

# Wenguang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6685514/publications.pdf>

Version: 2024-02-01

17  
papers

910  
citations

1040056

9  
h-index

996975

15  
g-index

21  
all docs

21  
docs citations

21  
times ranked

1342  
citing authors

#	ARTICLE	IF	CITATIONS
1	Antibiotics, Antibiotic Resistance Genes, and Bacterial Community Composition in Fresh Water Aquaculture Environment in China. <i>Microbial Ecology</i> , 2015, 70, 425-432.	2.8	322
2	Antibiotic-mediated changes in the fecal microbiome of broiler chickens define the incidence of antibiotic resistance genes. <i>Microbiome</i> , 2018, 6, 34.	11.1	185
3	Selective pressure of antibiotics on ARGs and bacterial communities in manure-polluted freshwater-sediment microcosms. <i>Frontiers in Microbiology</i> , 2015, 6, 194.	3.5	98
4	Metagenomic insights into the distribution of antibiotic resistome between the gut-associated environments and the pristine environments. <i>Environment International</i> , 2019, 126, 346-354.	10.0	82
5	Fate of potential indicator antimicrobial resistance genes (ARGs) and bacterial community diversity in simulated manure-soil microcosms. <i>Ecotoxicology and Environmental Safety</i> , 2018, 147, 817-823.	6.0	50
6	Metagenomic insights into the effect of oxytetracycline on microbial structures, functions and functional genes in sediment denitrification. <i>Ecotoxicology and Environmental Safety</i> , 2018, 161, 85-91.	6.0	45
7	Swine manure facilitates the spread of antibiotic resistome including tigecycline-resistant tet(X) variants to farm workers and receiving environment. <i>Science of the Total Environment</i> , 2022, 808, 152157.	8.0	35
8	Co-existence of the oxazolidinone resistance genes cfr and optrA on two transferable multi-resistance plasmids in one <i>Enterococcus faecalis</i> isolate from swine. <i>International Journal of Antimicrobial Agents</i> , 2020, 56, 105993.	2.5	20
9	Fate of antimicrobial resistance genes in response to application of poultry and swine manure in simulated manure-soil microcosms and manure-pond microcosms. <i>Environmental Science and Pollution Research</i> , 2017, 24, 20949-20958.	5.3	15
10	The co-occurrence of antibiotic resistance genes between dogs and their owners in families. , 2022, 1, .		14
11	Metagenomic Insights into Chicken Gut Antibiotic Resistomes and Microbiomes. <i>Microbiology Spectrum</i> , 2022, , e0190721.	3.0	10
12	Draft genome sequence of an OXA-23, OXA-66, ADC-25 and TEM-1D co-producing <i>Acinetobacter baumannii</i> ST195 isolated from a patient with neonatal pneumonia in China. <i>Journal of Global Antimicrobial Resistance</i> , 2019, 16, 1-3.	2.2	8
13	Presence and distribution of Macrolides-Lincosamide-Streptogramin resistance genes and potential indicator ARGs in the university ponds in Guangzhou, China. <i>Environmental Science and Pollution Research</i> , 2016, 23, 22937-22946.	5.3	7
14	Isopropoxy Benzene Guanidine Kills <i>Staphylococcus aureus</i> Without Detectable Resistance. <i>Frontiers in Microbiology</i> , 2021, 12, 633467.	3.5	7
15	Dramatic decrease in colistin resistance in <i>Escherichia coli</i> from a typical pig farm following restriction of colistin use in China. <i>International Journal of Antimicrobial Agents</i> , 2019, 53, 707-708.	2.5	6
16	Emergence of bla <sub>NDM</sub> -carrying IncX3 plasmid in <i>Klebsiella pneumoniae</i> and <i>Klebsiella quasipneumoniae</i> from duck farms in Guangdong Province, China. <i>Journal of Global Antimicrobial Resistance</i> , 2020, 22, 703-705.	2.2	6
17	A Natural Antimicrobial Agent: Analysis of Antibacterial Effect and Mechanism of Compound Phenolic Acid on <i>Escherichia coli</i> Based on Tandem Mass Tag Proteomics. <i>Frontiers in Microbiology</i> , 2021, 12, 738896.	3.5	0