Minggui Wang

List of Publications by Year
in descending order

113 papers	4,975 citations	117625 34 h-index	106344 65 g-index
115 all docs	115 docs citations	115 times ranked	5238 citing authors

#	Article	IF	Citations
1	Structures of Class I and Class II Transcription Complexes Reveal the Molecular Basis of RamAâ€Dependent Transcription Activation. Advanced Science, 2022, 9, e2103669.	11.2	13
2	Clinical outcomes and bacterial characteristics of carbapenem-resistant Klebsiella pneumoniae complex among patients from different global regions (CRACKLE-2): a prospective, multicentre, cohort study. Lancet Infectious Diseases, The, 2022, 22, 401-412.	9.1	122
3	Transmission barrier of the blaKPC plasmid mediated by type I restriction-modification systems in Escherichia coli. Journal of Antimicrobial Chemotherapy, 2022, 77, 952-956.	3.0	7
4	A Clinical Practice Guideline for the Emergency Management of Anaphylaxis (2020). Frontiers in Pharmacology, 2022, 13, 845689.	3.5	7
5	Treatment and economic burden of mucormycosis in China: Case report review and burden estimation. Journal of Clinical Pharmacy and Therapeutics, 2022, 47, 905-914.	1.5	5
6	Combined PK/PD Index May Be a More Appropriate PK/PD Index for Cefoperazone/Sulbactam against Acinetobacter baumannii in Patients with Hospital-Acquired Pneumonia. Antibiotics, 2022, 11, 703.	3.7	4
7	Serogroup Y Clonal Complex 23 Meningococcus in China Acquiring Penicillin Resistance from Commensal Neisseria lactamica Species. Antimicrobial Agents and Chemotherapy, 2022, 66, .	3.2	1
8	Antibiotic Exposure during the Preceding Six Months Is Related to Intestinal ESBL-Producing Enterobacteriaceae Carriage in the Elderly. Antibiotics, 2022, 11, 953.	3.7	2
9	Carbapenemase-Encoding Gene Copy Number Estimator (CCNE): a Tool for Carbapenemase Gene Copy Number Estimation. Microbiology Spectrum, 2022, 10, .	3.0	4
10	Determination of norvancomycin epidemiological cut-off values (ECOFFs) for Staphylococcus aureus, Staphylococcus epidermidis, Staphylococcus haemolyticus and Staphylococcus hominis. Journal of Antimicrobial Chemotherapy, 2021, 76, 152-159.	3.0	8
11	RamA upregulates multidrug resistance efflux pumps AcrAB and OqxAB in Klebsiella pneumoniae. International Journal of Antimicrobial Agents, 2021, 57, 106251.	2.5	19
12	An IncP-2 plasmid sublineage associated with dissemination of bla_{IMP-45} among carbapenem-resistant Pseudomonas aeruginosa. Emerging Microbes and Infections, 2021, 10, 442-449.	6.5	22
13	Development of a Fast Raman-Assisted Antibiotic Susceptibility Test (FRAST) for the Antibiotic Resistance Analysis of Clinical Urine and Blood Samples. Analytical Chemistry, 2021, 93, 5098-5106.	6.5	45
14	Clinical and molecular characteristics of Chryseobacterium indologenes isolates at a teaching hospital in Shanghai, China. Annals of Translational Medicine, 2021, 9, 668-668.	1.7	12
15	Molecular Characteristics of Escherichia coli Causing Bloodstream Infections During 2010â€“2015 in Tertiary Hospital, Shanghai, China. Infection and Drug Resistance, 2021, Volume 14, 2079-2086.	2.7	1
16	Establishment of epidemiological cut-off values for cefoselis, a new fourth-generation cephalosporin, against Escherichia coli, Klebsiella pneumoniae, Enterobacter cloacae, Proteus mirabilis and Pseudomonas aeruginosa. Journal of Antimicrobial Chemotherapy, 2021, 76, 2593-2599.	3.0	4
17	Identification of qnrE3 and qnrE4 , New Transferable Quinolone Resistance qnrE Family Genes Originating from Enterobacter mori and Enterobacter asburiae, Respectively. Antimicrobial Agents and Chemotherapy, 2021, 65, e0045621.	3.2	6
