Ruichao Li

List of Publications by Year
in descending order

126	3,006	218381	233125
papers	citations	26 h-index	45 g-index

128	128	128	2147
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Genetic characterization of mcr-1 -bearing plasmids to depict molecular mechanisms underlying dissemination of the colistin resistance determinant. Journal of Antimicrobial Chemotherapy, 2017, 72, 393-401.	1.3	198
2	Prevalence and characterization of Salmonella species isolated from pigs, ducks and chickens in Sichuan Province, China. International Journal of Food Microbiology, 2013, 163, 14-18.	2.1	162
3	Efficient generation of complete sequences of MDR-encoding plasmids by rapid assembly of MinION barcoding sequencing data. GigaScience, 2018, 7, 1-9.	3.3	140
4	Antibiotic adjuvants: an alternative approach to overcome multi-drug resistant Gram-negative bacteria. Critical Reviews in Microbiology, 2019, 45, 301-314.	2.7	118
5	Metformin Restores Tetracyclines Susceptibility against Multidrug Resistant Bacteria. Advanced Science, 2020, 7, 1902227.	5.6	104
6	Gut microbiome alterations in high-fat-diet-fed mice are associated with antibiotic tolerance. Nature Microbiology, 2021, 6, 874-884.	5.9	88
7	Deciphering the Structural Diversity and Classification of the Mobile Tigecycline Resistance Gene tet (X)-Bearing Plasmidome among Bacteria. MSystems, 2020, 5, .	1.7	85
8	Complete genetic analysis of plasmids carryingmcr-1and other resistance genes in anEscherichia coliisolate of animal origin. Journal of Antimicrobial Chemotherapy, 2017, 72, dkw509.	1.3	73
9	Drug repurposing for next-generation combination therapies against multidrug-resistant bacteria. Theranostics, 2021, 11, 4910-4928.	4.6	70
10	Melatonin overcomes MCR-mediated colistin resistance in Gram-negative pathogens. Theranostics, 2020, 10, 10697-10711.	4.6	60
11	Genome analysis of clinical multilocus sequence Type 11 Klebsiella pneumoniae from China. Microbial Genomics, 2018, 4, .	1.0	52
12	CARB-17 Family of Î²-Lactamases Mediates Intrinsic Resistance to Penicillins in Vibrio parahaemolyticus. Antimicrobial Agents and Chemotherapy, 2015, 59, 3593-3595.	1.4	49
13	Emergence of IncX3 Plasmid-Harboring blaNDMâ€“5 Dominated by Escherichia coli ST48 in a Goose Farm in Jiangsu, China. Frontiers in Microbiology, 2019, 10, 2002.	1.5	48
14	Recombination of plasmids in a carbapenem-resistant NDM-5-producing clinical Escherichia coli isolate. Journal of Antimicrobial Chemotherapy, 2018, 73, 1230-1234.	1.3	47
15	Exploring tet(X)-bearing tigecycline-resistant bacteria of swine farming environments. Science of the Total Environment, 2020, 733, 139306.	3.9	47
16	Genetic basis of chromosomally-encoded mcr-1 gene. International Journal of Antimicrobial Agents, 2018, 51, 578-585.	1.1	46
17	Emergence of a novel conjugative hybrid virulence multidrug-resistant plasmid in extensively drug-resistant Klebsiella pneumoniae ST15. International Journal of Antimicrobial Agents, 2020, 55, 105952.	1.1	45
18	Anti-HIV agent azidothymidine decreases Tet(X)-mediated bacterial resistance to tigecycline in Escherichia coli. Communications Biology, 2020, 3, 162.	2.0	41

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19	Prevalence and Molecular Characterization of mcr-1 -Positive Salmonella Strains Recovered from Clinical Specimens in China. Antimicrobial Agents and Chemotherapy, 2017, 61, .	1.4	40
20	Emergence of plasmid-mediated tigecycline resistance tet(X4) gene in Escherichia coli isolated from poultry, food and the environment in South Asia. Science of the Total Environment, 2021, 787, 147613.	3.9	40
21	Characterization of TMexCD3-TOprJ3, an RND-Type Efflux System Conferring Resistance to Tigecycline in Proteus mirabilis, and Its Associated Integrative Conjugative Element. Antimicrobial Agents and Chemotherapy, 2021, 65, e0271220.	1.4	38
