnosis and Management of Pneumonia. Journal of Molecular Diagnostics, 2021, 23, 1259-1268.	2.8	43
34	Detection of an <i>Escherichia coli</i> Sequence Type 167 Strain with Two Tandem Copies of bla _{NDM-1} in the Chromosome. Journal of Clinical Microbiology, 2017, 55, 199-205.	3.9	42
35	Global transcriptional response of <i>Acinetobacter baumannii</i> to a subinhibitory concentration of tigecycline. International Journal of Antimicrobial Agents, 2014, 44, 337-344.	2.5	39
36	Emergence of Ceftazidime- and Avibactam-Resistant <i>Klebsiella pneumoniae</i> Carbapenemase-Producing <i>Pseudomonas aeruginosa</i> in China. MSystems, 2021, 6, e0078721.	3.8	39

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37	A Biological Inventory of Prophages in <i>A. baumannii</i> Genomes Reveal Distinct Distributions in Classes, Length, and Genomic Positions. <i>Frontiers in Microbiology</i> , 2020, 11, 579802.	3.5	38
38	BacAnt: A Combination Annotation Server for Bacterial DNA Sequences to Identify Antibiotic Resistance Genes, Integrons, and Transposable Elements. <i>Frontiers in Microbiology</i> , 2021, 12, 649969.	3.5	38
39	Dissemination of NDM-1-Producing Enterobacteriaceae Mediated by the IncX3-Type Plasmid. <i>PLoS ONE</i> , 2015, 10, e0129454.	2.5	38
40	Step-Wise Increase in Tigecycline Resistance in <i>Klebsiella pneumoniae</i> Associated with Mutations in <i>ramR</i> , <i>lon</i> and <i>rpsJ</i> . <i>PLoS ONE</i> , 2016, 11, e0165019.	2.5	38
41	A 10 year surveillance for antimicrobial susceptibility of <i>Escherichia coli</i> and <i>Klebsiella pneumoniae</i> in community- and hospital-associated intra-abdominal infections in China. <i>Journal of Medical Microbiology</i> , 2013, 62, 1343-1349.	1.8	36
42	Molecular Epidemiology and Mechanism of Sulbactam Resistance in <i>Acinetobacter baumannii</i> Isolates with Diverse Genetic Backgrounds in China. <i>Antimicrobial Agents and Chemotherapy</i> , 2018, 62, .	3.2	35
43	High prevalence of colistin resistance and <i>mcr-9/10</i> genes in <i>Enterobacter</i> spp. in a tertiary hospital over a decade. <i>International Journal of Antimicrobial Agents</i> , 2022, 59, 106573.	2.5	35
44	In vitro activity of flomoxef and comparators against <i>Escherichia coli</i> , <i>Klebsiella pneumoniae</i> and <i>Proteus mirabilis</i> producing extended-spectrum β -lactamases in China. <i>International Journal of Antimicrobial Agents</i> , 2015, 45, 485-490.	2.5	34
45	Detection and characterization of a clinical <i>Escherichia coli</i> ST3204 strain coproducing NDM-16 and MCR-1. <i>Infection and Drug Resistance</i> , 2018, Volume 11, 1189-1195.	2.7	33
46	Target-oriented design and biosynthesis of thioStrepton-derived thiopeptide antibiotics with improved pharmaceutical properties. <i>Organic Chemistry Frontiers</i> , 2015, 2, 106-109.	4.5	32
47	Comparison of the ability to identify arterial stiffness between two new anthropometric indices and classical obesity indices in Chinese adults. <i>Atherosclerosis</i> , 2017, 263, 263-271.	0.8	31
48	Epidemiology, evolution and cryptic susceptibility of methicillin-resistant <i>Staphylococcus aureus</i> in China: a whole-genome-based survey. <i>Clinical Microbiology and Infection</i> , 2022, 28, 85-92.	6.0	31
49	Tigecycline resistance caused by <i>rpsJ</i> evolution in a 59-year-old male patient infected with KPC-producing <i>Klebsiella pneumoniae</i> during tigecycline treatment. <i>Infection, Genetics and Evolution</i> , 2018, 66, 188-191.	2.3	30
50	Cointegration as a mechanism for the evolution of a KPC-producing multidrug resistance plasmid in <i>Proteus mirabilis</i> . <i>Emerging Microbes and Infections</i> , 2020, 9, 1206-1218.	6.5	30
51	Basis of Virulence in a Panton-Valentine Leukocidin-Negative Community-Associated Methicillin-Resistant <i>Staphylococcus aureus</i> Strain. <i>Journal of Infectious Diseases</i> , 2015, 211, 472-480.	4.0	29
52	Toxin profiles, PCR ribotypes and resistance patterns of <i>Clostridium difficile</i> : a multicentre study in China, 2012-2013. <i>International Journal of Antimicrobial Agents</i> , 2016, 48, 736-739.	2.5	29