18	Characterization of the novel plasmid-encoded MBL gene blaAFM-1, integrated into a blaIMP-45-bearing transposon Tn6485e in a carbapenem-resistant Pseudomonas aeruginosa clinical isolate. Journal of Antimicrobial Chemotherapy, 2021, 77, 83-88.	3.0	10

#	Article	IF	Citations
19	Efflux Pump AcrAB Confers Decreased Susceptibility to Piperacillinâ€“Tazobactam and Ceftolozaneâ€“Tazobactam in Tigecycline-Non-Susceptible Klebsiella pneumoniae . Infection and Drug Resistance, 2020, Volume 13, 4309-4319.	2.7	10
20	In Vivo Evolution of CTX-M-215, a Novel Narrow-Spectrum Î²-Lactamase in an Escherichia coli Clinical Isolate Conferring Resistance to Mecillinam. Antimicrobial Agents and Chemotherapy, 2020, 64, .	3.2	4
21	Acinetobacter baumannii Sequence Types Harboring Genes Encoding Aminoglycoside Modifying Enzymes and 16SrRNA Methylase; a Multicenter Study from Pakistan . Infection and Drug Resistance, 2020, Volume 13, 2855-2862.	2.7	15
22	The Predominance of Strain Replacement Among Enterobacteriaceae Pairs With Emerging Carbapenem Resistance During Hospitalization. Journal of Infectious Diseases, 2020, 221, S215-S219.	4.0	3
23	The Colonization of Carbapenem-Resistant Klebsiella pneumoniae: Epidemiology, Resistance Mechanisms, and Risk Factors in Patients Admitted to Intensive Care Units in China. Journal of Infectious Diseases, 2020, 221, S206-S214.	4.0	83
24	CHINET efforts to control antimicrobial resistance in China. Journal of Global Antimicrobial Resistance, 2020, 21, 76-77.	2.2	19
25	Emergence of a Plasmid-Encoded Resistance-Nodulation-Division Efflux Pump Conferring Resistance to Multiple Drugs, Including Tigecycline, in Klebsiella pneumoniae. MBio, 2020, 11, .	4.1	153
26	Molecular and clinical epidemiology of carbapenem-resistant Enterobacterales in the USA (CRACKLE-2): a prospective cohort study. Lancet Infectious Diseases, The, 2020, 20, 731-741.	9.1	174
27	Dissemination of blaOXA-23-harbouring carbapenem-resistant Acinetobacter baumannii clones in Pakistan. Journal of Global Antimicrobial Resistance, 2020, 21, 357-362.	2.2	25
28	Molecular Characteristics and Antimicrobial Susceptibility Profiles of Elizabethkingia Clinical Isolates in Shanghai, China . Infection and Drug Resistance, 2020, Volume 13, 247-256.	2.7	17
29	Vancomycin Heteroresistance in vanM-type Enterococcus faecium. Microbial Drug Resistance, 2020, 26, 776-782.	2.0	8
30	Plasmids and genes contributing to high-level quinolone resistance in Escherichia coli. International Journal of Antimicrobial Agents, 2020, 56, 105987.	2.5	10
31	Evidence-based Guideline for Therapeutic Drug Monitoring of Vancomycin: 2020 Update by the Division of Therapeutic Drug Monitoring, Chinese Pharmacological Society. Clinical Infectious Diseases, 2020, 71, S363-S371.	5.8	109
32	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. Clinical Infectious Diseases, 2020, 71, S337-S362.	5.8	9
33	toxin B-induced colonic inflammation is mediated by the FOXO3/PPM1B pathway in fetal human colon epithelial cells. American Journal of Translational Research (discontinued), 2020, 12, 6204-6219.	0.0	0
34	Evaluation of anidulafungin in the treatment of intra-abdominal candidiasis: a pooled analysis of patient-level data from 5 prospective studies. European Journal of Clinical Microbiology and Infectious Diseases, 2019, 38, 1849-1856.	2.9	12
35	Outbreak Of Klebsiella pneumoniae Carbapenemase-Producing Klebsiella aerogenes Strains In A Tertiary Hospital In China . Infection and Drug Resistance, 2019, Volume 12, 3283-3290.	2.7	11
36	Efflux pumps AcrAB and OqxAB contribute to nitrofurantoin resistance in an uropathogenic Klebsiella pneumoniae isolate. International Journal of Antimicrobial Agents, 2019, 54, 223-227.	2.5	27

