Yunxue Guo

List of Publications by Year in descending order

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		218381	197535
55	2,652 citations	26	49
papers	citations	h-index	g-index
55	55	55	3011
33	33	33	3011
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	The HipAB Toxin–Antitoxin System Stabilizes a Composite Genomic Island in Shewanella putrefaciens CN-32. Frontiers in Microbiology, 2022, 13, 858857.	1.5	5
2	Filamentous prophage capsid proteins contribute to superinfection exclusion and phage defence in <i>Pseudomonas aeruginosa</i> . Environmental Microbiology, 2022, 24, 4285-4298.	1.8	10
3	Antitoxin CrlA of CrlTA Toxin–Antitoxin System in a Clinical Isolate Pseudomonas aeruginosa Inhibits Lytic Phage Infection. Frontiers in Microbiology, 2022, 13, .	1.5	7
4	The coral pathogen Vibrio corallilyticus kills non-pathogenic holobiont competitors by triggering prophage induction. Nature Ecology and Evolution, 2022, 6, 1132-1144.	3.4	20
5	Type VII Toxin/Antitoxin Classification System for Antitoxins that Enzymatically Neutralize Toxins. Trends in Microbiology, 2021, 29, 388-393.	3.5	58
6	Xenogeneic silencing relies on temperature-dependent phosphorylation of the host H-NS protein in <i>Shewanella</i> . Nucleic Acids Research, 2021, 49, 3427-3440.	6.5	11
7	Rapid detection of temperate bacteriophage using a simple motility assay. Environmental Microbiology Reports, 2021, 13, 728-734.	1.0	2
8	Prophage Tracer: precisely tracing prophages in prokaryotic genomes using overlapping split-read alignment. Nucleic Acids Research, 2021, 49, e128-e128.	6.5	12
9	Conjugative plasmid-encoded toxin–antitoxin system PrpT/PrpA directly controls plasmid copy number. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	25
10	Antagonism between coral pathogen Vibrio coralliilyticus and other bacteria in the gastric cavity of scleractinian coral Galaxea fascicularis. Science China Earth Sciences, 2020, 63, 157-166.	2.3	21
11	Novel polyadenylylation-dependent neutralization mechanism of the HEPN/MNT toxin/antitoxin system. Nucleic Acids Research, 2020, 48, 11054-11067.	6.5	27
12	Upregulation of a marine fungal biosynthetic gene cluster by an endobacterial symbiont. Communications Biology, 2020, 3, 527.	2.0	12
13	Identification of bacteria-derived urease in the coral gastric cavity. Science China Earth Sciences, 2020, 63, 1553-1563.	2.3	10
14	Prophage encoding toxin/antitoxin system PfiT/PfiA inhibits Pf4 production in <i>Pseudomonas aeruginosa </i> . Microbial Biotechnology, 2020, 13, 1132-1144.	2.0	30
15	Symbiosis of a P2â€family phage and deepâ€sea <i>Shewanella putrefaciens</i> . Environmental Microbiology, 2019, 21, 4212-4232.	1.8	16
16	Resistance to oxidative stress by inner membrane protein ElaB is regulated by OxyR and RpoS. Microbial Biotechnology, 2019, 12, 392-404.	2.0	21
17	Eliminating mcr-1-harbouring plasmids in clinical isolates using the CRISPR/Cas9 system. Journal of Antimicrobial Chemotherapy, 2019, 74, 2559-2565.	1.3	48
18	Biofilm formation in Pseudoalteromonas lipolytica is related to IS5-like insertions in the capsular polysaccharide operon. FEMS Microbiology Ecology, 2019, 95, .	1.3	7