53	Evolution of <i>Acinetobacter baumannii</i> In Vivo: International Clone II, More Resistance to Ceftazidime, Mutation in <i>ptk</i> . <i>Frontiers in Microbiology</i> , 2017, 8, 1256.	3.5	29
54	Emergence of High-Level Cefiderocol Resistance in Carbapenem-Resistant <i>Klebsiella pneumoniae</i> from Bloodstream Infections in Patients with Hematologic Malignancies in China. <i>Microbiology Spectrum</i> , 2022, 10, e0008422.	3.0	29

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55	Clinical relevance and plasmid dynamics of <i>mcr-1</i> -positive <i>Escherichia coli</i> in China: a multicentre case-control and molecular epidemiological study. <i>Lancet Microbe</i> , The, 2020, 1, e24-e33.	7.3	28
56	Cefepime combined with amoxicillin/clavulanic acid: a new choice for the KPC-producing <i>K. pneumoniae</i> infection. <i>International Journal of Infectious Diseases</i> , 2015, 38, 108-114.	3.3	27
57	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
58	Emergence of tigecycline resistance in <i>Escherichia coli</i> ; co-producing MCR-1 and NDM-5 during tigecycline salvage treatment. <i>Infection and Drug Resistance</i> , 2018, Volume 11, 2241-2248.	2.7	27
59	Transferable <i>Acinetobacter baumannii</i> plasmid pDETAB2 encodes OXA-58 and NDM-1 and represents a new class of antibiotic resistance plasmids. <i>Journal of Antimicrobial Chemotherapy</i> , 2021, 76, 1130-1134.	3.0	27
60	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
61	Update of incidence and antimicrobial susceptibility trends of <i>Escherichia coli</i> and <i>Klebsiella pneumoniae</i> isolates from Chinese intra-abdominal infection patients. <i>BMC Infectious Diseases</i> , 2017, 17, 776.	2.9	26
62	Mechanism of eravacycline resistance in <i>Acinetobacter baumannii</i> mediated by a deletion mutation in the sensor kinase <i>adeS</i> , leading to elevated expression of the efflux pump <i>AdeABC</i> . <i>Infection, Genetics and Evolution</i> , 2020, 80, 104185.	2.3	26
63	High percentage of the ceftriaxone-resistant <i>Neisseria gonorrhoeae</i> FC428 clone among isolates from a single hospital in Hangzhou, China. <i>Journal of Antimicrobial Chemotherapy</i> , 2021, 76, 936-939.	3.0	26
64	A Genomics Based Discovery of Secondary Metabolite Biosynthetic Gene Clusters in <i>Aspergillus ustus</i> . <i>PLoS ONE</i> , 2015, 10, e0116089.	2.5	25
65	Capsule Thickness, Not Biofilm Formation, Gives Rise to Mucoïd <i>Acinetobacter baumannii</i> Phenotypes That are More Prevalent in Long-Term Infections: A Study of Clinical Isolates from a Hospital in China. <i>Infection and Drug Resistance</i> , 2020, Volume 13, 99-109.	2.7	25
66	Complete genome sequence of <i>Acinetobacter baumannii</i> XH386 (ST208), a multi-drug resistant bacteria isolated from pediatric hospital in China. <i>Genomics Data</i> , 2016, 7, 269-274.	1.3	24
67	Molecular characteristics of extended-spectrum β -lactamase-producing <i>Escherichia coli</i> and <i>Klebsiella pneumoniae</i> causing intra-abdominal infections from 9 tertiary hospitals in China. <i>Diagnostic Microbiology and Infectious Disease</i> , 2017, 87, 45-48.	1.8	23
68	Dual Role of <i>gnaA</i> in Antibiotic Resistance and Virulence in <i>Acinetobacter baumannii</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2019, 63, .	3.2	23
69	Mechanical penetration of β -lactam-resistant Gram-negative bacteria by programmable nanowires. <i>Science Advances</i> , 2020, 6, .	10.3	23
70	Defining persistent critical illness based on growth trajectories in patients with sepsis. <i>Critical Care</i> , 2020, 24, 57.	5.8	23
71	A Novel SXT/R391 Integrative and Conjugative Element Carries Two Copies of the <i>bla</i> _{NDM-1} Gene in <i>Proteus mirabilis</i> . <i>MSphere</i> , 2021, 6, e0058821.	2.9	23
72	Risk factors for infection and mortality caused by carbapenem-resistant <i>Klebsiella pneumoniae</i> : A large multicentre case-control and cohort study. <i>Journal of Infection</i> , 2022, 84, 637-647.	3.3	23

