

# Zhiyong Zong

## List of Publications by Year in descending order

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168  
papers

5,365  
citations

101543

36  
h-index

128289

60  
g-index

175  
all docs

175  
docs citations

175  
times ranked

5659  
citing authors

#	ARTICLE	IF	CITATIONS
1	Clinical outcomes and risk factors for mortality from ventilator-associated events: A registry-based cohort study among 30,830 intensive care unit patients. <i>Infection Control and Hospital Epidemiology</i> , 2022, 43, 48-55.	1.8	14
2	Association between blood transfusion and ventilator-associated events: a nested case-control study. <i>Infection Control and Hospital Epidemiology</i> , 2022, 43, 597-602.	1.8	5
3	Fluid Balance and Ventilator-Associated Events Among Patients Admitted to ICUs in China: A Nested Case-Control Study*. <i>Critical Care Medicine</i> , 2022, 50, 307-316.	0.9	8
4	Impact of Allergic Rhinitis and Asthma on COVID-19 Infection, Hospitalization, and Mortality. <i>Journal of Allergy and Clinical Immunology: in Practice</i> , 2022, 10, 124-133.	3.8	53
5	<i>Klebsiella oxytoca</i> Complex: Update on Taxonomy, Antimicrobial Resistance, and Virulence. <i>Clinical Microbiology Reviews</i> , 2022, 35, e0000621.	13.6	48
6	An integrated IncFIB/IncFII plasmid confers hypervirulence and its fitness cost and stability. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2022, 41, 681-684.	2.9	6
7	Genome Analysis of <i>Klebsiella oxytoca</i> Complex for Antimicrobial Resistance and Virulence Genes. <i>Antimicrobial Agents and Chemotherapy</i> , 2022, 66, aac0218321.	3.2	8
8	Fine-Scale Reconstruction of the Evolution of FII-33 Multidrug Resistance Plasmids Enables High-Resolution Genomic Surveillance. <i>MSystems</i> , 2022, 7, e0083121.	3.8	9
9	Characterization of phage resistance and phages capable of intestinal decolonization of carbapenem-resistant <i>Klebsiella pneumoniae</i> in mice. <i>Communications Biology</i> , 2022, 5, 48.	4.4	32
10	<i>Aliidiomarina shirensis</i> as Possible Source of the Integron- and Plasmid-Mediated Fosfomycin Resistance Gene <i>fosC2</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2022, 66, aac0222721.	3.2	2
11	NDM-5-producing carbapenem-resistant <i>Klebsiella pneumoniae</i> of sequence type 789 emerged as a threat for neonates: a multicentre, genome-based study. <i>International Journal of Antimicrobial Agents</i> , 2022, 59, 106508.	2.5	5
12	Lytic Phages against ST11 K47 Carbapenem-Resistant <i>Klebsiella pneumoniae</i> and the Corresponding Phage Resistance Mechanisms. <i>MSphere</i> , 2022, 7, e0008022.	2.9	7
13	Strongyloidiasis in a Patient Diagnosed by Metagenomic Next-Generation Sequencing: A Case Report. <i>Frontiers in Medicine</i> , 2022, 9, 835252.	2.6	4
14	Conjugation of a Hybrid Plasmid Encoding Hypervirulence and Carbapenem Resistance in <i>Klebsiella pneumoniae</i> of Sequence Type 592. <i>Frontiers in Microbiology</i> , 2022, 13, 852596.	3.5	6
15	Potential Mobilization of <i>mcr-10</i> by an Integrative Mobile Element via Site-Specific Recombination in <i>Cronobacter sakazakii</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2021, 65, .	3.2	13
16	Precise Species Identification by Whole-Genome Sequencing of <i>Enterobacter</i> Bloodstream Infection, China. <i>Emerging Infectious Diseases</i> , 2021, 27, 161-169.	4.3	14
17	Molecular Basis of the Versatile Regulatory Mechanism of HtrA-Type Protease AlgW from <i>Pseudomonas aeruginosa</i> . <i>MBio</i> , 2021, 12, .	4.1	5
18	The epidemiology and clinical outcomes of ventilator-associated events among 20,769 mechanically ventilated patients at intensive care units: an observational study. <i>Critical Care</i> , 2021, 25, 44.	5.8	40

