

# 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  | 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        |
| 2  | 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        |
| 3  | 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        |
| 4  | 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         |
| 5  | <i>Alcaligenes faecalis</i> metallo- $\beta$ -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        |
| 6  | 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       |
| 7  | 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         |
| 8  | 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) <i>Tj ETQq0 0 0 rgBT /Overlook 10 Tf 5</i>                                  |     |           |
| 9  | 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        |
| 10 | 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         |
| 11 | 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        |
| 12 | 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         |
| 13 | 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        |
| 14 | 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        |
| 15 | 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         |
| 16 | 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         |
| 17 | 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         |
| 18 | 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         |

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|----|--|------|-----------|
| 19 | Complete Genome Sequence of <i>Vibrio harveyi</i> Strain ATCC 33866. <i>Microbiology Resource Announcements</i> , 2022, 11, .  | 0.6  | 2         |
| 20 | 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         |
| 21 | 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         |
| 22 | 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         |
| 23 | 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         |
| 24 | 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       |
| 25 | 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         |
| 26 | Acquisition of a genomic resistance island (AbGRI5) 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        |
| 27 | 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        |
| 28 | 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        |
| 29 | 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        |
| 30 | 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         |
| 31 | In vitro Effect of the Combination of Aztreonam and Amoxicillin/Clavulanic Acid Against Carbapenem-Resistant Gram-Negative Organisms Producing Metallo- $\beta$ -Lactamase. <i>Infection and Drug Resistance</i> , 2021, Volume 14, 833-839.   | 2.7  | 4         |
| 32 | Emergence of carbapenem-resistant <i>Klebsiella pneumoniae</i> harbouring bla OXA-48-like genes in China. <i>Journal of Medical Microbiology</i> , 2021, 70, .   | 1.8  | 13        |
| 33 | Household Transmission of Community-Associated Methicillin-Resistant <i>Staphylococcus Aureus</i> . <i>Frontiers in Public Health</i> , 2021, 9, 658638.   | 2.7  | 8         |
| 34 | 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        |
| 35 | A Sequence Type 23 Hypervirulent <i>Klebsiella pneumoniae</i> Strain Presenting Carbapenem Resistance by Acquiring an IncP1 blaKPC-2 Plasmid. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021, 11, 641830.  | 3.9  | 10        |
| 36 | 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       |

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|----|---|-----|-----------|
| 37 | Clinical Impact of Metagenomic Next-Generation Sequencing of Bronchoalveolar Lavage in the Diagnosis and Management of Pneumonia. <i>Journal of Molecular Diagnostics</i> , 2021, 23, 1259-1268.  | 2.8 | 43        |
| 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 | 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         |
| 40 | Co-harboring of Novel blaKPC $\epsilon$ 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        |
| 41 | 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         |
| 42 | 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        |
| 43 | A Novel SXT/R391 Integrative and Conjugative Element Carries Two Copies of the bla <sub>NDM-1</sub> Gene in <i>Proteus mirabilis</i> . <i>MSphere</i> , 2021, 6, e0058821.  | 2.9 | 23        |
| 44 | 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        |
| 45 | 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         |
| 46 | Genetic diversity of siderophores and hypermucoviscosity phenotype in <i>Klebsiella pneumoniae</i> . <i>Microbial Pathogenesis</i> , 2021, 158, 105014.   | 2.9 | 6         |
| 47 | <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        |
| 48 | 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        |
| 49 | Resistance evolution of hypervirulent carbapenem-resistant <i>Klebsiella pneumoniae</i> ST11 during treatment with tigecycline and polymyxin. <i>Emerging Microbes and Infections</i> , 2021, 10, 1129-1136.  | 6.5 | 49        |
| 50 | 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         |
| 51 | 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         |
| 52 | Distribution of erm 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         |
| 53 | Colistin-phage combinations decrease antibiotic resistance in <i>Acinetobacter baumannii</i> via changes in envelope architecture. <i>Emerging Microbes and Infections</i> , 2021, 10, 2205-2219.   | 6.5 | 50        |
| 54 | 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         |

