

Linhuan Wu

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4911571/publications.pdf>

Version: 2024-02-01

19
papers

1,211
citations

759233

12
h-index

888059

17
g-index

23
all docs

23
docs citations

23
times ranked

1532
citing authors

#	ARTICLE	IF	CITATIONS
1	Enlightening the taxonomy darkness of human gut microbiomes with a cultured biobank. <i>Microbiome</i> , 2021, 9, 119.	11.1	479
2	Mycobank gearing up for new horizons. <i>IMA Fungus</i> , 2013, 4, 371-379.	3.8	170
3	Dynamic reassortments and genetic heterogeneity of the human-infecting influenza A (H7N9) virus. <i>Nature Communications</i> , 2014, 5, 3142.	12.8	145
4	gcMeta: a Global Catalogue of Metagenomics platform to support the archiving, standardization and analysis of microbiome data. <i>Nucleic Acids Research</i> , 2019, 47, D637-D648.	14.5	70
5	The Global Catalogue of Microorganisms (GCM) 10K type strain sequencing project: providing services to taxonomists for standard genome sequencing and annotation. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 69, 895-898.	1.7	58
6	The Mouse Gut Microbial Biobank expands the coverage of cultured bacteria. <i>Nature Communications</i> , 2020, 11, 79.	12.8	55
7	gcType: a high-quality type strain genome database for microbial phylogenetic and functional research. <i>Nucleic Acids Research</i> , 2021, 49, D694-D705.	14.5	53
8	World data centre for microorganisms: an information infrastructure to explore and utilize preserved microbial strains worldwide. <i>Nucleic Acids Research</i> , 2017, 45, D611-D618.	14.5	46
9	The global catalogue of microorganisms 10K type strain sequencing project: closing the genomic gaps for the validly published prokaryotic and fungi species. <i>GigaScience</i> , 2018, 7, .	6.4	35
10	VarEPS: an evaluation and prewarning system of known and virtual variations of SARS-CoV-2 genomes. <i>Nucleic Acids Research</i> , 2022, 50, D888-D897.	14.5	18
11	Mycobacterial biomaterials and resources for researchers. <i>Pathogens and Disease</i> , 2018, 76, .	2.0	14
12	Genetic tracing of HCoV-19 for the re-emerging outbreak of COVID-19 in Beijing, China. <i>Protein and Cell</i> , 2021, 12, 4-6.	11.0	13
13	Web Resources for Microbial Data. <i>Genomics, Proteomics and Bioinformatics</i> , 2015, 13, 69-72.	6.9	12
14	Description of the First Strain of 2019-nCoV, C-Tan-nCoV Wuhan Strain - National Pathogen Resource Center, China, 2020. <i>China CDC Weekly</i> , 2020, 2, 81-82.	2.3	10
15	<i>Allobacillus salarius</i> sp. nov., and <i>Allobacillus saliphilus</i> sp. nov., isolated from shrimp paste (ka-pi) in Thailand. <i>Archives of Microbiology</i> , 2022, 204, 71.	2.2	10
16	Laboratory information management system for biosafety laboratory: Safety and efficiency. <i>Journal of Biosafety and Biosecurity</i> , 2021, 3, 28-34.	2.8	8
17	Computational predicting the human infectivity of H7N9 influenza viruses isolated from avian hosts. <i>Transboundary and Emerging Diseases</i> , 2021, 68, 846-856.	3.0	6
18	The monkey microbial biobank brings previously uncultivated bioresources for nonhuman primate and human gut microbiomes. , 2022, 1, 210-217.		3

#	ARTICLE	IF	CITATIONS
19	gcCov: Linked open data for global coronavirus studies. , 2022, 1, 92-95.		2