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19	KPC-12 with a L169M substitution in the Î© loop has reduced carbapenemase activity. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2021, 40, 1761-1766.	2.9	6
20	Carbapenem and Colistin Resistance in Enterobacter: Determinants and Clones. <i>Trends in Microbiology</i> , 2021, 29, 473-476.	7.7	29
21	Precise Species Identification for Acinetobacter: a Genome-Based Study with Description of Two Novel Acinetobacter Species. <i>MSystems</i> , 2021, 6, e0023721.	3.8	13
22	KPC-2-Producing Carbapenem-Resistant <i>Klebsiella pneumoniae</i> of the Uncommon ST29 Type Carrying OXA-926, a Novel Narrow-Spectrum OXA Î²-Lactamase. <i>Frontiers in Microbiology</i> , 2021, 12, 701513.	3.5	8
23	Genome-Based Taxonomy of <i>Brevundimonas</i> with Reporting <i>Brevundimonas huaxiensis</i> sp. nov.. <i>Microbiology Spectrum</i> , 2021, 9, e0011121.	3.0	15
24	Re-examining the association of AmpC variants with Enterobacter species in the context of updated taxonomy. <i>Antimicrobial Agents and Chemotherapy</i> , 2021, 65, e0159621.	3.2	6
25	Spread of Carbapenem-Resistant <i>Klebsiella pneumoniae</i> in an Intensive Care Unit: A Whole-Genome Sequence-Based Prospective Observational Study. <i>Microbiology Spectrum</i> , 2021, 9, e0005821.	3.0	12
26	Ubiquitous Conjugative Mega-Plasmids of Acinetobacter Species and Their Role in Horizontal Transfer of Multi-Drug Resistance. <i>Frontiers in Microbiology</i> , 2021, 12, 728644.	3.5	11
27	IS Î²1294 Reorganizes Plasmids in a Multidrug-Resistant <i>Escherichia coli</i> Strain. <i>Microbiology Spectrum</i> , 2021, 9, e0050321.	3.0	7
28	Cefoperazone-sulbactam and risk of coagulation disorders or bleeding: a retrospective cohort study. <i>Expert Opinion on Drug Safety</i> , 2020, 19, 339-347.	2.4	26
29	A Cluster of Colistin- and Carbapenem-Resistant <i>Klebsiella pneumoniae</i> Carrying bla <sub>NDM-1</sub> and mcr-8.2. <i>Journal of Infectious Diseases</i> , 2020, 221, S237-S242.	4.0	18
30	Selection of homemade mask materials for preventing transmission of COVID-19: A laboratory study. <i>PLoS ONE</i> , 2020, 15, e0240285.	2.5	30
31	Precise Species Identification and Taxonomy Update for the Genus <i>Kluyvera</i> With Reporting <i>Kluyvera sichuanensis</i> sp. nov.. <i>Frontiers in Microbiology</i> , 2020, 11, 579306.	3.5	13
32	Clinical characteristics and outcomes of patients with multidrug-resistant Gram-negative bacterial infections treated with ceftazidime/avibactam. <i>Journal of Global Antimicrobial Resistance</i> , 2020, 23, 404-407.	2.2	13
33	Precise Species Identification for <i>Enterobacter</i> : a Genome Sequence-Based Study with Reporting of Two Novel Species, <i>Enterobacter quasiroggenkampii</i> sp. nov. and <i>Enterobacter quasimori</i> sp. nov. <i>MSystems</i> , 2020, 5, .	3.8	95
34	Contamination of SARS-CoV-2 in patient surroundings and on personal protective equipment in a non-ICU isolation ward for COVID-19 patients with prolonged PCR positive status. <i>Antimicrobial Resistance and Infection Control</i> , 2020, 9, 167.	4.1	26
35	Risk factor for intestinal carriage of carbapenem-resistant <i>Acinetobacter baumannii</i> and the impact on subsequent infection among patients in an intensive care unit: an observational study. <i>BMJ Open</i> , 2020, 10, e035893.	1.9	9
36	Genome-based Taxonomy for Bacteria: A Recent Advance. <i>Trends in Microbiology</i> , 2020, 28, 871-874.	7.7	16