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|----|--|------|-----------|
| 55 | Emergence of Ceftazidime- and Avibactam-Resistant <i>Klebsiella pneumoniae</i> Carbapenemase-Producing <i>Pseudomonas aeruginosa</i> in China. <i>MSystems</i> , 2021, 6, e0078721.  | 3.8  | 39        |
| 56 | 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         |
| 57 | 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         |
| 58 | 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         |
| 59 | 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         |
| 60 | Genetic Characterization and Passage Instability of a Hybrid Plasmid Co-Harboring <i>bla</i> <sub>IMP-4</sub> and <i>bla</i> <sub>NDM-1</sub> Reveal the Contribution of Insertion Sequences During Plasmid Formation and Evolution. <i>Microbiology Spectrum</i> , 2021, 9, e0157721. | 3.0  | 13        |
| 61 | 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         |
| 62 | 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         |
| 63 | fIDBAC: A Platform for Fast Bacterial Genome Identification and Typing. <i>Frontiers in Microbiology</i> , 2021, 12, 723577.   | 3.5  | 9         |
| 64 | The global dissemination of bacterial infections necessitates the study of reverse genomic epidemiology. <i>Briefings in Bioinformatics</i> , 2020, 21, 741-750.   | 6.5  | 56        |
| 65 | 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         |
| 66 | Clinical characteristic of 15 cases of cryptococcal meningitis treated with Ommaya reservoir. <i>Acta Neurologica Belgica</i> , 2020, 120, 1139-1145.  | 1.1  | 4         |
| 67 | 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         |
| 68 | 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        |
| 69 | 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        |
| 70 | Coexistence of <i>bla</i> <sub>KPC-2</sub> - <i>IncN</i> and <i>mcr-1</i> - <i>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        |
| 71 | 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       |
| 72 | Emergence of a Clinical <i>Escherichia coli</i> Sequence Type 131 Strain Carrying a Chromosomal <i>bla</i> <sub>KPC-2</sub> Gene. <i>Frontiers in Microbiology</i> , 2020, 11, 586764.   | 3.5  | 4         |

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|----|--|------|-----------|
| 73 | The Characterization of OXA-232 Carbapenemase-Producing ST437 <i>Klebsiella pneumoniae</i> in China. Canadian Journal of Infectious Diseases and Medical Microbiology, 2020, 2020, 1-5.  | 1.9  | 18        |
| 74 | A Biological Inventory of Prophages in <i>A. baumannii</i> Genomes Reveal Distinct Distributions in Classes, Length, and Genomic Positions. Frontiers in Microbiology, 2020, 11, 579802.   | 3.5  | 38        |
| 75 | Plasmid Dynamics of <i>mcr-1</i> -Positive <i>Salmonella</i> spp. in a General Hospital in China. Frontiers in Microbiology, 2020, 11, 604710.   | 3.5  | 13        |
| 76 | Molecular characteristics of PaLoc and acquired antimicrobial resistance in epidemic <i>Clostridioides difficile</i> isolates revealed by whole-genome sequencing. Journal of Global Antimicrobial Resistance, 2020, 23, 194-196.  | 2.2  | 3         |
| 77 | 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. International Journal of Medical Microbiology, 2020, 310, 151464.     | 3.6  | 11        |
| 78 | Discovery of a Novel Hypervirulent <i>Acinetobacter baumannii</i> Strain in a Case of Community-Acquired Pneumonia. Infection and Drug Resistance, 2020, Volume 13, 1147-1153.   | 2.7  | 5         |
| 79 | 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. Lancet Microbe, The, 2020, 1, e24-e33.   | 7.3  | 28        |
| 80 | Population Biology and Epidemiological Studies of <i>Acinetobacter baumannii</i> in the Era of Whole Genome Sequencing: Is the Oxford Scheme Still Appropriate?. Frontiers in Microbiology, 2020, 11, 775.   | 3.5  | 10        |
| 81 | Nonclassical Biofilms Induced by DNA Breaks in <i>Klebsiella pneumoniae</i> . MSphere, 2020, 5, .  | 2.9  | 6         |
| 82 | Phenotypic Variation and Carbapenem Resistance Potential in OXA-499-Producing <i>Acinetobacter pittii</i> . Frontiers in Microbiology, 2020, 11, 1134.   | 3.5  | 5         |
| 83 | 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. Journal of Infectious Diseases, 2020, 221, S263-S271.                              | 4.0  | 17        |
| 84 | In-Host Evolution of Daptomycin Resistance and Heteroresistance in Methicillin-Resistant <i>Staphylococcus aureus</i> Strains From Three Endocarditis Patients. Journal of Infectious Diseases, 2020, 221, S243-S252.  | 4.0  | 16        |
| 85 | Prevalence and characteristics of <i>pks</i> gene cluster harbouring <i>Klebsiella pneumoniae</i> from bloodstream infection in China. Epidemiology and Infection, 2020, 148, e69.   | 2.1  | 7         |
| 86 | Mechanical penetration of $\beta$ -lactam-resistant Gram-negative bacteria by programmable nanowires. Science Advances, 2020, 6, .   | 10.3 | 23        |
| 87 | 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. Infection and Drug Resistance, 2020, Volume 13, 99-109. | 2.7  | 25        |
| 88 | Defining persistent critical illness based on growth trajectories in patients with sepsis. Critical Care, 2020, 24, 57.  | 5.8  | 23        |
| 89 | The mismatch repair system ( <i>mutS</i> and <i>mutL</i> ) in <i>Acinetobacter baylyi</i> ADP1. BMC Microbiology, 2020, 20, 40.  | 3.3  | 12        |
| 90 | Effect of <i>ramR</i> loss-of-function insertion on tigecycline resistance in clinical isolates of carbapenem-resistant <i>Klebsiella pneumoniae</i> . Journal of Global Antimicrobial Resistance, 2020, 21, 410-413.  | 2.2  | 10        |

