## Xiaofei Jiang

## List of Publications by Year in descending order

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471509 395702 1,224 33 17 33 citations h-index g-index papers 33 33 33 1431 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Resistance reported from China antimicrobial surveillance network (CHINET) in 2018. European Journal of Clinical Microbiology and Infectious Diseases, 2019, 38, 2275-2281.	2.9	185
2	Complete Genome Sequence of Klebsiella pneumoniae subsp. pneumoniae HS11286, a Multidrug-Resistant Strain Isolated from Human Sputum. Journal of Bacteriology, 2012, 194, 1841-1842.	2.2	152
3	Detection of Extended-Spectrum $\hat{l}^2$ -Lactamases in Clinical Isolates of Pseudomonas aeruginosa. Antimicrobial Agents and Chemotherapy, 2006, 50, 2990-2995.	3.2	124
4	Pandemic spread of bla among Klebsiella pneumoniae ST11 in China is associated with horizontal transfer mediated by IncFII-like plasmids. International Journal of Antimicrobial Agents, 2019, 54, 117-124.	2.5	67
5	Outbreak of Infection Caused by <i>Enterobacter cloacae</i> Producing the Novel VEB-3 Beta-Lactamase in China. Journal of Clinical Microbiology, 2005, 43, 826-831.	3.9	62
6	Mapping the resistance-associated mobilome of a carbapenem-resistant <i>Klebsiella pneumoniae</i> strain reveals insights into factors shaping these regions and facilitates generation of a â€resistance-disarmed' model organism. Journal of Antimicrobial Chemotherapy, 2015, 70, 2770-2774.	3.0	55
7	Contribution of $\hat{l}^2$ -Lactamases and Porin Proteins OmpK35 and OmpK36 to Carbapenem Resistance in Clinical Isolates of KPC-2-Producing Klebsiella pneumoniae. Antimicrobial Agents and Chemotherapy, 2014, 58, 1214-1217.	3.2	54
8	First Report of a Clinical, Multidrug-Resistant Enterobacteriaceae Isolate Coharboring Fosfomycin Resistance GenefosA3and Carbapenemase GeneblaKPC-2on the Same Transposon, Tn1721. Antimicrobial Agents and Chemotherapy, 2015, 59, 338-343.	3.2	37
9	Human Mesenchymal Stem Cell-derived Exosomes Reduce Ischemia/Reperfusion Injury by the Inhibitions of Apoptosis and Autophagy. Current Pharmaceutical Design, 2019, 24, 5334-5341.	1.9	37
10	Characterization of the genetic environment of the blaKPC-2 gene among Klebsiella pneumoniae isolates from a Chinese Hospital. Brazilian Journal of Infectious Diseases, 2016, 20, 384-388.	0.6	36
11	The association of metabolic syndrome components and chronic kidney disease in patients with hypertension. Lipids in Health and Disease, 2019, 18, 229.	3.0	35
12	Salidroside inhibits high-glucose induced proliferation of vascular smooth muscle cells via inhibiting mitochondrial fission and oxidative stress. Experimental and Therapeutic Medicine, 2017, 14, 515-524.	1.8	33
13	The type I-E CRISPR-Cas system influences the acquisition of <i>bla</i> <sub>KPC</sub> -IncF plasmid in <i>Klebsiella pneumonia</i> Emerging Microbes and Infections, 2020, 9, 1011-1022.	6.5	33
14	Absence of the type I-E CRISPR-Cas system in Klebsiella pneumoniae clonal complex 258 is associated with dissemination of IncF epidemic resistance plasmids in this clonal complex. Journal of Antimicrobial Chemotherapy, 2020, 75, 890-895.	3.0	33
15	Genetic diversity and evolution of the virulence plasmids encoding aerobactin and salmochelin in <i>Klebsiella pneumoniae</i> i>Nirulence, 2021, 12, 1323-1333.	4.4	33
16	Prevalence of hypervirulent and carbapenem-resistant <i>Klebsiella pneumoniae</i> under divergent evolutionary patterns. Emerging Microbes and Infections, 2022, 11, 1936-1949.	6.5	32
17	Identification of hypervirulent Klebsiella pneumoniae isolates using the string test in combination with Galleria mellonella infectivity. European Journal of Clinical Microbiology and Infectious Diseases, 2020, 39, 1673-1679.	2.9	31
18	Anti-Restriction Protein, KlcAHS, Promotes Dissemination of Carbapenem Resistance. Frontiers in Cellular and Infection Microbiology, 2017, 7, 150.	3.9	25