#	Article	IF	Citations
37	2037. A Novel Strategy of Antimicrobial Stewardship in Shanghai: Preliminary Practice with Integration of Three Surveillance Networks. Open Forum Infectious Diseases, 2019, 6, S685-S685.	0.9	0
38	Porin Deficiency in Carbapenem-ResistantEnterobacter aerogenesStrains. Microbial Drug Resistance, 2018, 24, 1277-1283.	2.0	27
39	New Subclass B1 Metallo-Î²-Lactamase Gene from a Clinical PathogenicMyroides odoratusStrain. Microbial Drug Resistance, 2018, 24, 909-914.	2.0	3
40	Antimicrobial Resistance in China: Challenges and Actions. Clinical Infectious Diseases, 2018, 67, S127-S127.	5.8	9
41	Current Status and Trends of Antibacterial Resistance in China. Clinical Infectious Diseases, 2018, 67, S128-S134.	5.8	205
42	A Case-Control Study: Clinical Characteristics of Nosocomial Bloodstream Infections Versus Non-bloodstream Infections ofAcinetobacterspp.. Clinical Infectious Diseases, 2018, 67, S189-S195.	5.8	4
43	Chinaâ€“United States Research Collaborations in Antimicrobial Resistance. Clinical Infectious Diseases, 2018, 67, S142-S145.	5.8	3
44	Comparison of empirical therapy with cefoperazone/sulbactam or a carbapenem for bloodstream infections due to ESBL-producing Enterobacteriaceae. Journal of Antimicrobial Chemotherapy, 2018, 73, 3176-3180.	3.0	18
45	Evolution of Sequence Type 4821 Clonal Complex Meningococcal Strains in China from Prequinolone to Quinolone Era, 1972â€“2013. Emerging Infectious Diseases, 2018, 24, 683-690.	4.3	11
46	Molecular characteristics of Clostridium difficile strains from patients with a first recurrence more than 8Âweeks after the primary infection. Journal of Microbiology, Immunology and Infection, 2017, 50, 532-536.	3.1	3
47	IncX2 and IncX1-X2 Hybrid Plasmids Coexisting in a FosA6-Producing Escherichia coli Strain. Antimicrobial Agents and Chemotherapy, 2017, 61, .	3.2	14
48	High ceftazidime hydrolysis activity and porin OmpK35 deficiency contribute to the decreased susceptibility to ceftazidime/avibactam in KPC-producing Klebsiella pneumoniae. Journal of Antimicrobial Chemotherapy, 2017, 72, 1930-1936.	3.0	64
49	Characterization of a blaNDM-1-harboring plasmid from a Salmonella enterica clinical isolate in China. Molecular Medicine Reports, 2017, 16, 1087-1092.	2.4	13
50	In vivo development of tigecycline resistance in Klebsiella pneumoniae owing to deletion of the ramR ribosomal binding site. International Journal of Antimicrobial Agents, 2017, 50, 523-528.	2.5	20
51	CTX-M-190, a Novel Î²-Lactamase Resistant to Tazobactam and Sulbactam, Identified in an Escherichia coli Clinical Isolate. Antimicrobial Agents and Chemotherapy, 2017, 61, .	3.2	21
52	High-Level Fosfomycin Resistance in Vancomycin-Resistant Enterococcus faecium. Emerging Infectious Diseases, 2017, 23, 1902-1904.	4.3	23
53	Mutations of the Transporter Proteins GlpT and UhpT Confer Fosfomycin Resistance in Staphylococcus aureus. Frontiers in Microbiology, 2017, 8, 914.	3.5	35
54	Macrolide-resistant Mycoplasma pneumoniae prevalence and clinical aspects in adult patients with community-acquired pneumonia in China: a prospective multicenter surveillance study. Journal of Thoracic Disease, 2017, 9, 3774-3781.	1.4	21

#	Article	IF	Citations
55	Clinical and Genomic Analysis of Liver Abscess-Causing Klebsiella pneumoniae Identifies New Liver Abscess-Associated Virulence Genes. Frontiers in Cellular and Infection Microbiology, 2016, 6, 165.	3.9	82
56	In vivo Acquisition of Carbapenemase Gene blaKPC-2 in Multiple Species of Enterobacteriaceae through Horizontal Transfer of Insertion Sequence or Plasmid. Frontiers in Microbiology, 2016, 7, 1651.	3.5	11