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73	Production of a Promising Biosynthetic Self-Assembled Nanoconjugate Vaccine against <i>Klebsiella Pneumoniae</i> Serotype O2 in a General <i>Escherichia Coli</i> Host. <i>Advanced Science</i> , 2021, 8, e2100549.	11.2	22
74	Identification of Novel Conjugative Plasmids with Multiple Copies of <i>fosB</i> that Confer High-Level Fosfomycin Resistance to Vancomycin-Resistant Enterococci. <i>Frontiers in Microbiology</i> , 2017, 8, 1541.	3.5	21
75	Using Core-genome Multilocus Sequence Typing to Monitor the Changing Epidemiology of Methicillin-resistant <i>Staphylococcus aureus</i> in a Teaching Hospital. <i>Clinical Infectious Diseases</i> , 2018, 67, S241-S248.	5.8	21
76	Antimicrobial Susceptibilities of Aerobic and Facultative Gram-Negative Bacilli from Intra-abdominal Infections in Patients from Seven Regions in China in 2012 and 2013. <i>Antimicrobial Agents and Chemotherapy</i> , 2016, 60, 245-251.	3.2	20
77	Novel tigecycline resistance mechanisms in <i>Acinetobacter baumannii</i> mediated by mutations in <i>adeS</i> , <i>rpoB</i> and <i>rrf</i> . <i>Emerging Microbes and Infections</i> , 2021, 10, 1404-1417.	6.5	20
78	In Vitro Activity of Imipenem/Relebactam Against Enterobacteriaceae Isolates Obtained from Intra-abdominal, Respiratory Tract, and Urinary Tract Infections in China: Study for Monitoring Antimicrobial Resistance Trends (SMART), 2015–2018. <i>Clinical Infectious Diseases</i> , 2020, 71, S427-S435.	5.8	20
79	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
80	Concurrent modifications of the C-terminus and side ring of thiostrepton and their synergistic effects with respect to improving antibacterial activities. <i>Organic Chemistry Frontiers</i> , 2016, 3, 496-500.	4.5	19
81	Molecular Characterization of Carbapenem-Resistant <i>Serratia marcescens</i> ; Clinical Isolates in a Tertiary Hospital in Hangzhou, China. <i>Infection and Drug Resistance</i> , 2020, Volume 13, 999-1008.	2.7	19
82	Whole Genome Analysis of a Community-Associated Methicillin-Resistant <i>Staphylococcus aureus</i> ST59 Isolate from a Case of Human Sepsis and Severe Pneumonia in China. <i>PLoS ONE</i> , 2014, 9, e89235.	2.5	19
83	Complete Genome Sequence of <i>Klebsiella pneumoniae</i> Sequence Type 17, a Multidrug-Resistant Strain Isolated during Tigecycline Treatment. <i>Genome Announcements</i> , 2014, 2, .	0.8	18
84	The Characterization of OXA-232 Carbapenemase-Producing ST437 <i>Klebsiella pneumoniae</i> in China. <i>Canadian Journal of Infectious Diseases and Medical Microbiology</i> , 2020, 2020, 1-5.	1.9	18
85	<i>Alcaligenes faecalis</i> metallo- β -lactamase in extensively drug-resistant <i>Pseudomonas aeruginosa</i> isolates. <i>Clinical Microbiology and Infection</i> , 2022, 28, 880.e1-880.e8.	6.0	18
86	Multiplication of <i>bla</i> _{OXA-23} is common in clinical <i>Acinetobacter baumannii</i> , but does not enhance carbapenem resistance. <i>Journal of Antimicrobial Chemotherapy</i> , 2016, 71, 3381-3385.	3.0	17
87	Rapid emergence of high-level tigecycline resistance in <i>Escherichia coli</i> strains harbouring <i>bla</i> _{NDM-5} in vivo. <i>International Journal of Antimicrobial Agents</i> , 2016, 47, 324-327.	2.5	17
88	Antimicrobial susceptibilities of aerobic and facultative gram-negative bacilli isolated from Chinese patients with urinary tract infections between 2010 and 2014. <i>BMC Infectious Diseases</i> , 2017, 17, 192.	2.9	17
89	Increasing prevalence of <i>Neisseria gonorrhoeae</i> with decreased susceptibility to ceftriaxone and resistance to azithromycin in Hangzhou, China (2015–17). <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 29-37.	3.0	17
90	Predominance of ST5-II-t311 clone among healthcare-associated methicillin-resistant <i>Staphylococcus aureus</i> isolates recovered from Zhejiang, China. <i>International Journal of Infectious Diseases</i> , 2018, 71, 107-112.	3.3	17

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91	Core Genome Allelic Profiles of Clinical <i>Klebsiella pneumoniae</i> Strains Using a Random Forest Algorithm Based on Multilocus Sequence Typing Scheme for Hypervirulence Analysis. <i>Journal of Infectious Diseases</i> , 2020, 221, S263-S271.	4.0	17
