Yun-Zeng Zhang

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8929895/publications.pdf

Version: 2024-02-01

22 2,856 13 papers citations h-index

22 22 6284
all docs docs citations times ranked citing authors

22

g-index

#	Article	IF	Citations
1	Closely related Salmonella Derby strains triggered distinct gut microbiota alteration. Gut Pathogens, 2022, 14, 6.	1.6	5
2	The phyllosphere microbiome shifts toward combating melanose pathogen. Microbiome, 2022, 10, 56.	4.9	54
3	Salmonella Pullorum spiC mutant is a desirable LASV candidate with proper virulence, high immune protection and easy-to-use oral administration. Vaccine, 2021, 39, 1383-1391.	1.7	8
4	The Citrus Microbiome: From Structure and Function to Microbiome Engineering and Beyond. Phytobiomes Journal, 2021, 5, 249-262.	1.4	16
5	Gut microbiota mediates cognitive impairment in young mice after multiple neonatal exposures to sevoflurane. Aging, 2021, 13, 16733-16748.	1.4	8
6	Implication of immune cell signature of tumor microenvironment in diffuse large Bâ€eell lymphoma. Hematological Oncology, 2021, 39, 616-624.	0.8	7
7	Transcriptomic Analysis of Streptococcus suis in Response to Ferrous Iron and Cobalt Toxicity. Genes, 2020, 11, 1035.	1.0	2
8	Mechanisms Underlying the Rhizosphere-To-Rhizoplane Enrichment of Cellvibrio Unveiled by Genome-Centric Metagenomics and Metatranscriptomics. Microorganisms, 2020, 8, 583.	1.6	14
9	Diffusible signal factor (DSF)-mediated quorum sensing modulates expression of diverse traits in Xanthomonas citri and responses of citrus plants to promote disease. BMC Genomics, 2019, 20, 55.	1.2	35
10	Origin and diversification of Xanthomonas citri subsp. citri pathotypes revealed by inclusive phylogenomic, dating, and biogeographic analyses. BMC Genomics, 2019, 20, 700.	1.2	33
11	The structure and function of the global citrus rhizosphere microbiome. Nature Communications, 2018, 9, 4894.	5.8	304
12	Tale of the Huanglongbing Disease Pyramid in the Context of the Citrus Microbiome. Phytopathology, 2017, 107, 380-387.	1.1	79
13	The <i>Candidatus</i> Liberibacter–Host Interface: Insights into Pathogenesis Mechanisms and Disease Control. Annual Review of Phytopathology, 2017, 55, 451-482.	3.5	246
14	Genome editing of the disease susceptibility gene <i>Cs<scp>LOB</scp>1</i> in citrus confers resistance to citrus canker. Plant Biotechnology Journal, 2017, 15, 817-823.	4.1	371
15	Taxonomic structure and functional association of foxtail millet root microbiome. GigaScience, 2017, 6, 1-12.	3.3	1,228
16	Editing Citrus Genome via SaCas9/sgRNA System. Frontiers in Plant Science, 2017, 8, 2135.	1.7	87
17	Characterization of Antimicrobial-Producing Beneficial Bacteria Isolated from Huanglongbing Escape Citrus Trees. Frontiers in Microbiology, 2017, 8, 2415.	1.5	48
18	Huanglongbing impairs the rhizosphere-to-rhizoplane enrichment process of the citrus root-associated microbiome. Microbiome, 2017, 5, 97.	4.9	177

#	Article	IF	CITATION
19	SEC-Translocon Dependent Extracytoplasmic Proteins of Candidatus Liberibacter asiaticus. Frontiers in Microbiology, 2016, 7, 1989.	1.5	72
20	Temporal Transcription Profiling of Sweet Orange in Response to PthA4-Mediated Xanthomonas citri subsp. citri Infection. Phytopathology, 2016, 106, 442-451.	1.1	12
21	Transcriptome analysis of root response to citrus blight based on the newly assembled Swingle citrumelo draft genome. BMC Genomics, 2016, 17, 485.	1.2	15
22	Positive selection is the main driving force for evolution of citrus canker-causing <i>Xanthomonas</i> . ISME Journal, 2015, 9, 2128-2138.	4.4	35