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37	Why did so few healthcare workers in China get COVID-19 infection. QJM - Monthly Journal of the Association of Physicians, 2020, 114, 225-226.	0.5	5
38	Struggle To Survive: the Choir of Target Alteration, Hydrolyzing Enzyme, and Plasmid Expression as a Novel Aztreonam-Avibactam Resistance Mechanism. MSystems, 2020, 5, .	3.8	22
39	Enhanced survival of ST-11 carbapenem-resistant <i>Klebsiella pneumoniae</i> in the intensive care unit. Infection Control and Hospital Epidemiology, 2020, 41, 740-742.	1.8	10
40	A Pandrug-Resistant <i>Providencia</i> Carrying Two blaIMP Carbapenemase-Encoding Genes Including blaIMP-69, a New blaIMP Variant, on a Newly Identified Worldwide-Distributed IncC Plasmid. Journal of Infectious Diseases, 2020, 221, S253-S256.	4.0	3
41	New evidence-based clinical practice guideline timely supports hospital infection control of coronavirus disease 2019. Precision Clinical Medicine, 2020, 3, 1-2.	3.3	6
42	Emergence of a Plasmid-Encoded Resistance-Nodulation-Division Efflux Pump Conferring Resistance to Multiple Drugs, Including Tigecycline, in <i>Klebsiella pneumoniae</i> . MBio, 2020, 11, .	4.1	153
43	Asymptomatic COVID-19 Patients Can Contaminate Their Surroundings: an Environment Sampling Study. MSphere, 2020, 5, .	2.9	81
44	Prolonged intermittent fever and massive splenomegaly in a miner working in the tropical jungle, China. PLoS Neglected Tropical Diseases, 2020, 14, e0008278.	3.0	2
45	Identification of novel mobile colistin resistance gene <i>mcr-10</i> . Emerging Microbes and Infections, 2020, 9, 508-516.	6.5	346
46	A precision medicine approach to managing 2019 novel coronavirus pneumonia. Precision Clinical Medicine, 2020, 3, 14-21.	3.3	34
47	Leprosy in a low-incidence setting. Wiener Klinische Wochenschrift, 2020, 132, 589-592.	1.9	2
48	Characterization of <i>Acinetobacter chengduensis</i> sp. nov., isolated from hospital sewage and capable of acquisition of carbapenem resistance genes. Systematic and Applied Microbiology, 2020, 43, 126092.	2.8	14
49	Infection Control in the Era of Antimicrobial Resistance in China: Progress, Challenges, and Opportunities. Clinical Infectious Diseases, 2020, 71, S372-S378.	5.8	12
50	<i>Enterobacter wuhouensis</i> sp. nov. and <i>Enterobacter quasihormaechei</i> sp. nov. recovered from human sputum. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 874-881.	1.7	22
51	Genome analysis-based reclassification of <i>Enterobacter tabaci</i> Duan et al. 2016 as a later heterotypic synonym of <i>Enterobacter mori</i> Zhu et al. 2011. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 1055-1058.	1.7	6
52	<i>Pseudomonas defluvii</i> sp. nov., isolated from hospital sewage. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 4199-4203.	1.7	10
53	An unhealing wound and subcutaneous nodules due to <i>Sporothrix globosa</i> after a cat bite. PLoS Neglected Tropical Diseases, 2020, 14, e0008859.	3.0	3
54	Handwashing Sink Contamination and Carbapenem-resistant <i>Klebsiella</i> Infection in the Intensive Care Unit: A Prospective Multicenter Study. Clinical Infectious Diseases, 2020, 71, S379-S385.	5.8	25

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55	Description of <i>Klebsiella spallanzanii</i> sp. nov. and of <i>Klebsiella pasteurii</i> sp. nov.. <i>Frontiers in Microbiology</i> , 2019, 10, 2360.	3.5	49
56	Key evolutionary events in the emergence of a globally disseminated, carbapenem resistant clone in the <i>Escherichia coli</i> ST410 lineage. <i>Communications Biology</i> , 2019, 2, 322.	4.4	36
57	The co-transfer of plasmid-borne colistin-resistant genes <i>mcr-1</i> and <i>mcr-3.5</i> , the carbapenemase gene <i>bla</i> NDM-5 and the 16S methylase gene <i>rmtB</i> from <i>Escherichia coli</i> . <i>Scientific Reports</i> , 2019, 9, 696.	3.3	48
58	NDM Metallo-β-Lactamases and Their Bacterial Producers in Health Care Settings. <i>Clinical Microbiology Reviews</i> , 2019, 32, .	13.6	406
59	Development and evaluation of the method for detecting metallo-carbapenemases among carbapenemase-producing Enterobacteriaceae. <i>Journal of Microbiological Methods</i> , 2019, 163, 105652.	1.6	8
60	Heterogeneous resistance to colistin in <i>Enterobacter cloacae</i> complex due to a new small transmembrane protein. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 2551-2558.	3.0	30
61	Nosocomial bloodstream infection and the emerging carbapenem-resistant pathogen <i>Ralstonia insidiosa</i> . <i>BMC Infectious Diseases</i> , 2019, 19, 334.	2.9	33
62	A Genomic, Evolutionary, and Mechanistic Study of MCR Action Suggests Functional Unification across the MCR Family of Colistin Resistance. <i>Advanced Science</i> , 2019, 6, 1900034.	11.2	29
63	Coexistence of three <i>bla</i> KPC-2 genes on an <i>IncF/IncR</i> plasmid in ST11 <i>Klebsiella pneumoniae</i> . <i>Journal of Global Antimicrobial Resistance</i> , 2019, 17, 90-93.	2.2	23
64	Reply to Kaier, Mutters, and Wolkewitz. <i>Clinical Infectious Diseases</i> , 2019, 69, 1082-1084.	5.8	0
65	<i>Acinetobacter chinensis</i> , a novel <i>Acinetobacter</i> species, carrying <i>bla</i> NDM-1, recovered from hospital sewage. <i>Journal of Microbiology</i> , 2019, 57, 350-355.	2.8	13
66	<i>Acinetobacter cumulans</i> sp. nov., isolated from hospital sewage and capable of acquisition of multiple antibiotic resistance genes. <i>Systematic and Applied Microbiology</i> , 2019, 42, 319-325.	2.8	24
67	The clinical impacts and risk factors for non-central line-associated bloodstream infection in 5046 intensive care unit patients: an observational study based on electronic medical records. <i>Critical Care</i> , 2019, 23, 52.	5.8	26
68	&lt;p&gt;Developing a Registry of Healthcare-Associated Infections at Intensive Care Units in West China: Study Rationale and Patient Characteristics&lt;/p&gt;. <i>Clinical Epidemiology</i> , 2019, Volume 11, 1035-1045.	3.0	14
69	Risk factors for ventilator-associated events: A prospective cohort study. <i>American Journal of Infection Control</i> , 2019, 47, 744-749.	2.3	22
70	Characterization of a strain representing a new <i>Enterobacter</i> species, <i>Enterobacter chengduensis</i> sp. nov.. <i>Antonie Van Leeuwenhoek</i> , 2019, 112, 491-500.	1.7	26
71	<i>Klebsiella huaxiensis</i> sp. nov., recovered from human urine. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 69, 333-336.	1.7	31
72	<i>Pseudomonas sichuanensis</i> sp. nov., isolated from hospital sewage. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 69, 517-522.	1.7	12