92	Etiology and prevalence of ESBLs in adult community-onset urinary tract infections in East China: A prospective multicenter study. <i>Journal of Infection</i> , 2021, 83, 175-181.	3.3	17
93	Pooled Plasmid Sequencing Reveals the Relationship Between Mobile Genetic Elements and Antimicrobial Resistance Genes in Clinically Isolated <i>Klebsiella pneumoniae</i> . <i>Genomics, Proteomics and Bioinformatics</i> , 2020, 18, 539-548.	6.9	17
94	Diversity and evolution of oligopeptide permease systems in staphylococcal species. <i>Genomics</i> , 2014, 104, 8-13.	2.9	16
95	The Effect of Hepatosteatosis on Response to Antiviral Treatment in Patients with Chronic Hepatitis B: A Meta-Analysis. <i>Gastroenterology Research and Practice</i> , 2017, 2017, 1-12.	1.5	16
96	Prevalence and molecular characteristics of <i>mcr-1</i> gene in <i>Salmonella typhimurium</i> in a tertiary hospital of Zhejiang Province. <i>Infection and Drug Resistance</i> , 2019, Volume 12, 105-110.	2.7	16
97	Tandem amplification of the <i>vanM</i> gene cluster drives vancomycin resistance in vancomycin-variable enterococci. <i>Journal of Antimicrobial Chemotherapy</i> , 2020, 75, 283-291.	3.0	16
98	In-Host Evolution of Daptomycin Resistance and Heteroresistance in Methicillin-Resistant <i>Staphylococcus aureus</i> Strains From Three Endocarditis Patients. <i>Journal of Infectious Diseases</i> , 2020, 221, S243-S252.	4.0	16
99	Correlation between <i>Ureaplasma</i> Subgroup 2 and Genitourinary Tract Disease Outcomes Revealed by an Expanded Multilocus Sequence Typing (eMLST) Scheme. <i>PLoS ONE</i> , 2014, 9, e104347.	2.5	15
100	In Vitro Activities of Ceftaroline/Avibactam, Ceftazidime/Avibactam, and Other Comparators Against Pathogens From Various Complicated Infections in China. <i>Clinical Infectious Diseases</i> , 2018, 67, S206-S216.	5.8	15
101	Risk factors and outcomes of bloodstream infections caused by <i>Acinetobacter baumannii</i> : a case-control study. <i>Diagnostic Microbiology and Infectious Disease</i> , 2021, 99, 115229.	1.8	15
102	Safety and immunogenicity of a new glycoengineered vaccine against <i>Acinetobacter baumannii</i> in mice. <i>Microbial Biotechnology</i> , 2022, 15, 703-716.	4.2	15
103	Phenotypic and Genotypic Characterization of a Hypervirulent Carbapenem-Resistant <i>Klebsiella pneumoniae</i> ST17-KL38 Clinical Isolate Harboring the Carbapenemase IMP-4. <i>Microbiology Spectrum</i> , 2022, 10, e0213421.	3.0	15
104	Characterization of <i>vanM</i> carrying clinical <i>Enterococcus</i> isolates and diversity of the suppressed <i>vanM</i> gene cluster. <i>Infection, Genetics and Evolution</i> , 2019, 68, 145-152.	2.3	14
105	Emergence of a KPC Variant Conferring Resistance to Ceftazidime-Avibactam in a Widespread ST11 Carbapenem-Resistant <i>Klebsiella pneumoniae</i> Clone in China. <i>Frontiers in Microbiology</i> , 2021, 12, 724272.	3.5	14
106	Clinical and Microbiological Characteristics of Community-Onset Carbapenem-Resistant Enterobacteriaceae Isolates. <i>Infection and Drug Resistance</i> , 2020, Volume 13, 3131-3143.	2.7	14
107	Carbapenem susceptibilities of Gram-negative pathogens in intra-abdominal and urinary tract infections: updated report of SMART 2015 in China. <i>BMC Infectious Diseases</i> , 2018, 18, 493.	2.9	13
108	Detection and analysis of two cases of the internationally spreading ceftriaxone-resistant <i>Neisseria gonorrhoeae</i> FC428 clone in China. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 3635-3636.	3.0	13

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109	Plasmid Dynamics of <i>mcr-1</i> -Positive <i>Salmonella</i> spp. in a General Hospital in China. <i>Frontiers in Microbiology</i> , 2020, 11, 604710.	3.5	13
110	Acquisition of a genomic resistance island (AbGR15) from global clone 2 through homologous recombination in a clinical <i>Acinetobacter baumannii</i> isolate. <i>Journal of Antimicrobial Chemotherapy</i> , 2021, 76, 65-69.	3.0	13