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|-----|---|-----|-----------|
| 91  | Bautype: Capsule and Lipopolysaccharide Serotype Prediction for <i>Acinetobacter baumannii</i> Genome. <i>Infectious Microbes &amp; Diseases</i> , 2020, 2, 18-25.  | 1.3 | 7         |
| 92  | 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         |
| 93  | 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         |
| 94  | &lt;p&gt;Molecular Characterization of Carbapenem-Resistant &lt;em&gt; <i>Serratia marcescens</i> &lt;/em&gt; Clinical Isolates in a Tertiary Hospital in Hangzhou, China&lt;/p&gt;. <i>Infection and Drug Resistance</i> , 2020, Volume 13, 999-1008.  | 2.7 | 19        |
| 95  | 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         |
| 96  | Cointegration as a mechanism for the evolution of a KPC-producing multidrug resistance plasmid in <i> <i>Proteus mirabilis</i> </i>. <i>Emerging Microbes and Infections</i> , 2020, 9, 1206-1218.  | 6.5 | 30        |
| 97  | 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), 2015â€“2018. <i>Clinical Infectious Diseases</i> , 2020, 71, S427-S435. | 5.8 | 20        |
| 98  | &lt;p&gt;Clinical and Microbiological Characteristics of Community-Onset Carbapenem-Resistant Enterobacteriaceae Isolates&lt;/p&gt;. <i>Infection and Drug Resistance</i> , 2020, Volume 13, 3131-3143.   | 2.7 | 14        |
| 99  | 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         |
| 100 | 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        |
| 101 | Prevalence and molecular characteristics of &lt;em&gt; <i>mcr-1</i> &lt;/em&gt; gene in &lt;em&gt; <i>Salmonella typhimurium</i> &lt;/em&gt; in a tertiary hospital of Zhejiang Province. <i>Infection and Drug Resistance</i> , 2019, Volume 12, 105-110.  | 2.7 | 16        |
| 102 | Increasing prevalence of <i>Neisseria gonorrhoeae</i> with decreased susceptibility to ceftriaxone and resistance to azithromycin in Hangzhou, China (2015â€“17). <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 29-37.   | 3.0 | 17        |
| 103 | Risk factors for acquisition and mortality of multidrug-resistant <i>Acinetobacter baumannii</i> bacteremia. <i>Medicine (United States)</i> , 2019, 98, e14937.  | 1.0 | 50        |
| 104 | Dual Role of <i> <i>gnaA</i> </i> in Antibiotic Resistance and Virulence in <i>Acinetobacter baumannii</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2019, 63, .   | 3.2 | 23        |
| 105 | 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        |
| 106 | A Highly Efficient CRISPR-Cas9-Based Genome Engineering Platform in <i>Acinetobacter baumannii</i> to Understand the H2O2-Sensing Mechanism of OxyR. <i>Cell Chemical Biology</i> , 2019, 26, 1732-1742.e5.   | 5.2 | 55        |
| 107 | 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         |
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