57	Emergence of tigecycline- and carbapenem-nonsusceptible Klebsiella pneumoniae ST11 clone in patients without exposure to tigecycline. Journal of Microbiology, Immunology and Infection, 2016, 49, 962-968.	3.1	5
58	Characterization of a Novel IncHI2 Plasmid Carrying Tandem Copies of bla _{CTX-M-2} in a fosA6 -Harboring Escherichia coli Sequence Type 410 Strain. Antimicrobial Agents and Chemotherapy, 2016, 60, 6742-6747.	3.2	12
59	Achieving global targets for antimicrobial resistance. Science, 2016, 353, 874-875.	12.6	233
60	Glutathione-S-transferase FosA6 ofKlebsiella pneumoniaeorigin conferring fosfomycin resistance in ESBL-producingEscherichia coli. Journal of Antimicrobial Chemotherapy, 2016, 71, 2460-2465.	3.0	49
61	Laboratory diagnosis, clinical management and infection control of the infections caused by extensively drug-resistant Gram-negative bacilli: a Chinese consensus statement. Clinical Microbiology and Infection, 2016, 22, S15-S25.	6.0	85
62	Characterization of Fosfomycin Resistance Gene, fosB, in Methicillin-Resistant Staphylococcus aureus Isolates. PLoS ONE, 2016, 11, e0154829.	2.5	42
63	Shifts in the Antibiotic Susceptibility, Serogroups, and Clonal Complexes of Neisseria meningitidis in Shanghai, China: A Time Trend Analysis of the Pre-Quinolone and Quinolone Eras. PLoS Medicine, 2015, 12, e1001838.	8.4	35
64	Four Carbapenem-Resistant Gram-Negative Species Carrying Distinct Carbapenemases in a Single Patient. Journal of Clinical Microbiology, 2015, 53, 1031-1033.	3.9	11
65	Characterization of fosA5 , a new plasmid-mediated fosfomycin resistance gene in Escherichia coli. Letters in Applied Microbiology, 2015, 60, 259-264.	2.2	47
66	Mapping the resistance-associated mobilome of a carbapenem-resistantKlebsiella pneumoniaestrain reveals insights into factors shaping these regions and facilitates generation of a â€˜resistance-disarmedâ€™ model organism. Journal of Antimicrobial Chemotherapy, 2015, 70, 2770-2774.	3.0	55
67	High Prevalence of vanM in Vancomycin-Resistant Enterococcus faecium Isolates from Shanghai, China. Antimicrobial Agents and Chemotherapy, 2015, 59, 7795-7798.	3.2	34
68	Clonal dissemination of extensively drug-resistant Acinetobacter baumannii producing an OXA-23 Î²-lactamase at a teaching hospital in Shanghai, China. Journal of Microbiology, Immunology and Infection, 2015, 48, 101-108.	3.1	22
69	Prevalence of Fosfomycin Resistance and Mutations in murA, glpT, and uhpT in Methicillin-Resistant Staphylococcus aureus Strains Isolated from Blood and Cerebrospinal Fluid Samples. Frontiers in Microbiology, 2015, 6, 1544.	3.5	39
70	Prevalence of the fosfomycin-resistance determinant, fosB3, in Enterococcus faecium clinical isolates from China. Journal of Medical Microbiology, 2014, 63, 1484-1489.	1.8	26
71	Draft Genome Sequence of Linezolid-Resistant Enterococcus faecalis Clinical Isolate HS0914. Genome Announcements, 2014, 2, .	0.8	0
72	Selection and characterisation of Staphylococcus aureus mutants with reduced susceptibility to the investigational oxazolidinone MRX-I. International Journal of Antimicrobial Agents, 2014, 43, 418-422.	2.5	19

#	Article	IF	Citations
73	Genetic diversity of fluoroquinolone-nonsusceptible Streptococcus pneumoniae clinical isolates and the first identification of serotype 20B in China. European Journal of Clinical Microbiology and Infectious Diseases, 2014, 33, 465-470.	2.9	3
74	Association Between Single-Nucleotide Polymorphism in CISH Gene and Susceptibility to Tuberculosis in Chinese Han Population. Cell Biochemistry and Biophysics, 2014, 68, 529-534.	1.8	14