22	First Detection of AmpC Î²-Lactamase bla _{CMY-2} on a Conjugative IncA/C Plasmid in a Vibrio parahaemolyticus Isolate of Food Origin. Antimicrobial Agents and Chemotherapy, 2015, 59, 4106-4111.	1.4	37
23	Emergence of a hybrid plasmid derived from IncN1-F33:Aâ^':Bâ^' and mcr-1-bearing plasmids mediated by IS26. Journal of Antimicrobial Chemotherapy, 2019, 74, 3184-3189.	1.3	37
24	Unique Class 1 Integron and Multiple Resistance Genes Co-located on IncHI2 Plasmid Is Associated with the Emerging Multidrug Resistance of Salmonella Indiana Isolated from Chicken in China. Foodborne Pathogens and Disease, 2013, 10, 581-588.	0.8	36
25	IncHI2 Plasmids Are the Key Vectors Responsible for oqxAB Transmission among Salmonella Species. Antimicrobial Agents and Chemotherapy, 2016, 60, 6911-6915.	1.4	35
26	Characterization of a porcine Proteus cibarius strain co-harbouring tet(X6) and cfr. Journal of Antimicrobial Chemotherapy, 2020, 75, 1652-1654.	1.3	34
27	Novel IS26-mediated hybrid plasmid harbouring tet(X4) in Escherichia coli. Journal of Global Antimicrobial Resistance, 2020, 21, 162-168.	0.9	31
28	Characterization of an IncA/C Multidrug Resistance Plasmid in Vibrio alginolyticus. Antimicrobial Agents and Chemotherapy, 2016, 60, 3232-3235.	1.4	30
29	Comprehensive Genomic Investigation of Tigecycline Resistance Gene tet (X4)-Bearing Strains Expanding among Different Settings. Microbiology Spectrum, 2021, 9, e0163321.	1.2	27
30	Emergence of mcr-8.2-bearing Klebsiella quasipneumoniae of animal origin. Journal of Antimicrobial Chemotherapy, 2019, 74, 2814-2817.	1.3	26
31	Acetaminophen promotes horizontal transfer of plasmid-borne multiple antibiotic resistance genes. Science of the Total Environment, 2021, 782, 146916.	3.9	26
32	Plasmids Shape the Current Prevalence of tmexCD1-toprJ1 among Klebsiella pneumoniae in Food Production Chains. MSystems, 2021, 6, e0070221.	1.7	26
33	Molecules that Inhibit Bacterial Resistance Enzymes. Molecules, 2019, 24, 43.	1.7	25
34	Cooccurrence of Two tet (X) Variants in an Empedobacter brevis Strain of Shrimp Origin. Antimicrobial Agents and Chemotherapy, 2019, 63, .	1.4	24
35	Characterisation of a cointegrate plasmid harbouring blaNDM-1 in a clinical Salmonella Lomita strain. International Journal of Antimicrobial Agents, 2020, 55, 105817.	1.1	23
36	Co-existence of tet(X4) and mcr-1 in two porcine Escherichia coli isolates. Journal of Antimicrobial Chemotherapy, 2020, 75, 764-766.	1.3	23

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37	Combined Linkage Mapping and BSA to Identify QTL and Candidate Genes for Plant Height and the Number of Nodes on the Main Stem in Soybean. International Journal of Molecular Sciences, 2020, 21, 42.	1.8	23
38	Emergence of a multidrug resistance efflux pump with carbapenem resistance gene blaVIM-2 in a Pseudomonas putida megaplasmid of migratory bird origin. Journal of Antimicrobial Chemotherapy, 2021, 76, 1455-1458.	1.3	23
39	Widespread Prevalence of Plasmid-Mediated Colistin Resistance Gene mcr-1 in Escherichia coli from PÃ¨re David's Deer in China. MSphere, 2020, 5, .	1.3	23
40	IncFII Conjugative Plasmid-Mediated Transmission of bla _{NDM-1} Elements among Animal-Borne Escherichia coli Strains. Antimicrobial Agents and Chemotherapy, 2017, 61, .	1.4	22
41	Bacterial metabolism-inspired molecules to modulate antibiotic efficacy. Journal of Antimicrobial Chemotherapy, 2019, 74, 3409-3417.	1.3	22
42	Identification of a novel hybrid plasmid coproducing MCR-1 and MCR-3 variant from an Escherichia coli strain. Journal of Antimicrobial Chemotherapy, 2019, 74, 1517-1520.	1.3	21