111	Emergence of carbapenem-resistant <i>Klebsiella pneumoniae</i> harbouring <i>bla</i> OXA-48-like genes in China. <i>Journal of Medical Microbiology</i> , 2021, 70, .	1.8	13
112	Genetic Characterization and Passage Instability of a Hybrid Plasmid Co-Harboring <i>bla</i> IMP-4 and <i>bla</i> NDM-1 Reveal the Contribution of Insertion Sequences During Plasmid Formation and Evolution. <i>Microbiology Spectrum</i> , 2021, 9, e0157721.	3.0	13
113	The mismatch repair system (<i>mutS</i> and <i>mutL</i>) in <i>Acinetobacter baylyi</i> ADP1. <i>BMC Microbiology</i> , 2020, 20, 40.	3.3	12
114	<i>mcr-1</i> Gene Has No Effect on Colistin Resistance When It Coexists with Inactivated <i>mgrB</i> Gene in <i>Klebsiella pneumoniae</i> . <i>Microbial Drug Resistance</i> , 2018, 24, 1117-1120.	2.0	11
115	Prevalence of Fosfomycin Resistance in Methicillin-Resistant <i>Staphylococcus aureus</i> ; Isolated from Patients in a University Hospital in China from 2013 to 2015. <i>Japanese Journal of Infectious Diseases</i> , 2018, 71, 312-314.	1.2	11
116	Coexistence of <i>bla</i> KPC-2- <i>IncN</i> and <i>mcr-1-IncX4</i> plasmids in a ST48 <i>Escherichia coli</i> strain in China. <i>Journal of Global Antimicrobial Resistance</i> , 2020, 23, 149-153.	2.2	11
117	The distribution of mutations and hotspots in transcription regulators of resistance-nodulation-cell division efflux pumps in tigecycline non-susceptible <i>Acinetobacter baumannii</i> in China. <i>International Journal of Medical Microbiology</i> , 2020, 310, 151464.	3.6	11
118	Co-harboring of Novel <i>bla</i> KPC-2 Plasmid and Integrative and Conjugative Element Carrying Tn6203 in Multidrug-Resistant <i>Pseudomonas aeruginosa</i> . <i>Frontiers in Microbiology</i> , 2021, 12, 674974.	3.5	11
119	Metagenomic sequencing with spiked-in internal control to monitor cellularity and diagnosis of pneumonia. <i>Journal of Infection</i> , 2022, 84, e13-e17.	3.3	11
120	Characterization of the Staphylococcal Cassette Chromosome Composite Island of <i>Staphylococcus haemolyticus</i> SH32, a Methicillin-Resistant Clinical Isolate from China. <i>PLoS ONE</i> , 2014, 9, e87346.	2.5	10
121	Decreased Susceptibility to Tigecycline Mediated by a Mutation in <i>mlaA</i> in <i>Escherichia coli</i> Strains. <i>Antimicrobial Agents and Chemotherapy</i> , 2016, 60, 7530-7531.	3.2	10
122	Population Biology and Epidemiological Studies of <i>Acinetobacter baumannii</i> in the Era of Whole Genome Sequencing: Is the Oxford Scheme Still Appropriate?. <i>Frontiers in Microbiology</i> , 2020, 11, 775.	3.5	10
123	Effect of <i>ramR</i> loss-of-function insertion on tigecycline resistance in clinical isolates of carbapenem-resistant <i>Klebsiella pneumoniae</i> . <i>Journal of Global Antimicrobial Resistance</i> , 2020, 21, 410-413.	2.2	10
124	A Sequence Type 23 Hypervirulent <i>Klebsiella pneumoniae</i> Strain Presenting Carbapenem Resistance by Acquiring an <i>IncP1 bla</i> KPC-2 Plasmid. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021, 11, 641830.	3.9	10
125	<i>Acinetobacter baumannii</i> strains isolated from cerebrospinal fluid (CSF) and bloodstream analysed by cgMLST: the dominance of clonal complex CC92 in CSF infections. <i>International Journal of Antimicrobial Agents</i> , 2021, 58, 106404.	2.5	10
126	Genome Sequences of Two Multidrug-Resistant <i>Acinetobacter baumannii</i> Strains Isolated from a Patient before and after Treatment with Tigecycline. <i>Journal of Bacteriology</i> , 2012, 194, 6979-6980.	2.2	9

#	ARTICLE	IF	CITATIONS
127	Spread of a common bla _{NDM-1} -carrying plasmid among diverse <i>Acinetobacter</i> species. <i>Infection, Genetics and Evolution</i> , 2015, 32, 30-33.	2.3	9
128	Coexistence of <i>mcr-1</i> , <i>bla</i> KPC-2 and two copies of <i>fosA3</i> in a clinical <i>Escherichia coli</i> strain isolated from urine. <i>Infection, Genetics and Evolution</i> , 2018, 60, 77-79.	2.3	9
129	Characterization of a community-acquired methicillin-resistant sequence type 338 <i>Staphylococcus aureus</i> strain containing a staphylococcal cassette chromosome <i>mec</i> type VT. <i>International Journal of Infectious Diseases</i> , 2020, 90, 181-187.	3.3	9