75	Mechanisms of Tigecycline Resistance among Klebsiella pneumoniae Clinical Isolates. Antimicrobial Agents and Chemotherapy, 2014, 58, 6982-6985.	3.2	71
76	High-level tetracycline resistance mediated by efflux pumps Tet(A) and Tet(A)-1 with two start codons. Journal of Medical Microbiology, 2014, 63, 1454-1459.	1.8	21
77	Type II and type IV topoisomerase mutations in clinical isolates of Morganella morganii harbouring the qnrD gene. Annals of Clinical Microbiology and Antimicrobials, 2014, 13, 34.	3.8	12
78	Detection of Mycoplasma pneumoniae P1 subtype variations by denaturing gradient gel electrophoresis. Diagnostic Microbiology and Infectious Disease, 2014, 78, 24-28.	1.8	21
79	Treatment of extensively drug-resistant Gram-negative infections in critically ill patients: Outcome of a consensus meeting at the 13th Asia-Pacific Congress of Clinical Microbiology and Infection, October 2012. Journal of Global Antimicrobial Resistance, 2013, 1, 117-122.	2.2	5
80	Sulbactam-based therapy for Acinetobacter baumannii infection: a systematic review and meta-analysis. Brazilian Journal of Infectious Diseases, 2013, 17, 389-394.	0.6	53
81	New Delhi Metallo-Î²-Lactamase-1 in Carbapenem-ResistantSalmonellaStrain, China. Emerging Infectious Diseases, 2013, 19, 2049-2051.	4.3	30
82	Prevalence of fusB in Staphylococcus aureus clinical isolates. Journal of Medical Microbiology, 2013, 62, 1199-1203.	1.8	10
83	The Fosfomycin Resistance Gene fosB3 Is Located on a Transferable, Extrachromosomal Circular Intermediate in Clinical Enterococcus faecium Isolates. PLoS ONE, 2013, 8, e78106.	2.5	31
84	Comparison of Adhesin Genes and Antimicrobial Susceptibilities between Uropathogenic and Intestinal Commensal Escherichia coli Strains. PLoS ONE, 2013, 8, e61169.	2.5	70
85	Co-production of SFO-1 and DHA-1 Â-lactamases and 16S rRNA methylase ArmA in clinical isolates of Klebsiella pneumoniae. Journal of Antimicrobial Chemotherapy, 2012, 67, 2361-2366.	3.0	16
86	Prevalence of the oqxAB gene complex in Klebsiella pneumoniae and Escherichia coli clinical isolates. Journal of Antimicrobial Chemotherapy, 2012, 67, 1655-1659.	3.0	86
87	Establishment of norvancomycin fluorescence polarization immunoassay for therapeutic drug monitoring. Journal of Antibiotics, 2012, 65, 35-39.	2.0	14
88	Multiclonal Origin of Macrolide-Resistant Mycoplasma pneumoniae Isolates as Determined by Multilocus Variable-Number Tandem-Repeat Analysis. Journal of Clinical Microbiology, 2012, 50, 2793-2795.	3.9	26
89	Comparison of the genetic structures surrounding qnrA1 in Korean Enterobacter cloacae and Chinese Escherichia coli Strains isolated in the early 2000s: Evidence for qnrA mobilization via Inc HI2 type plasmid. Journal of Microbiology, 2012, 50, 166-169.	2.8	1
90	Coexistence of armA and genes encoding aminoglycoside-modifying enzymes in Acinetobacter baumannii. African Journal of Microbiology Research, 2012, 6, .	0.4	1

#	Article	IF	Citations
91	Susceptibility of Extended-Spectrum-Î²-Lactamase-Producing Enterobacteriaceae According to the New CLSI Breakpoints. Journal of Clinical Microbiology, 2011, 49, 3127-3131.	3.9	54
92	Liquid chromatography/tandem mass spectrometry assay for the simultaneous determination of cefoperazone and sulbactam in plasma and its application to a pharmacokinetic study. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2010, 878, 3119-3124.	2.3	27
93	Distribution of 16S rRNA methylases among different species of Gram-negative bacilli with high-level resistance to aminoglycosides. European Journal of Clinical Microbiology and Infectious Diseases, 2010, 29, 1349-1353.	2.9	83