43	Emergence of Carbapenem- and Tigecycline-Resistant Proteus cibarius of Animal Origin. Frontiers in Microbiology, 2020, 11, 1940.	1.5	21
44	Genomic Epidemiology Insights on NDM-Producing Pathogens Revealed the Pivotal Role of Plasmids on bla _{NDM} Transmission. Microbiology Spectrum, 2022, 10, e0215621.	1.2	21
45	Molecular Characterization of Escherichia coli Isolates Carrying mcr-1 , fosA3 , and Extended-Spectrum-Î²-Lactamase Genes from Food Samples in China. Antimicrobial Agents and Chemotherapy, 2017, 61, .	1.4	20
46	A Novel PCR-Based Approach for Accurate Identification of Vibrio parahaemolyticus. Frontiers in Microbiology, 2016, 7, 44.	1.5	19
47	Characterization of a Multidrug-Resistant Porcine Klebsiella pneumoniae Sequence Type 11 Strain Coharboring bla _{KPC-2} and fosA3 on Two Novel Hybrid Plasmids. MSphere, 2019, 4, .	1.3	19
48	Comprehensive Genomic Investigation of Coevolution of mcr genes in Escherichia coli Strains via Nanopore Sequencing. Global Challenges, 2021, 5, 2000014.	1.8	19
49	Characterization of novel ISAba1-bounded tet(X15)-bearing composite transposon Tn6866 in Acinetobacter variabilis. Journal of Antimicrobial Chemotherapy, 2021, 76, 2481-2483.	1.3	19
50	Identification and Characterization of Conjugative Plasmids That Encode Ciprofloxacin Resistance in Salmonella. Antimicrobial Agents and Chemotherapy, 2018, 62, .	1.4	18
51	Coexistence of tet (X4), mcr-1 , and bla _{NDM-5} in ST6775 Escherichia coli Isolates of Animal Origin in China. Microbiology Spectrum, 2022, 10, e0019622.	1.2	18
52	Distinct mechanisms of acquisition of mcr-1 â€“bearing plasmid by Salmonella strains recovered from animals and food samples. Scientific Reports, 2017, 7, 13199.	1.6	17
53	Widespread prevalence and molecular epidemiology of tet(X4) and mcr-1 harboring Escherichia coli isolated from chickens in Pakistan. Science of the Total Environment, 2022, 806, 150689.	3.9	17
54	Characterization of Acinetobacter indicus co-harbouring tet(X3) and blaNDM-1 of dairy cow origin. Journal of Antimicrobial Chemotherapy, 2020, 75, 2693-2696.	1.3	17

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55	Phenotypic and genomic analysis reveals Riemerella anatipestifer as the potential reservoir of tet(X) variants. Journal of Antimicrobial Chemotherapy, 2022, 77, 374-380.	1.3	17
56	Development of a Novel Hexa-plex PCR Method for Identification and Serotyping of Salmonella Species. Foodborne Pathogens and Disease, 2014, 11, 75-77.	0.8	16
57	Antagonizing Vancomycin Resistance in Enterococcus by Surface Localized Antimicrobial Display-Derived Peptides. ACS Infectious Diseases, 2020, 6, 761-767.	1.8	15
58	Co-occurrence of plasmid-mediated resistance genes tet(X4) and blaNDM-5 in a multidrug-resistant Escherichia coli isolate recovered from chicken in China. Journal of Global Antimicrobial Resistance, 2021, 24, 415-417.	0.9	15
59	Identification of Two Plasmids Coharboring Carbapenemase Genes and tmexCD1-toprJ1 in Clinical Klebsiella pneumoniae ST2667. Antimicrobial Agents and Chemotherapy, 2021, 65, .	1.4	15
60	Subinhibitory Concentration of Colistin Promotes the Conjugation Frequencies of Mcr-1- and bla _{NDM-5} -Positive Plasmids. Microbiology Spectrum, 2022, 10, e0216021.	1.2	15
61	Identification and Characterization of IncA/C Conjugative, bla _{NDM-1} -Bearing Plasmid in Vibrio alginolyticus of Food Origin. Antimicrobial Agents and Chemotherapy, 2018, 62, .	1.4	14
62	Characterization of the stability and dynamics of Tn6330 in an Escherichia coli strain by nanopore long reads. Journal of Antimicrobial Chemotherapy, 2019, 74, 1807-1811.	1.3	14
63	Polymorphism Existence of Mobile Tigecycline Resistance Gene tet (X4) in Escherichia coli. Antimicrobial Agents and Chemotherapy, 2020, 64, .	1.4	14