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19	Translocation of Carbapenemase Gene <i>bla</i> <sub>KPC-2</sub> both Internal and External to Transposons Occurs via Novel Structures of Tn <i>1721</i> and Exhibits Distinct Movement Patterns. Antimicrobial Agents and Chemotherapy, 2017, 61, .	3.2	24
20	Acquisition of the Conjugative Virulence Plasmid From a CG23 Hypervirulent Klebsiella pneumoniae Strain Enhances Bacterial Virulence. Frontiers in Cellular and Infection Microbiology, 2021, 11, 752011.	3.9	21
21	Molecular Epidemiology of Hypervirulent Carbapenemase-Producing Klebsiella pneumoniae. Frontiers in Cellular and Infection Microbiology, 2021, 11, 661218.	3.9	18
22	Dynamin-related protein inhibitor downregulates reactive oxygen species levels to indirectly suppress high glucose-induced hyperproliferation of vascular smooth muscle cells. Biochemical and Biophysical Research Communications, 2016, 471, 474-478.	2.1	13
23	A putative multi-replicon plasmid co-harboring beta-lactamase genes blaKPC-2, blaCTX-M-14 and blaTEM-1 and trimethoprim resistance gene dfrA25 from a Klebsiella pneumoniae sequence type (ST) 11 strain in China. PLoS ONE, 2017, 12, e0171339.	2.5	13
24	Convergence of carbapenem resistance and hypervirulence leads to high mortality in patients with postoperative Klebsiella pneumoniae meningitis. Journal of Global Antimicrobial Resistance, 2021, 27, 95-100.	2,2	11
25	ST-segment elevation myocardial infarction in patient with essential thrombocythemia without associated risk. International Journal of Cardiology, 2015, 180, 223-225.	1.7	10
26	Prevention options for ventriculoperitoneal shunt infections: a retrospective analysis during a five-year period. International Journal of Clinical and Experimental Medicine, 2015, 8, 19775-80.	1.3	10
27	High-risk KPC-producing Klebsiella pneumoniae lack type I R-M systems. International Journal of Antimicrobial Agents, 2020, 56, 106050.	2.5	8
28	Replicative transposition contributes to the evolution and dissemination of KPC-2-producing plasmid in <i>Enterobacterales</i> . Emerging Microbes and Infections, 2022, 11, 113-122.	6.5	8
29	Co-occurrence of a novel VIM-1 and FosA3-encoding multidrug-resistant plasmid and a KPC-2-encoding pKPO48-like plasmid in a clinical isolate of Klebsiella pneumoniae sequence type 11. Infection, Genetics and Evolution, 2020, 85, 104479.	2.3	7
30	A Site-Specific Integrative Plasmid Found in Pseudomonas aeruginosa Clinical Isolate HS87 along with A Plasmid Carrying an Aminoglycoside-Resistant Gene. PLoS ONE, 2016, 11, e0148367.	2.5	7
31	Microbial Contamination of Rigid Gas Permeable (RGP) Trial Lenses and Lens Cases in China. Current Eye Research, 2020, 45, 550-555.	1.5	5
32	Molecular dissection of <i>bla </i> <csub>KPC-2 -bearing plasmids evolving in <i>Klebsiella pneumoniae </i><li>isolated at one teaching hospital in Shanghai, China. FEMS Microbiology Letters, 2016, 363, fnw142.</li></csub>	1.8	3
33	KlcAHS genes are ubiquitous in clinical, blaKPC-2-positive, Klebsiella pneumoniae isolates. Infection, Genetics and Evolution, 2019, 70, 84-89.	2.3	2