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73	<i>Enterobacter huaxiensis</i> sp. nov. and <i>Enterobacter chuandaensis</i> sp. nov., recovered from human blood. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 69, 708-714.	1.7	26
74	Genome analysis-based reclassification of <i>Lelliottia aquatilis</i> as a later heterotypic synonym of <i>Lelliottia jeotgali</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 69, 998-1000.	1.7	10
75	<i>Providencia huaxiensis</i> sp. nov., recovered from a human rectal swab. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 69, 2638-2643.	1.7	18
76	<i>Kosakonia quasisacchari</i> sp. nov. recovered from human wound secretion in China. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 69, 3155-3160.	1.7	11
77	<i>Pseudomonas huaxiensis</i> sp. nov., isolated from hospital sewage. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 69, 3281-3286.	1.7	9
78	Sequence Type 273 Carbapenem-Resistant <i>Klebsiella pneumoniae</i> Carrying <i>bla</i> NDM-1 and <i>bla</i> IMP-4. <i>Antimicrobial Agents and Chemotherapy</i> , 2018, 62, .	3.2	17
79	Infection prevention and control in outpatient settings in China—structure, resources, and basic practices. <i>American Journal of Infection Control</i> , 2018, 46, 802-807.	2.3	5
80	Coexistence of Two <i>bla</i> NDM-5 Genes on an IncF Plasmid as Revealed by Nanopore Sequencing. <i>Antimicrobial Agents and Chemotherapy</i> , 2018, 62, .	3.2	26
81	Cryptic transmission of ST405 <i>Escherichia coli</i> carrying <i>bla</i> NDM-4 in hospital. <i>Scientific Reports</i> , 2018, 8, 390.	3.3	25
82	Carbapenem-Resistant Hypervirulent <i>Klebsiella pneumoniae</i> of Sequence Type 36. <i>Antimicrobial Agents and Chemotherapy</i> , 2018, 62, .	3.2	66
83	Occurrence of <i>Enterobacter hormaechei</i> carrying <i>bla</i> NDM-1 and <i>bla</i> KPC-2 in China. <i>Diagnostic Microbiology and Infectious Disease</i> , 2018, 90, 139-142.	1.8	24
84	In-hospital Medical Costs of Infections Caused by Carbapenem-resistant <i>Klebsiella pneumoniae</i> . <i>Clinical Infectious Diseases</i> , 2018, 67, S225-S230.	5.8	45
85	Carbapenem-resistant Isolates of the <i>Klebsiella pneumoniae</i> Complex in Western China: The Common ST11 and the Surprising Hospital-specific Types. <i>Clinical Infectious Diseases</i> , 2018, 67, S263-S265.	5.8	36
86	Fitness cost of a <i>mcr-1</i> -carrying IncHI2 plasmid. <i>PLoS ONE</i> , 2018, 13, e0209706.	2.5	48
87	Antimicrobial stewardship for acute-care hospitals: An Asian perspective. <i>Infection Control and Hospital Epidemiology</i> , 2018, 39, 1237-1245.	1.8	31
88	<i>Klebsiella grimontii</i> , a New Species Acquired Carbapenem Resistance. <i>Frontiers in Microbiology</i> , 2018, 9, 2170.	3.5	12
89	The Occurrence of Colistin-Resistant Hypervirulent <i>Klebsiella pneumoniae</i> in China. <i>Frontiers in Microbiology</i> , 2018, 9, 2568.	3.5	36
90	Identification of <i>Mycobacterium chimaera</i> in heater-cooler units in China. <i>Scientific Reports</i> , 2018, 8, 7843.	3.3	10