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19	Structure and allosteric coupling of type â; antitoxin CopASO. Biochemical and Biophysical Research Communications, 2019, 514, 1122-1127.	1.0	5
20	Phages Mediate Bacterial Self-Recognition. Cell Reports, 2019, 27, 737-749.e4.	2.9	20
21	Characterization of Two Toxin-Antitoxin Systems in Deep-Sea Streptomyces sp. SCSIO 02999. Marine Drugs, 2019, 17, 211.	2,2	4
22	Antitoxin HigA inhibits virulence gene <i>mvfR</i> expression in <i>Pseudomonas aeruginosa</i> Environmental Microbiology, 2019, 21, 2707-2723.	1.8	39
23	Recent progress on signalling molecules of coral-associated microorganisms. Science China Earth Sciences, 2019, 62, 609-618.	2.3	6
24	Excisionase in Pf filamentous prophage controls lysisâ€lysogeny decisionâ€making in <i>Pseudomonas aeruginosa</i> . Molecular Microbiology, 2019, 111, 495-513.	1.2	34
25	Colistin Resistance Gene mcr-1 Mediates Cell Permeability and Resistance to Hydrophobic Antibiotics. Frontiers in Microbiology, 2019, 10, 3015.	1.5	49
26	Type II toxin/antitoxin system ParE _{SO} /CopA _{SO} stabilizes prophage CP4So in <i>Shewanella oneidensis</i> . Environmental Microbiology, 2018, 20, 1224-1239.	1.8	39
27	Structure–function analyses reveal the molecular architecture and neutralization mechanism of a bacterial HEPN–MNT toxin–antitoxin system. Journal of Biological Chemistry, 2018, 293, 6812-6823.	1.6	24
28	Characteristics and Trends of Ambient Ozone and Nitrogen Oxides at Urban, Suburban, and Rural Sites from 2011 to 2017 in Shenzhen, China. Sustainability, 2018, 10, 4530.	1.6	23
29	Antimicrobial Resistance Profile of mcr-1 Positive Clinical Isolates of Escherichia coli in China From 2013 to 2016. Frontiers in Microbiology, 2018, 9, 2514.	1.5	28
30	Marine Bacteria Provide Lasting Anticorrosion Activity for Steel via Biofilm-Induced Mineralization. ACS Applied Materials & Samp; Interfaces, 2018, 10, 40317-40327.	4.0	87
31	Tail-Anchored Inner Membrane Protein ElaB Increases Resistance to Stress While Reducing Persistence in Escherichia coli. Journal of Bacteriology, 2017, 199, .	1.0	31
32	Dissemination and loss of a biofilmâ€related genomic island in marine <i>Pseudoalteromonas</i> mediated by integrative and conjugative elements. Environmental Microbiology, 2017, 19, 4620-4637.	1.8	10
33	Pyomelanin from <i>Pseudoalteromonas lipolytica</i> reduces biofouling. Microbial Biotechnology, 2017, 10, 1718-1731.	2.0	35
34	Interaction of Type IV Toxin/Antitoxin Systems in Cryptic Prophages of Escherichia coli K-12. Toxins, 2017, 9, 77.	1.5	27
35	Quantifying Grain-Size Variability of Metal Pollutants in Road-Deposited Sediments Using the Coefficient of Variation. International Journal of Environmental Research and Public Health, 2017, 14, 850.	1.2	21
36	Characterization of the Deep-Sea Streptomyces sp. SCSIO 02999 Derived VapC/VapB Toxin-Antitoxin System in Escherichia coli. Toxins, 2016, 8, 195.	1.5	10

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37	Cryptic prophages as targets for drug development. Drug Resistance Updates, 2016, 27, 30-38.	6.5	58
38	Complete genome sequence of Vibrio alginolyticus ATCC 33787T isolated from seawater with three native megaplasmids. Marine Genomics, 2016, 28, 45-47.	0.4	16
39	Complete genome sequence of Pseudoalteromonas rubra SCSIO 6842, harboring a putative conjugative plasmid pMBL6842. Journal of Biotechnology, 2016, 224, 66-67.	1.9	8
40	Cold adaptation regulated by cryptic prophage excision in <i>Shewanella oneidensis</i> . ISME Journal, 2016, 10, 2787-2800.	4.4	72
41	Development and application of watershed-scale indicator to quantify non-point source P losses in semi-humid and semi-arid watershed, China. Ecological Indicators, 2016, 63, 374-385.	2.6	11
42	Physiological Function of Rac Prophage During Biofilm Formation and Regulation of Rac Excision in Escherichia coli K-12. Scientific Reports, 2015, 5, 16074.	1.6	28
43	Identification and characterization of a <scp>HEPNâ€MNT</scp> family type <scp>II</scp> toxin–antitoxin in <scp><i>S</i></scp> <i>hewanella oneidensis</i> . Microbial Biotechnology, 2015, 8, 961-973.	2.0	34
44	Spatial and temporal characteristics of droughts in the Northeast China Transect. Natural Hazards, 2015, 76, 601-614.	1.6	28
45	Development of an efficient conjugation-based genetic manipulation system for Pseudoalteromonas. Microbial Cell Factories, 2015, 14, 11.	1.9	81
46	Characterization of self-generated variants in Pseudoalteromonas lipolytica biofilm with increased antifouling activities. Applied Microbiology and Biotechnology, 2015, 99, 10127-10139.	1.7	39
47	RalR (a DNase) and RalA (a small RNA) form a type I toxin–antitoxin system in Escherichia coli. Nucleic Acids Research, 2014, 42, 6448-6462.	6.5	98
48	Estimation of the nonpoint source nitrogen load in a strongly disturbed watershed of the North China Plain. Water Science and Technology, 2014, 69, 1304-1311.	1.2	6
49	Precipitation partitioning in a diverse stand and a monospecific stand of regenerating forest in eastern China. Southern Forests, 2013, 75, 81-87.	0.2	0
50	Toxin-Antitoxin Systems Influence Biofilm and Persister Cell Formation and the General Stress Response. Applied and Environmental Microbiology, 2011, 77, 5577-5583.	1.4	368
51	Antitoxin MqsA helps mediate the bacterial general stress response. Nature Chemical Biology, 2011, 7, 359-366.	3.9	201
52	IS <i>>5</i> in a quasi-Lamarckian way. ISME Journal, 2011, 5, 1517-1525.	4.4	46
53	Controlling biofilm formation, prophage excision and cell death by rewiring global regulator Hâ€NS of <i>Escherichia coli</i> . Microbial Biotechnology, 2010, 3, 344-356.	2.0	66
54	Cryptic prophages help bacteria cope with adverse environments. Nature Communications, 2010, 1, 147.	5.8	560

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55	Control and benefits of CP4-57 prophage excision in <i>Escherichia coli</i> biofilms. ISME Journal, 2009, 3, 1164-1179.	4.4	98