64	Characterization of Three Porcine Acinetobacter towneri Strains Co-Harboring tet(X3) and blaOXA-58. Frontiers in Cellular and Infection Microbiology, 2020, 10, 586507.	1.8	14
65	Reorganization of mcr-1-bearing large MDR plasmids resolved by nanopore sequencing. Journal of Antimicrobial Chemotherapy, 2020, 75, 1645-1647.	1.3	14
66	Rapid detection and characterization of tet(X4)-positive Escherichia coli strains with nanopore sequencing. Journal of Antimicrobial Chemotherapy, 2020, 75, 1068-1070.	1.3	14
67	Distribution and genomic characterization of tigecycline-resistant tet(X4)-positive Escherichia coli of swine farm origin. Microbial Genomics, 2021, 7, .	1.0	14
68	Complete Nucleotide Sequence of a Conjugative Plasmid Carrying bla _{PER-1}. Antimicrobial Agents and Chemotherapy, 2015, 59, 3582-3584.	1.4	13
69	Resolution of dynamic MDR structures among the plasmidome of Salmonella using MinION single-molecule, long-read sequencing. Journal of Antimicrobial Chemotherapy, 2018, 73, 2691-2695.	1.3	13
70	Recombination of NDM-5-producing plasmids mediated by IS26 among Escherichia coli. International Journal of Antimicrobial Agents, 2020, 55, 105815.	1.1	13
71	Genomic Epidemiology of ST34 Monophasic Salmonella enterica Serovar Typhimurium from Clinical Patients from 2008 to 2017 in Henan, China. Engineering, 2022, 15, 34-44.	3.2	13
72	Genetic Characterization of blaCTXâ€“Mâ€“55 -Bearing Plasmids Harbored by Food-Borne Cephalosporin-Resistant Vibrio parahaemolyticus Strains in China. Frontiers in Microbiology, 2019, 10, 1338.	1.5	12

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73	Genetic characterization of an MDR/virulence genomic element carrying two T6SS gene clusters in a clinical Klebsiella pneumoniae isolate of swine origin. Journal of Antimicrobial Chemotherapy, 2019, 74, 1539-1544.	1.3	12
74	SVM+KF Target Tracking Strategy Using the Signal Strength in Wireless Sensor Networks. Sensors, 2020, 20, 3832.	2.1	12
75	Emerging Opportunity and Destiny of mcr-1 - and tet (X4)-Coharboring Plasmids in Escherichia coli. Microbiology Spectrum, 2021, 9, e0152021.	1.2	12
76	Evolution and comparative genomics of pAQU-like conjugative plasmids in Vibrio species. Journal of Antimicrobial Chemotherapy, 2017, 72, 2503-2506.	1.3	11
77	Chromosome-mediated mcr-1 in Escherichia coli strain L73 from a goose. International Journal of Antimicrobial Agents, 2019, 54, 99-101.	1.1	11
78	Emergence of Plasmid-Mediated Resistance Genes tet (X) and mcr-1 in Escherichia coli Clinical Isolates from Pakistan. MSphere, 2021, 6, e0069521.	1.3	11
79	Structural Diversity, Fitness Cost, and Stability of a BlaNDM-1-Bearing Cointegrate Plasmid in Klebsiella pneumoniae and Escherichia coli. Microorganisms, 2021, 9, 2435.	1.6	11
80	Deciphering the Epidemiological Characteristics and Molecular Features of blaKPCâ€“2- or blaNDMâ€“1-Positive Klebsiella pneumoniae Isolates in a Newly Established Hospital. Frontiers in Microbiology, 2021, 12, 741093.	1.5	11
81	Selection of target mutation in rat gastrointestinal tract E. coli by minute dosage of enrofloxacin. Frontiers in Microbiology, 2014, 5, 468.	1.5	10
82	Genetic Characterization of a bla VEB-2 -Carrying Plasmid in Vibrio parahaemolyticus. Antimicrobial Agents and Chemotherapy, 2016, 60, 6965-6968.	1.4	10
83	Loss of mcr Genes Mediated by Plasmid Elimination and IS Apl1. Antimicrobial Agents and Chemotherapy, 2019, 63, .	1.4	9
84	Contribution of biofilm formation genetic locus, pgaABCD, to antibiotic resistance development in gut microbiome. Gut Microbes, 2020, 12, 1842992.	4.3	9
85	Genomic features of a high-risk mcr-1.1-positive Escherichia coli ST10 isolated from cattle farm environment. Environmental Science and Pollution Research, 2021, 28, 54147-54152.	2.7	9