130	Association of D-dimer elevation with inflammation and organ dysfunction in ICU patients with COVID-19 in Wuhan, China: a retrospective observational study. <i>Aging</i> , 2021, 13, 4794-4810.	3.1	9
131	The Role of <i>mprF</i> Mutations in Seesaw Effect of Daptomycin-Resistant Methicillin-Resistant <i>Staphylococcus aureus</i> Isolates. <i>Antimicrobial Agents and Chemotherapy</i> , 2022, 66, AAC0129521.	3.2	9
132	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
133	Epidemiological and antibiotic resistant study on extended-spectrum beta-lactamase-producing <i>Escherichia coli</i> and <i>Klebsiella pneumoniae</i> in Zhejiang Province. <i>Chinese Medical Journal</i> , 2002, 115, 1479-82.	2.3	9
134	fIDBAC: A Platform for Fast Bacterial Genome Identification and Typing. <i>Frontiers in Microbiology</i> , 2021, 12, 723577.	3.5	9
135	<i>Acinetobacter baumannii</i> Outer Membrane Protein A Induces Pulmonary Epithelial Barrier Dysfunction and Bacterial Translocation Through The TLR2/IQGAP1 Axis. <i>Frontiers in Immunology</i> , 0, 13, .	4.8	9
136	Evaluation of a quantitative serum <i>Aspergillus fumigatus</i> -specific IgM assay for diagnosis of chronic pulmonary aspergillosis. <i>Clinical Respiratory Journal</i> , 2018, 12, 2566-2572.	1.6	8
137	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
138	Household Transmission of Community-Associated Methicillin-Resistant <i>Staphylococcus Aureus</i> . <i>Frontiers in Public Health</i> , 2021, 9, 658638.	2.7	8
139	The Emergence of Novel Sequence Type Strains Reveals an Evolutionary Process of Intraspecies Clone Shifting in ICU-Spreading Carbapenem-Resistant <i>Klebsiella pneumoniae</i> . <i>Frontiers in Microbiology</i> , 2021, 12, 691406.	3.5	8
140	Multicenter Evaluation of Xpert Carba-R Assay for Detection and Identification of the Carbapenemase Genes in Rectal Swabs and Clinical Isolates. <i>Journal of Molecular Diagnostics</i> , 2021, 23, 111-119.	2.8	8
141	Emergence of Ceftazidime/Avibactam and Tigecycline Resistance in Carbapenem-Resistant <i>Klebsiella pneumoniae</i> Due to In-Host Microevolution. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021, 11, 757470.	3.9	8
142	Genomic epidemiology study of <i>Klebsiella pneumoniae</i> causing bloodstream infections in China. <i>Clinical and Translational Medicine</i> , 2021, 11, e624.	4.0	8
143	Epidemiological Characteristics and Clinical Manifestations of Hepatitis E in a Tertiary Hospital in China: A Retrospective Study. <i>Frontiers in Microbiology</i> , 2021, 12, 831968.	3.5	8
144	GR13-type plasmids in <i>Acinetobacter</i> potentiate the accumulation and horizontal transfer of diverse accessory genes. <i>Microbial Genomics</i> , 2022, 8, .	2.0	8

#	ARTICLE	IF	CITATIONS
145	Relocation of Tn2009 and characterization of an ABGR13-2 from re-sequenced genome sequence of <i>Acinetobacter baumannii</i> MDR-ZJ06. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 1153-1155.	3.0	7
146	Prevalence and characteristics of <i>pks</i> gene cluster harbouring <i>Klebsiella pneumoniae</i> from bloodstream infection in China. <i>Epidemiology and Infection</i> , 2020, 148, e69.	2.1	7
147	Bautype: Capsule and Lipopolysaccharide Serotype Prediction for <i>Acinetobacter baumannii</i> Genome. <i>Infectious Microbes & Diseases</i> , 2020, 2, 18-25.	1.3	7
148	Genome-Based Analysis of a Sequence Type 1049 Hypervirulent <i>Klebsiella pneumoniae</i> Causing Bacteremic Neck Abscess. <i>Frontiers in Microbiology</i> , 2020, 11, 617651.	3.5	7
149	Effect of pneumococcal conjugate vaccine availability on <i>Streptococcus pneumoniae</i> infections and genetic recombination in Zhejiang, China from 2009 to 2019. <i>Emerging Microbes and Infections</i> , 2022, 11, 606-615.	6.5	7
150	Molecular Mechanisms Driving the <i>In Vivo</i> Development of KPC-71-Mediated Resistance to Ceftazidime-Avibactam during Treatment of Carbapenem-Resistant <i>Klebsiella pneumoniae</i> Infections. <i>MSphere</i> , 2021, 6, e0085921.	2.9	7