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91	Methicillin-resistant <i>Staphylococcus aureus</i> nasal colonization and infection in an intensive care unit of a university hospital in China. <i>Journal of International Medical Research</i> , 2018, 46, 3698-3708.	1.0	11
92	Occurrence of colistin-resistant hypervirulent <i>Klebsiella variicola</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2018, 73, 3001-3004.	3.0	38
93	A P7 Phage-Like Plasmid Carrying <i>mcr-1</i> in an ST15 <i>Klebsiella pneumoniae</i> Clinical Isolate. <i>Frontiers in Microbiology</i> , 2018, 9, 11.	3.5	33
94	Two New Lytic Bacteriophages of the Myoviridae Family Against Carbapenem-Resistant <i>Acinetobacter baumannii</i> . <i>Frontiers in Microbiology</i> , 2018, 9, 850.	3.5	47
95	Complete genomic characterization of two <i>Escherichia coli</i> lineages responsible for a cluster of carbapenem-resistant infections in a Chinese hospital. <i>Journal of Antimicrobial Chemotherapy</i> , 2018, 73, 2340-2346.	3.0	44
96	<i>bla</i> NDM-21, a new variant of <i>bla</i> NDM in an <i>Escherichia coli</i> clinical isolate carrying <i>bla</i> CTX-M-55 and <i>rmtB</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2018, 73, 2336-2339.	3.0	46
97	<i>Acinetobacter wuhouensis</i> sp. nov., isolated from hospital sewage. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 3212-3216.	1.7	12
98	<i>Acinetobacter sichuanensis</i> sp. nov., recovered from hospital sewage in China. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 3897-3901.	1.7	8
99	<i>Enterobacter sichuanensis</i> sp. nov., recovered from human urine. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 3922-3927.	1.7	31
100	A large-scale survey on sharp injuries among hospital-based healthcare workers in China. <i>Scientific Reports</i> , 2017, 7, 42620.	3.3	16
101	Draft Genome Sequence of a Colistin-Resistant <i>Klebsiella pneumoniae</i> Clinical Strain Carrying the <i>bla</i> NDM-1 Carbapenemase Gene. <i>Genome Announcements</i> , 2017, 5, .	0.8	1
102	Whole genome sequences of three Clade 3 <i>Clostridium difficile</i> strains carrying binary toxin genes in China. <i>Scientific Reports</i> , 2017, 7, 43555.	3.3	21
103	Draft Genome Sequence of a Sequence Type 11 <i>Klebsiella pneumoniae</i> Clinical Strain Carrying a <i>bla</i> KPC-2 Carbapenemase Gene and an <i>rmtB</i> 16S rRNA Methylase Gene. <i>Genome Announcements</i> , 2017, 5, .	0.8	0
104	Draft Genome Sequence of a High-Level Colistin-Resistant Clinical Strain of the <i>Enterobacter cloacae</i> Complex. <i>Genome Announcements</i> , 2017, 5, .	0.8	1
105	IncP Plasmid Carrying Colistin Resistance Gene <i>mcr-1</i> in <i>Klebsiella pneumoniae</i> from Hospital Sewage. <i>Antimicrobial Agents and Chemotherapy</i> , 2017, 61, .	3.2	74
106	Surveillance of Dialysis Events: one-year experience at 33 outpatient hemodialysis centers in China. <i>Scientific Reports</i> , 2017, 7, 249.	3.3	7
107	New Variant of <i>mcr-3</i> in an Extensively Drug-Resistant <i>Escherichia coli</i> Clinical Isolate Carrying <i>mcr-1</i> and <i>bla</i> NDM-5. <i>Antimicrobial Agents and Chemotherapy</i> , 2017, 61, .	3.2	64
108	Impact of Infectious Disease Consultation on Clinical Management and Outcome of Patients with Bloodstream Infection: a Retrospective Cohort Study. <i>Scientific Reports</i> , 2017, 7, 12898.	3.3	17