86	Characterization of a blaNDM-1-Bearing IncHI5-Like Plasmid From Klebsiella pneumoniae of Infant Origin. Frontiers in Cellular and Infection Microbiology, 2021, 11, 738053.	1.8	9
87	Paclitaxel and its derivative facilitate the transmission of plasmid-mediated antibiotic resistance genes through conjugative transfer. Science of the Total Environment, 2022, 810, 152245.	3.9	9
88	Extensive spread of tet(X4) in multidrug-resistant Escherichia coli of animal origin in western China. Veterinary Microbiology, 2022, 269, 109420.	0.8	9
89	Complete Sequence of a F33:A-:B- Conjugative Plasmid Carrying the oqxAB, fosA3, and blaCTX-M-55 Elements from a Foodborne Escherichia coli Strain. Frontiers in Microbiology, 2016, 7, 1729.	1.5	8
90	First detection of a bla CTX-M-15 -carrying plasmid in Vibrio alginolyticus. Journal of Global Antimicrobial Resistance, 2018, 13, 206-208.	0.9	8

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91	Emergence of a Novel tet (L) Variant in Campylobacter spp. of Chicken Origin in China. Antimicrobial Agents and Chemotherapy, 2020, 65, .	1.4	8
92	Histone-Like Nucleoid Structuring Protein Modulates the Fitness of tet(X4)-Bearing IncX1 Plasmids in Gram-Negative Bacteria. Frontiers in Microbiology, 2021, 12, 763288.	1.5	8
93	QitanTech Nanopore Long-Read Sequencing Enables Rapid Resolution of Complete Genomes of Multi-Drug Resistant Pathogens. Frontiers in Microbiology, 2022, 13, 778659.	1.5	8
94	Characterization of Carbapenem-Resistant Enterobacteriaceae Cultured From Retail Meat Products, Patients, and Porcine Excrement in China. Frontiers in Microbiology, 2021, 12, 743468.	1.5	8
95	Conjugative transfer of mcr-1-bearing plasmid from Salmonella to Escherichia coli in vitro on chicken meat and in mouse gut. Food Research International, 2022, 157, 111263.	2.9	8
96	Genetic Characterization of Broad-Host-Range IncQ Plasmids Harboring bla _{VEB-18} in Vibrio Species. Antimicrobial Agents and Chemotherapy, 2017, 61, .	1.4	7
97	Identification of a Novel Plasmid-Mediated Carbapenemase-Encoding Gene, bla _{VMB-2} , in Vibrio diabolicus. Antimicrobial Agents and Chemotherapy, 2021, 65, e0020621.	1.4	7
98	Genomic characterization of conjugative plasmids carrying the mcr-1 gene in foodborne and clinical strains of Salmonella and Escherichia coli. Food Control, 2021, 125, 108032.	2.8	7
99	Emergence of blaNDM-9-bearing tigecycline-resistant Klebsiella aerogenes of chicken origin. Journal of Global Antimicrobial Resistance, 2021, 26, 66-68.	0.9	7
100	IS 1294 Reorganizes Plasmids in a Multidrug-Resistant Escherichia coli Strain. Microbiology Spectrum, 2021, 9, e0050321.	1.2	7
101	Emergence and Characterization of Tigecycline Resistance Gene tet (X4) in ST609 Escherichia coli Isolates from Wastewater in Turkey. Microbiology Spectrum, 2022, 10, .	1.2	7
102	Complete Genetic Analysis of Plasmids Carried by Two Nonclonal bla _{NDM-5} - and mcr-1 -Bearing Escherichia coli Strains: Insight into Plasmid Transmission among Foodborne Bacteria. Microbiology Spectrum, 2021, 9, e0021721.	1.2	6
103	Prevalence, toxin-typing and antimicrobial susceptibility of Clostridium perfringens in sheep with different feeding modes from Gansu and Qinghai provinces, China. Anaerobe, 2022, 73, 102516.	1.0	6
104	A PM2.5 concentration estimation method based on multi-feature combination of image patches. Environmental Research, 2022, 211, 113051.	3.7	6
105	Characterization of blaNDM-positive Enterobacteriaceae reveals the clonal dissemination of Enterobacter hormaechei coharboring blaNDM and tet(X4) along the pork production chain. International Journal of Food Microbiology, 2022, 372, 109692.	2.1	6