94	A mutational analysis and molecular dynamics simulation of quinolone resistance proteins QnrA1 and QnrC from Proteus mirabilis. BMC Structural Biology, 2010, 10, 33.	2.3	14
95	Prevalence of plasmid-mediated quinolone-resistance determinants in Shigella flexneri isolates from Anhui Province, China. Journal of Antibiotics, 2010, 63, 187-189.	2.0	21
96	First Report of the Multidrug Resistance Gene cfr and the Phenicol Resistance Gene fexA in a Bacillus Strain from Swine Feces. Antimicrobial Agents and Chemotherapy, 2010, 54, 3953-3955.	3.2	71
97	vanM , a New Glycopeptide Resistance Gene Cluster Found in Enterococcus faecium. Antimicrobial Agents and Chemotherapy, 2010, 54, 4643-4647.	3.2	169
98	Decreased quinolone susceptibility in high percentage of Enterobacter cloacae clinical isolates caused only by Qnr determinants. Diagnostic Microbiology and Infectious Disease, 2010, 67, 110-113.	1.8	26
99	Characterization of macrolide resistance in Mycoplasma pneumoniae isolated from children in Shanghai, China. Diagnostic Microbiology and Infectious Disease, 2010, 67, 355-358.	1.8	74
100	The prevalence of plasmid-mediated quinolone resistance determinants among clinical isolates of ESBL or AmpC-producingEscherichia colifrom Chinese pediatric patients. Microbiology and Immunology, 2010, 54, 123-128.	1.4	21
101	New Plasmid-Mediated Quinolone Resistance Gene, qnrC , Found in a Clinical Isolate of Proteus mirabilis. Antimicrobial Agents and Chemotherapy, 2009, 53, 1892-1897.	3.2	279
102	Antimicrobial Susceptibility of Mycoplasma pneumoniae Isolates and Molecular Analysis of Macrolide-Resistant Strains from Shanghai, China. Antimicrobial Agents and Chemotherapy, 2009, 53, 2160-2162.	3.2	118
103	The prevalence of plasmid-mediated quinolone resistance determinants among clinical isolates of ESBL or AmpC-producing Escherichia coli from Chinese pediatric patients. Microbiology and Immunology, 2009, 54, 123-128.	1.4	0
104	Clostridium difficile infections in a Shanghai hospital: antimicrobial resistance, toxin profiles and ribotypes. International Journal of Antimicrobial Agents, 2009, 33, 339-342.	2.5	84
105	High Prevalence of Plasmid-Mediated Quinolone Resistance Determinants qnr , aac(6 â€² Tj ETQq1 1 0.784314 rgBT /Ov from Companion and Food-Producing Animals. Antimicrobial Agents and Chemotherapy, 2009, 53, 519-524.	3.2	188
106	Coexistence of qnrB4 and qnrS1 in a clinical strain of Klebsiella pneumoniae. Acta Pharmacologica Sinica, 2008, 29, 320-324.	6.1	36
107	qnr Gene Nomenclature. Antimicrobial Agents and Chemotherapy, 2008, 52, 2297-2299.	3.2	192
108	Molecular and Clinical Characteristics ofClostridium difficileInfection in a University Hospital in Shanghai, China. Clinical Infectious Diseases, 2008, 47, 1606-1608.	5.8	32

#	Article	IF	Citations
109	Prevalence and Expression of the Plasmid-Mediated Quinolone Resistance Determinant qnrA1. Antimicrobial Agents and Chemotherapy, 2007, 51, 4105-4110.	3.2	32
110	Activities of Newer Quinolones against Escherichia coli and Klebsiella pneumoniae Containing the Plasmid-Mediated Quinolone Resistance Determinant qnr. Antimicrobial Agents and Chemotherapy, 2004, 48, 1400-1401.	3.2	36
111	Emerging Plasmid-Mediated Quinolone Resistance Associated with the qnr Gene in Klebsiella pneumoniae Clinical Isolates in the United States. Antimicrobial Agents and Chemotherapy, 2004, 48, 1295-1299.	3.2	219
112	Plasmid-Mediated Quinolone Resistance in Clinical Isolates of Escherichia coli from Shanghai, China. Antimicrobial Agents and Chemotherapy, 2003, 47, 2242-2248.	3.2	423
113	Prevalence and phenotypes of erythromycin-resistant Streptococcus pneumoniae in Shanghai, China. Diagnostic Microbiology and Infectious Disease, 2001, 39, 187-189.	1.8	9