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109	Antimicrobial susceptibility of <i>Clostridium difficile</i> isolates from ICU colonized patients revealed alert to ST-37 (RT 017) isolates. <i>Diagnostic Microbiology and Infectious Disease</i> , 2017, 89, 161-163.	1.8	6
110	Remarkable Diversity of <i>Escherichia coli</i> Carrying <i>mcr-1</i> from Hospital Sewage with the Identification of Two New <i>mcr-1</i> Variants. <i>Frontiers in Microbiology</i> , 2017, 8, 2094.	3.5	63
111	In Vitro Activity of Neomycin, Streptomycin, Paromomycin and Apramycin against Carbapenem-Resistant Enterobacteriaceae Clinical Strains. <i>Frontiers in Microbiology</i> , 2017, 8, 2275.	3.5	46
112	<i>Acinetobacter defluvii</i> sp. nov., recovered from hospital sewage. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 1709-1713.	1.7	37
113	The Impact of Infectious Disease Consultation on Clinical Management and Outcome of Patients With Bacteremia in China: A Retrospective Cohort Study. <i>Open Forum Infectious Diseases</i> , 2016, 3, .	0.9	0
114	Draft Genome Sequence of a <i>Pseudomonas</i> sp. Strain Carrying <i>bla</i> IMP-25 and <i>bla</i> VIM-2 Carbapenemase Genes from Hospital Sewage. <i>Genome Announcements</i> , 2016, 4, .	0.8	2
115	Genome sequence and virulence factors of a group G <i>Streptococcus dysgalactiae</i> subsp. <i>equisimilis</i> strain with a new element carrying <i>erm(B)</i> . <i>Scientific Reports</i> , 2016, 6, 20389.	3.3	10
116	<i>Kluyvera ascorbata</i> Strain from Hospital Sewage Carrying the <i>mcr-1</i> Colistin Resistance Gene. <i>Antimicrobial Agents and Chemotherapy</i> , 2016, 60, 7498-7501.	3.2	55
117	Colonization of toxigenic <i>Clostridium difficile</i> among ICU patients: a prospective study. <i>BMC Infectious Diseases</i> , 2016, 16, 397.	2.9	9
118	<i>Citrobacter freundii</i> carrying <i>bla</i> KPC-2 and <i>bla</i> NDM-1: characterization by whole genome sequencing. <i>Scientific Reports</i> , 2016, 6, 30670.	3.3	28
119	APSIC guide for prevention of Central Line Associated Bloodstream Infections (CLABSI). <i>Antimicrobial Resistance and Infection Control</i> , 2016, 5, 16.	4.1	88
120	Predictability of Phenotype in Relation to Common $\beta$ -Lactam Resistance Mechanisms in <i>Escherichia coli</i> and <i>Klebsiella pneumoniae</i> . <i>Journal of Clinical Microbiology</i> , 2016, 54, 1243-1250.	3.9	38
121	Increase in bacteraemia cases in the East Midlands region of the UK due to MDREscherichia coliST73: high levels of genomic and plasmid diversity in causative isolates. <i>Journal of Antimicrobial Chemotherapy</i> , 2016, 71, 339-343.	3.0	24
122	Characterization of <i>Acinetobacter johnsonii</i> isolate XBB1 carrying nine plasmids and encoding NDM-1, OXA-58 and PER-1 by genome sequencing. <i>Journal of Antimicrobial Chemotherapy</i> , 2016, 71, 71-75.	3.0	70
123	The Clinical Impact of Ventilator-Associated Events: A Prospective Multi-Center Surveillance Study. <i>Infection Control and Hospital Epidemiology</i> , 2015, 36, 1388-1395.	1.8	28
124	Biliary Tract Infection or Colonization with <i>Elizabethkingia meningoseptica</i> after Endoscopic Procedures Involving the Biliary Tract. <i>Internal Medicine</i> , 2015, 54, 11-15.	0.7	6
125	First identification of an IMI-1 carbapenemase-producing colistin-resistant <i>Enterobacter cloacae</i> in China. <i>Annals of Clinical Microbiology and Antimicrobials</i> , 2015, 14, 51.	3.8	26
126	Severe infections as the leading complication after the Lushan earthquake. <i>Intensive Care Medicine</i> , 2015, 41, 560-561.	8.2	2