106	Coexistence of tmexCD-toprJ , bla _{NDM-1} , and bla _{IMP-4} in One Plasmid Carried by Clinical Klebsiella spp.. Microbiology Spectrum, 2022, 10, .	1.2	6
107	Characterization of a Novel mcr-8.2-Bearing Plasmid in ST395 Klebsiella pneumoniae of Chicken Origin . Infection and Drug Resistance, 2020, Volume 13, 1781-1784.	1.1	5
108	Investigation of tigecycline resistant Escherichia coli from raw meat reveals potential transmission among food-producing animals. Food Control, 2021, 121, 107633.	2.8	5

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109	PK/PD integration and pharmacodynamic cutoff of cefquinome against cow mastitis due to Escherichia coli. Journal of Veterinary Pharmacology and Therapeutics, 2021, , .	0.6	5
110	Emergence of the cfr Gene in Vibrio diabolicus of Seafood Origin. Antimicrobial Agents and Chemotherapy, 2022, 66, AAC0181921.	1.4	5
111	Occurrence and Molecular Characterization of Abundant tet(X) Variants Among Diverse Bacterial Species of Chicken Origin in Jiangsu, China. Frontiers in Microbiology, 2021, 12, 751006.	1.5	5
112	Rapid and Accurate Antibiotic Susceptibility Determination of tet (X)-Positive E. coli Using RNA Biomarkers. Microbiology Spectrum, 2021, 9, e0064821.	1.2	4
113	Formation, Transmission, and Dynamic Evolution of a Multidrug-Resistant Chromosomally Integrated Plasmid in Salmonella Spp.. Frontiers in Microbiology, 2022, 13, 846954.	1.5	4
114	Sodium dehydroacetate confers broad antibiotic tolerance by remodeling bacterial metabolism. Journal of Hazardous Materials, 2022, 432, 128645.	6.5	4
115	Small clone dissemination of tmexCD1-toprJ1â€“carrying Klebsiella pneumoniae isolates in a chicken farm. Journal of Global Antimicrobial Resistance, 2022, 29, 105-112.	0.9	4
116	Distribution, antimicrobial resistance and genomic characterization of Salmonella along the pork production chain in Jiangsu, China. LWT - Food Science and Technology, 2022, 163, 113516.	2.5	4
117	In vivo Pharmacokinetic and Pharmacodynamic (PK/PD) Modeling and Establishment of the PK/PD Cutoff of Florfenicol Against Pasteurella multocida in Ducks. Frontiers in Microbiology, 2020, 11, 616685.	1.5	3
118	Cunning plasmid fusion mediates antibiotic resistance genes represented by ESBLs encoding genes transfer in foodborne Salmonella. International Journal of Food Microbiology, 2021, 355, 109336.	2.1	3
119	Metformin Reverses tmexCD1-toprJ1- and tet(A)-Mediated High-Level Tigecycline Resistance in K. pneumoniae. Antibiotics, 2022, 11, 162.	1.5	3
120	Characterisation of a chromosomally-encoded extended-spectrum Î²-lactamase gene blaPER-3 in Aeromonas caviae of chicken origin. International Journal of Antimicrobial Agents, 2016, 47, 103-105.	1.1	2
121	Molecular characterization of two novel NDM-1-producing atypical enteroaggregative Escherichia coli isolates from patients. Plasmid, 2021, 115, 102568.	0.4	2
122	Whole-genome sequencing of strains of Vibrio spp. from China reveals different genetic contexts of blaCTX-M-14 among diverse lineages. Journal of Antimicrobial Chemotherapy, 2021, 76, 950-956.	1.3	2
123	Editorial: Investigating Antimicrobial Resistance With Single-Molecule Sequencing Technologies: Opportunities and Challenges. Frontiers in Microbiology, 2022, 13, .	1.5	2
124	Emergence of Mobilized Colistin Resistance Gene mcr-8.2 in Multidrug-Resistant Enterobacter cloacae Isolated from a Patient in China. Microbiology Spectrum, 0, , .	1.2	1
125	Rapid resolution of multi-drug resistance bacterial genome harbouring mcr-1 and blaCMY-2 using MinION sequencing platform. International Journal of Antimicrobial Agents, 2018, 52, 303-304.	1.1	0
126	Plasmid-mediated ciprofloxacin, carbapenem and colistin resistance of a foodborne Escherichia coli isolate. Food Control, 2022, 137, 108937.	2.8	0