151	Genomic and transcriptome analysis of triclosan response of a multidrug-resistant <i>Acinetobacter baumannii</i> strain, MDR-ZJ06. <i>Archives of Microbiology</i> , 2017, 199, 223-230.	2.2	6
152	Serum <i>Aspergillus fumigatus</i> -specific IgG antibody decreases after antifungal treatment in chronic pulmonary aspergillosis patients. <i>Clinical Respiratory Journal</i> , 2018, 12, 1772-1774.	1.6	6
153	Nonclassical Biofilms Induced by DNA Breaks in <i>Klebsiella pneumoniae</i> . <i>MSphere</i> , 2020, 5, .	2.9	6
154	Characterization of an ST5-SCCmec II-t311 methicillin-resistant <i>Staphylococcus aureus</i> strain with a widespread cfr-positive plasmid. <i>Journal of Infection and Chemotherapy</i> , 2020, 26, 699-705.	1.7	6
155	Genetic diversity of siderophores and hypermucoviscosity phenotype in <i>Klebsiella pneumoniae</i> . <i>Microbial Pathogenesis</i> , 2021, 158, 105014.	2.9	6
156	Prevalence and Characteristics of Ceftriaxone-Resistant <i>Salmonella</i> in Children's Hospital in Hangzhou, China. <i>Frontiers in Microbiology</i> , 2021, 12, 764787.	3.5	6
157	Anticolonization of Carbapenem-Resistant <i>Klebsiella pneumoniae</i> by <i>Lactobacillus plantarum</i> LP1812 Through Accumulated Acetic Acid in Mice Intestinal. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021, 11, 804253.	3.9	6
158	Epidemiological Characteristics of OXA-232-Producing Carbapenem-Resistant <i>Klebsiella pneumoniae</i> Strains Isolated during Nosocomial Clonal Spread Associated with Environmental Colonization. <i>Microbiology Spectrum</i> , 2022, 10, .	3.0	6
159	Comparing core-genome MLST with PFGE and MLST for cluster analysis of carbapenem-resistant <i>Acinetobacter baumannii</i> . <i>Journal of Global Antimicrobial Resistance</i> , 2022, 30, 148-151.	2.2	6
160	Characterization of a PVL-negative community-acquired methicillin-resistant <i>Staphylococcus aureus</i> strain of sequence type 88 in China. <i>International Journal of Medical Microbiology</i> , 2017, 307, 346-352.	3.6	5
161	Discovery of a Novel Hypervirulent <i>Acinetobacter baumannii</i> Strain in a Case of Community-Acquired Pneumonia. <i>Infection and Drug Resistance</i> , 2020, Volume 13, 1147-1153.	2.7	5
162	Phenotypic Variation and Carbapenem Resistance Potential in OXA-499-Producing <i>Acinetobacter pittii</i> . <i>Frontiers in Microbiology</i> , 2020, 11, 1134.	3.5	5

#	ARTICLE	IF	CITATIONS
163	Distribution of <i>erm</i> genes among MRSA isolates with resistance to clindamycin in a Chinese teaching hospital. <i>Infection, Genetics and Evolution</i> , 2021, 96, 105127.	2.3	5
164	Complete Genome Sequences of Bacteriophages Kaya, Guyu, Kopi, and TehO, Which Target Clinical Strains of <i>Pseudomonas aeruginosa</i> . <i>Microbiology Resource Announcements</i> , 2021, 10, e0104321.	0.6	5
165	Molecular Genetic Characteristics of Plasmid-Borne <i>mcr-9</i> in <i>Salmonella enterica</i> Serotype Typhimurium and Thompson in Zhejiang, China. <i>Frontiers in Microbiology</i> , 2022, 13, 852434.	3.5	5
166	Co-evolutionary adaptations of <i>Acinetobacter baumannii</i> and a clinical carbapenemase-encoding plasmid during carbapenem exposure. <i>Evolutionary Applications</i> , 2022, 15, 1045-1061.	3.1	5
167	Accuracy of in vitro susceptibility tests for carbapenemase-producing Gram-negative bacteria. <i>Journal of Medical Microbiology</i> , 2015, 64, 620-622.	1.8	4
168	Complete genome sequence of <i>Acinetobacter baumannii</i> A1296 (ST1469) with a small plasmid harbouring the <i>tet(39)</i> tetracycline resistance gene. <i>Journal of Global Antimicrobial Resistance</i> , 2017, 11, 105-107.	2.2	4
169	Revisiting the contribution of gene duplication of <i>bla</i> OXA-23 in carbapenem-resistant <i>Acinetobacter baumannii</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2018, 73, 250-252.	3.0	4
170	Spleen thickness can predict significant liver pathology in patients with chronic hepatitis B with persistently normal alanine aminotransferase or minimally raised alanine aminotransferase: a retrospective study. <i>Journal of International Medical Research</i> , 2019, 47, 122-132.	1.0	4