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127	Different IncI1 plasmids from <i>Escherichia coli</i> carry ISEcp1-blaCTX-M-15 associated with different Tn2-derived elements. <i>Plasmid</i> , 2015, 80, 118-126.	1.4	39
128	<i>Escherichia coli</i> of sequence type 3835 carrying blaNDM-1, blaCTX-M-15, blaCMY-42 and blaSHV-12. <i>Scientific Reports</i> , 2015, 5, 12275.	3.3	33
129	Characterization of an <i>Enterobacter cloacae</i> Strain Producing both KPC and NDM Carbapenemases by Whole-Genome Sequencing. <i>Antimicrobial Agents and Chemotherapy</i> , 2015, 59, 6625-6628.	3.2	38
130	First Report of OXA-181-Producing <i>Escherichia coli</i> in China and Characterization of the Isolate Using Whole-Genome Sequencing. <i>Antimicrobial Agents and Chemotherapy</i> , 2015, 59, 5022-5025.	3.2	84
131	The Complex Genetic Context of blaPER-1 Flanked by Miniature Inverted-Repeat Transposable Elements in <i>Acinetobacter johnsonii</i> . <i>PLoS ONE</i> , 2014, 9, e90046.	2.5	28
132	ICU-Onset <i>Clostridium difficile</i> Infection in a University Hospital in China: A Prospective Cohort Study. <i>PLoS ONE</i> , 2014, 9, e111735.	2.5	26
133	Comparative genome analysis identifies few traits unique to the <i>Escherichia coli</i> ST131 H30Rx clade and extensive mosaicism at the capsule locus. <i>BMC Genomics</i> , 2014, 15, 830.	2.8	23
134	Impact of test methodology, media type and ion content on the susceptibility of <i>Acinetobacter</i> spp. to tigecycline. <i>Journal of Antimicrobial Chemotherapy</i> , 2014, 69, 1710-1711.	3.0	0
135	CTX-M-15-D-ST648 <i>Escherichia coli</i> from companion animals and horses: another pandemic clone combining multiresistance and extraintestinal virulence?. <i>Journal of Antimicrobial Chemotherapy</i> , 2014, 69, 1224-1230.	3.0	160
136	bla <sub>NDM-5</sub> Carried by an IncX3 Plasmid in <i>Escherichia coli</i> Sequence Type 167. <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 7548-7552.	3.2	85
137	Prediction of major antibiotic resistance in <i>Escherichia coli</i> and <i>Klebsiella pneumoniae</i> in Singapore, USA and China using a limited set of gene targets. <i>International Journal of Antimicrobial Agents</i> , 2014, 43, 563-565.	2.5	12
138	<i>Elizabethkingia meningoseptica</i> as an Unusual Pathogen Causing Healthcare-associated Bacteriuria. <i>Internal Medicine</i> , 2014, 53, 1877-1879.	0.7	8
139	Characterization of a complex context containing mecA but lacking genes encoding cassette chromosome recombinases in <i>Staphylococcus haemolyticus</i> . <i>BMC Microbiology</i> , 2013, 13, 64.	3.3	18
140	<i>Acinetobacter pittii</i> and <i>Acinetobacter nosocomialis</i> among clinical isolates of the <i>Acinetobacter calcoaceticus-baumannii</i> complex in Sichuan, China. <i>Diagnostic Microbiology and Infectious Disease</i> , 2013, 76, 392-395.	1.8	38
141	blaNDM-1-carrying <i>Acinetobacter johnsonii</i> detected in hospital sewage. <i>Journal of Antimicrobial Chemotherapy</i> , 2013, 68, 1007-1010.	3.0	83
142	Clonal diversity of <i>Acinetobacter baumannii</i> clinical isolates revealed by a snapshot study. <i>BMC Microbiology</i> , 2013, 13, 234.	3.3	45
143	Limited diversity in the gene pool allows prediction of third-generation cephalosporin and aminoglycoside resistance in <i>Escherichia coli</i> and <i>Klebsiella pneumoniae</i> . <i>International Journal of Antimicrobial Agents</i> , 2013, 42, 19-26.	2.5	32
144	The emergence of blaCTX-M-15-carrying <i>Escherichia coli</i> of ST131 and new sequence types in Western China. <i>Annals of Clinical Microbiology and Antimicrobials</i> , 2013, 12, 35.	3.8	15