171	Clinical characteristic of 15 cases of cryptococcal meningitis treated with Ommaya reservoir. <i>Acta Neurologica Belgica</i> , 2020, 120, 1139-1145.	1.1	4
172	Emergence of a Clinical <i>Escherichia coli</i> Sequence Type 131 Strain Carrying a Chromosomal <i>bla</i> KPC-2 Gene. <i>Frontiers in Microbiology</i> , 2020, 11, 586764.	3.5	4
173	In vitro Effect of the Combination of Aztreonam and Amoxicillin/Clavulanic Acid Against Carbapenem-Resistant Gram-Negative Organisms Producing Metallo- β -Lactamase. <i>Infection and Drug Resistance</i> , 2021, Volume 14, 833-839.	2.7	4
174	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
175	China-United States Research Collaborations in Antimicrobial Resistance. <i>Clinical Infectious Diseases</i> , 2018, 67, S142-S145.	5.8	3
176	Molecular characteristics of PaLoc and acquired antimicrobial resistance in epidemic <i>Clostridioides difficile</i> isolates revealed by whole-genome sequencing. <i>Journal of Global Antimicrobial Resistance</i> , 2020, 23, 194-196.	2.2	3
177	New insights into the mechanisms of colistin resistance in <i>Klebsiella aerogenes</i> of clinical origin. <i>International Journal of Antimicrobial Agents</i> , 2020, 55, 105990.	2.5	3
178	A random forest model based on core genome allelic profiles of MRSA for penicillin plus potassium clavulanate susceptibility prediction. <i>Microbial Genomics</i> , 2021, 7, .	2.0	3
179	The novel fosfomycin resistance gene <i>fosY</i> is present on a genomic island in CC1 methicillin-resistant <i>Staphylococcus aureus</i> . <i>Emerging Microbes and Infections</i> , 2022, 11, 1166-1173.	6.5	3
180	Staphylococcal cassette chromosome <i>mec</i> amplification as a mechanism for ceftobiprole resistance in clinical methicillin-resistant <i>Staphylococcus aureus</i> isolates. <i>Clinical Microbiology and Infection</i> , 2022, 28, 1151.e1-1151.e7.	6.0	3

#	ARTICLE	IF	CITATIONS
181	Evaluation of the in vitro synergy of polymyxin B-based combinations against polymyxin B-resistant gram-negative bacilli. <i>Microbial Pathogenesis</i> , 2022, 166, 105517.	2.9	3
182	Whole-Genome Analysis of an Extensive Drug-Resistant <i>Acinetobacter Baumannii</i> ST195 Isolate from a Recipient After DCD Renal Transplantation in China. <i>Kidney and Blood Pressure Research</i> , 2017, 42, 1247-1257.	2.0	2
183	A new variant of mcr-1 identified from an extended-spectrum β -lactamase-producing <i>Escherichia coli</i> . <i>Journal of Global Antimicrobial Resistance</i> , 2019, 18, 26-27.	2.2	2
184	Whole-genome sequencing for detecting linezolid resistance in a patient with persistent methicillin-resistant <i>Staphylococcus aureus</i> infection during linezolid exposure. <i>International Journal of Antimicrobial Agents</i> , 2020, 55, 105819.	2.5	2
185	Rare and persistent <i>Rhodococcus equi</i> infection in a diffuse large B cell lymphoma patient: case report and review of the literature. <i>Journal of Thoracic Disease</i> , 2014, 6, E281-8.	1.4	2
186	Complete Genome Sequence of <i>Vibrio harveyi</i> Strain ATCC 33866. <i>Microbiology Resource Announcements</i> , 2022, 11, .	0.6	2
187	Draft genome sequence of ST203 vanA <i>Enterococcus faecium</i> strain XH877 from China. <i>Journal of Global Antimicrobial Resistance</i> , 2016, 7, 24-25.	2.2	1
188	Complete Genome Sequence of the Virulent <i>Klebsiella pneumoniae</i> Phage Geezett Infecting Multidrug-Resistant Clinical Strains. <i>Microbiology Resource Announcements</i> , 2021, 10, e0068521.	0.6	1
189	The Value of Neutrophil-To-Lymphocyte Ratio for Evaluating Blood Stream Infection Caused by Carbapenem-Resistant <i>Klebsiella pneumoniae</i> : A Retrospective Cohort Study. <i>Frontiers in Medicine</i> , 2022, 9, 832655.	2.6	1
190	Complete Genome Sequence of a Rare Pigment-Producing Strain of <i>Acinetobacter johnsonii</i> , Isolated from the Bile of a Patient in Hangzhou, China. <i>Microbiology Resource Announcements</i> , 2022, 11, e0002522.	0.6	0
191	Community Origins of Healthcare-Associated CC59 Methicillin-Resistant <i>Staphylococcus aureus</i> in China. <i>Journal of Infectious Diseases</i> , 0, , .	4.0	0