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145	Complete Sequence of pJIE186-2, a Plasmid Carrying Multiple Virulence Factors from a Sequence Type 131 <i>Escherichia coli</i> O25 Strain. <i>Antimicrobial Agents and Chemotherapy</i> , 2013, 57, 597-600.	3.2	17
146	Kodamaea ohmerias an Emerging Pathogen in Mainland China: 3 Case Reports and Literature Review. <i>Laboratory Medicine</i> , 2013, 44, e1-e9.	1.2	5
147	Characterization of Extended-Spectrum $\beta$ -Lactamase Genes Found among <i>Escherichia coli</i> Isolates from Duck and Environmental Samples Obtained on a Duck Farm. <i>Applied and Environmental Microbiology</i> , 2012, 78, 3668-3673.	3.1	70
148	A Case of Loiasis in a Patient Returning to China Diagnosed by Nested PCR Using DNA Extracted From Tissue. <i>Journal of Travel Medicine</i> , 2012, 19, 314-316.	3.0	3
149	Misidentification of <i>Burkholderia pseudomallei</i> as <i>Burkholderia cepacia</i> by the VITEK 2 system. <i>Journal of Medical Microbiology</i> , 2012, 61, 1483-1484.	1.8	46
150	Enterobacteriaceae producing the KPC-2 carbapenemase from hospital sewage. <i>Diagnostic Microbiology and Infectious Disease</i> , 2012, 73, 204-206.	1.8	56
151	Discovery of blaOXA-199, a Chromosome-Based blaOXA-48-Like Variant, in <i>Shewanella xiamenensis</i> . <i>PLoS ONE</i> , 2012, 7, e48280.	2.5	41
152	The newly-recognized species <i>Staphylococcus massiliensis</i> is likely to be part of the human skin microflora. <i>Antonie Van Leeuwenhoek</i> , 2012, 101, 449-451.	1.7	6
153	blaCTX-M-carrying <i>Escherichia coli</i> of the O25b ST131 clonal group have emerged in China. <i>Diagnostic Microbiology and Infectious Disease</i> , 2011, 69, 228-231.	1.8	12
154	Tn2008 is a major vehicle carrying blaOXA-23 in <i>Acinetobacter baumannii</i> from China. <i>Diagnostic Microbiology and Infectious Disease</i> , 2011, 69, 218-222.	1.8	29
155	Diversity of SCCmec Elements in Methicillin-Resistant Coagulase-Negative Staphylococci Clinical Isolates. <i>PLoS ONE</i> , 2011, 6, e20191.	2.5	112
156	Recombination in IS <sub>26</sub> and Tn <sub>2</sub> in the Evolution of Multiresistance Regions Carrying bla <sub>CTX-M-15</sub> on Conjugative IncF Plasmids from <i>Escherichia coli</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2011, 55, 4971-4978.	3.2	105
157	Sequence type 38 <i>Escherichia coli</i> carrying bla <sub>CTX-M-14</sub> . <i>Journal of Medical Microbiology</i> , 2011, 60, 694-695.	1.8	7
158	bla <sub>CTX-M-65</sub> is carried by a Tn1722-like element on an IncN conjugative plasmid of ST131 <i>Escherichia coli</i> . <i>Journal of Medical Microbiology</i> , 2011, 60, 435-441.	1.8	18
159	Complete Sequence of pJIE143, a $\beta$ -Lactamase-Resistant Plasmid Carrying ISEcp1-bla <sub>CTX-M-15</sub> from an <i>Escherichia coli</i> ST131 Isolate. <i>Antimicrobial Agents and Chemotherapy</i> , 2011, 55, 5933-5935.	3.2	38
160	Nosocomial peripancreatic infection associated with <i>Shewanella xiamenensis</i> . <i>Journal of Medical Microbiology</i> , 2011, 60, 1387-1390.	1.8	37
161	Characterization of a New SCCmec Element in <i>Staphylococcus cohnii</i> . <i>PLoS ONE</i> , 2010, 5, e14016.	2.5	30
162	<i>Escherichia coli</i> carrying the bla <sub>CTX-M-15</sub> gene of ST648. <i>Journal of Medical Microbiology</i> , 2010, 59, 1536-1537.	1.8	20

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163	IS <i>Ecp1</i> -Mediated Transposition and Homologous Recombination Can Explain the Context of <i>bla</i> <sub>CTX-M-62</sub> Linked to <i>qnrB2</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2010, 54, 3039-3042.	3.2	38
164	A <i>bla</i> <sub>VEB-1</sub> Variant, <i>bla</i> <sub>VEB-6</sub> , Associated with Repeated Elements in a Complex Genetic Structure. <i>Antimicrobial Agents and Chemotherapy</i> , 2009, 53, 1693-1697.	3.2	23
165	An outbreak of carbapenem-resistant <i>Acinetobacter baumannii</i> producing OXA-23 carbapenemase in western China. <i>International Journal of Antimicrobial Agents</i> , 2008, 31, 50-54.	2.5	57
166	Dominance of <i>bla</i> <sub>CTX-M</sub> within an Australian Extended-Spectrum $\beta$ -Lactamase Gene Pool. <i>Antimicrobial Agents and Chemotherapy</i> , 2008, 52, 4198-4202.	3.2	87
167	RmtC 16S rRNA Methyltransferase in Australia. <i>Antimicrobial Agents and Chemotherapy</i> , 2008, 52, 794-795.	3.2	16
168	Should post-trial provision of beneficial experimental interventions be mandatory in developing countries?. <i>Journal of Medical Ethics</i> , 2008, 34, 188-192.	1.8	34